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PETSc/TAO Users Manual

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January 8, 2024

This work was supported by the Office of Advanced Scientific Computing Research, Office of Science, U.S. Department of Energy, under Contract DE-AC02-06CH11357.
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INTRODUCTION TO PETSC

1.1 About This Manual

This manual describes the use of the Portable, Extensible Toolkit for Scientific Computation (PETSc) and the Toolkit for Advanced Optimization (TAO) for the numerical solution of partial differential equations and related problems on high-performance computers. PETSc/TAO is a suite of data structures and routines that provide the building blocks for the implementation of large-scale application codes on parallel (and serial) computers. PETSc uses the MPI standard for all distributed memory communication.

PETSc/TAO includes a large suite of parallel linear solvers, nonlinear solvers, time integrators, and optimization that may be used in application codes written in Fortran, C, C++, and Python (via petsc4py; see Getting Started). PETSc provides many of the mechanisms needed within parallel application codes, such as parallel matrix and vector assembly routines. The library is organized hierarchically, enabling users to employ the level of abstraction that is most appropriate for a particular problem. By using techniques of object-oriented programming, PETSc provides enormous flexibility for users.

PETSc is a sophisticated set of software tools; as such, for some users it initially has a much steeper learning curve than packages such as MATLAB or a simple subroutine library. In particular, for individuals without some computer science background, experience programming in C, C++, python, or Fortran and experience using a debugger such as gdb or lldb, it may require a significant amount of time to take full advantage of the features that enable efficient software use. However, the power of the PETSc design and the algorithms it incorporates may make the efficient implementation of many application codes simpler than “rolling them” yourself.

- For many tasks a package such as MATLAB is often the best tool; PETSc is not intended for the classes of problems for which effective MATLAB code can be written.
- There are several packages (listed on https://petsc.org/), built on PETSc, that may satisfy your needs without requiring directly using PETSc. We recommend reviewing these packages functionality before starting to code directly with PETSc.
- PETSc can be used to provide a “MPI parallel linear solver” in an otherwise sequential, or OpenMP parallel code. This approach cannot provide extremely large improvements in the application time by utilizing large numbers of MPI processes but can still improve the performance. Certainly all parts of a previously sequential code need not be parallelized but the matrix generation portion must be parallelized to expect true scalability to large numbers of MPI processes. See PCMPI for details on how to utilize the PETSc MPI linear solver server.

Since PETSc is under continued development, small changes in usage and calling sequences of routines will occur. PETSc has been supported for twenty-five years; see mailing list information on our website for information on contacting support.
1.2 Getting Started

PETSc consists of a collection of classes, which are discussed in detail in later parts of the manual (The Solvers in PETSc/TAO and Additional Information). The important PETSc classes include:

- index sets (IS), for indexing into vectors, renumbering, permuting, etc;
- vectors (Vec); Vectors and Parallel Data
- matrices (Mat) (generally sparse); Matrices
- Krylov subspace methods (KSP); KSP: Linear System Solvers
- preconditioners, including multigrid, block solvers, patch solvers, and sparse direct solvers (PC);
- nonlinear solvers (SNES); SNES: Nonlinear Solvers
- timesteppers for solving time-dependent (nonlinear) PDEs, including support for differential algebraic equations, and the computation of adjoints (sensitivities/gradients of the solutions) (TS); TS: Scalable ODE and DAE Solvers
- scalable optimization algorithms including a rich set of gradient-based optimizers, Newton-based optimizers and optimization with constraints (Tao). TAO: Optimization Solvers
- code for managing interactions between mesh data structures and vectors, matrices, and solvers (DM); DM Basics

Each class consists of an abstract interface (simply a set of calling sequences; an abstract base class in C++) and an implementation for each algorithm and data structure. This design enables easy comparison and use of different algorithms (for example, to experiment with different Krylov subspace methods, preconditioners, or truncated Newton methods). Hence, PETSc provides a rich environment for modeling scientific applications as well as for rapid algorithm design and prototyping.

The classes enable easy customization and extension of both algorithms and implementations. This approach promotes code reuse and flexibility, and also separates the issues of parallelism from the choice of algorithms. The PETSc infrastructure creates a foundation for building large-scale applications.

It is useful to consider the interrelationships among different pieces of PETSc. Numerical Libraries in PETSc is a diagram of some of these pieces. The figure illustrates the library’s hierarchical organization, which enables users to employ the solvers that are most appropriate for a particular problem.

1.2.1 Suggested Reading

The manual is divided into four parts:

- Introduction to PETSc
- The Solvers in PETSc/TAO
- DM: Interfacing Between Solvers and Models/Discretizations
- Additional Information

Introduction to PETSc describes the basic procedure for using the PETSc library and presents simple examples of solving linear systems with PETSc. This section conveys the typical style used throughout the library and enables the application programmer to begin using the software immediately.

The Solvers in PETSc/TAO explains in detail the use of the various PETSc algebraic objects, such as vectors, matrices, index sets and the PETSc solvers including linear and nonlinear solvers, time integrators, and optimization support. DM: Interfacing Between Solvers and Models/Discretizations details how a user’s models and discretizations can easily be interfaced with the solvers by using the DM construct. The
Fig. 1.1: Numerical Libraries in PETSc
Additional Information describes a variety of useful information, including profiling, the options database, viewers, error handling, and some details of PETSc design.

PETSc has evolved to become quite a comprehensive package, and therefore this manual can be rather intimidating for new users. Bear in mind that PETSc can be used efficiently before one understands all of the material presented here. Furthermore, the definitive reference for any PETSc function is always the online manual page. Manual pages for all PETSc functions can be accessed here. The manual pages provide hyperlinked indices (organized by both concept and routine name) to the tutorial examples and enable easy movement among related topics.

Visual Studio Code, Eclipse, Emacs, and Vim users may find their development environment’s options for searching in the source code are useful for exploring the PETSc source code. Details of these feature are provided in Developer Environments.

The complete PETSc distribution, manual pages, and additional information are available via the PETSc home page. The PETSc home page also contains details regarding installation, new features and changes in recent versions of PETSc, machines that we currently support, and a frequently asked questions (FAQ) list.

Note to Fortran Programmers: In most of the manual, the examples and calling sequences are given for the C/C++ family of programming languages. However, Fortran programmers can use all of the functionality of PETSc from Fortran, with only minor differences in the user interface. PETSc for Fortran Users provides a discussion of the differences between using PETSc from Fortran and C, as well as several complete Fortran examples.

Note to Python Programmers: To program with PETSc in Python you need to enable Python bindings (i.e petsc4py) with the configure option --with-petsc4py=1. See the PETSc installation guide for more details.

1.2.2 Running PETSc Programs

Before using PETSc, the user must first set the environmental variable $PETSC_DIR, indicating the full path of the PETSc home directory. For example, under the Unix bash shell a command of the form

```
$ export PETSC_DIR=$HOME/petsc
```

can be placed in the user’s .bashrc or other startup file. In addition, the user may need to set the environment variable $PETSC_ARCH to specify a particular configuration of the PETSc libraries. Note that $PETSC_ARCH is just a name selected by the installer to refer to the libraries compiled for a particular set of compiler options and machine type. Using different values of $PETSC_ARCH allows one to switch between several different sets (say debug and optimized) of libraries easily. To determine if you need to set $PETSC_ARCH, look in the directory indicated by $PETSC_DIR, if there are subdirectories beginning with arch then those subdirectories give the possible values for $PETSC_ARCH.

See handson to immediately jump in and run PETSc code.

All PETSc programs use the MPI (Message Passing Interface) standard for message-passing communication [For94]. Thus, to execute PETSc programs, users must know the procedure for beginning MPI jobs on their selected computer system(s). For instance, when using the MPICH implementation of MPI and many others, the following command initiates a program that uses eight processors:

```
$ mpiexec -n 8 ./petsc_program_name petsc_options
```

PETSc also comes with a script that automatically uses the correct mpiexec for your configuration.

```
$ $PETSC_DIR/lib/petsc/bin/petscmpiexec -n 8 ./petsc_program_name petsc_options
```

All PETSc-compliant programs support the use of the -help option as well as the -version option.
Certain options are supported by all PETSc programs. We list a few particularly useful ones below; a complete list can be obtained by running any PETSc program with the option `-help`.

- `-log_view` - summarize the program’s performance (see Profiling)
- `-fp_trap` - stop on floating-point exceptions; for example divide by zero
- `-malloc_dump` - enable memory tracing; dump list of unfreed memory at conclusion of the run, see Detecting Memory Allocation Problems and Memory Usage,
- `-malloc_debug` - enable memory debugging (by default this is activated for the debugging version of PETSc), see Detecting Memory Allocation Problems and Memory Usage,
- `-start_in_debugger [noxterm,gdb,lldb] [-display name]` - start all processes in debugger. See Debugging, for more information on debugging PETSc programs.
- `-on_error_attach_debugger [noxterm,gdb,lldb] [-display name]` - start debugger only on encountering an error
- `-info` - print a great deal of information about what the program is doing as it runs

### 1.2.3 Writing PETSc Programs

Most PETSc programs begin with a call to

```c
PetscInitialize(int *argc, char ***argv, char *file, char *help);
```

which initializes PETSc and MPI. The arguments `argc` and `argv` are the command line arguments delivered in all C and C++ programs. The argument `file` optionally indicates an alternative name for the PETSc options file, `.petscrc`, which resides by default in the user’s home directory. Runtime Options provides details regarding this file and the PETSc options database, which can be used for runtime customization. The final argument, `help`, is an optional character string that will be printed if the program is run with the `-help` option. In Fortran the initialization command has the form

```fortran
call PetscInitialize(character(*) file, integer ierr)
```

Where the file argument is optional. `PetscInitialize()` automatically calls `MPI_Init()` if MPI has not been not previously initialized. In certain circumstances in which MPI needs to be initialized directly (or is initialized by some other library), the user can first call `MPI_Init()` (or have the other library do it), and then call `PetscInitialize()`. By default, `PetscInitialize()` sets the PETSc “world” communicator `PETSC_COMM_WORLD` to `MPI_COMM_WORLD`.

For those not familiar with MPI, a **communicator** is a way of indicating a collection of processes that will be involved together in a calculation or communication. Communicators have the variable type `MPI_Comm`. In most cases users can employ the communicator `PETSC_COMM_WORLD` to indicate all processes in a given run and `PETSC_COMM_SELF` to indicate a single process.

MPI provides routines for generating new communicators consisting of subsets of processors, though most users rarely need to use these. The book *Using MPI*, by Lusk, Gropp, and Skjellum [GLS94] provides an excellent introduction to the concepts in MPI. See also the MPI homepage. Note that PETSc users need not program much message passing directly with MPI, but they must be familiar with the basic concepts of message passing and distributed memory computing.

All PETSc programs should call `PetscFinalize()` as their final (or nearly final) statement. This routine handles options to be called at the conclusion of the program, and calls `MPI_Finalize()` if `PetscInitialize()` began MPI. If MPI was initiated externally from PETSc (by either the user or another software package), the user is responsible for calling `MPI_Finalize()`.
Error Checking

Most PETSc functions return a PetscErrorCode, which is an integer indicating whether an error has occurred during the call. The error code is set to be nonzero if an error has been detected; otherwise, it is zero. For the C/C++ interface, the error variable is the routine’s return value, while for the Fortran version, each PETSc routine has as its final argument an integer error variable.

One should always check these routine values as given below in the C/C++ and Fortran formats, respectively:

```
PetscCall(PetscFunction(Args));
```

or

```
! within the main program
PetscCallA(PetscFunction(Args,ierr))
```

```
! within any subroutine
PetscCall(PetscFunction(Args,ierr))
```

These macros check the returned error code and if it is nonzero they call the PETSc error handler and then return from the function with the error code. PetscCallA() calls abort after calling the error handler because it is not possible to return from a Fortran main program. The above macros should be used in all subroutines to enable a complete error traceback. See Error Checking for more details on PETSc error handling.

1.2.4 Simple PETSc Examples

To help the user start using PETSc immediately, we begin with a simple uniprocessor example that solves the one-dimensional Laplacian problem with finite differences. This sequential code, which can be found in $PETSC_DIR/src/ksp/ksp/tutorials/ex1.c, illustrates the solution of a linear system with KSP, the interface to the preconditioners, Krylov subspace methods, and direct linear solvers of PETSc. Following the code we highlight a few of the most important parts of this example.

Listing: KSP Tutorial src/ksp/ksp/tutorials/ex1.c

```
static char help[] = "Solves a tridiagonal linear system with KSP."

/*
 * Include "petscksp.h" so that we can use KSP solvers. Note that this file
 * automatically includes:
 *  petcsys.h    - base PETSc routines  petscvec.h - vectors
 *  petsmat.h   - matrices  petscpc.h - preconditioners
 *  petscis.h   - index sets
 *  petscviewer.h - viewers
 *
 * Note: The corresponding parallel example is ex23.c
 */
#include <petscksp.h>

int main(int argc, char **args)
{
  Vec x, b, u; /* approx solution, RHS, exact solution */
  Mat A; /* linear system matrix */
  KSP ksp; /* linear solver context */
  PC pc; /* preconditioner context */
```
PetscReal  norm;  /* norm of solution error */
PetscInt  i, n = 10, col[3], its;
PetscMPIInt size;
PetscScalar value[3];

PetscFunctionBeginUser;
PetscCall(PetscInitialize(&argc, &args, (char *)0, help));
PetscCallMPI(MPI_Comm_size(PETSC_COMM_WORLD, &size));
PetscCheck(size == 1, PETSC_COMM_WORLD, PETSC_ERR_WRONG_MPI_SIZE, "This is a uniprocessor example only!");
PetscCall(PetscOptionsGetInt(NULL, NULL, "-n", &n, NULL));

/* Compute the matrix and right-hand-side vector that define the linear system, Ax = b. */
/** Create vectors. Note that we form 1 vector from scratch and then duplicate as needed. */
PetscCall(VecCreate(PETSC_COMM_SELF, &x));
PetscCall(PetscObjectSetName((PetscObject)x, "Solution");
PetscCall(VecSetSizes(x, PETSC_DECIDE, n));
PetscCall(VecSetFromOptions(x));
PetscCall(VecDuplicate(x, &b));
PetscCall(VecDuplicate(x, &u));

/* Create matrix. When using MatCreate(), the matrix format can be specified at runtime. */
PetscCall(MatCreate(PETSC_COMM_SELF, &A));
PetscCall(MatSetSizes(A, PETSC_DECIDE, PETSC_DECIDE, n, n));
PetscCall(MatSetFromOptions(A));
PetscCall(MatSetUp(A));

/* Assemble matrix */
value[0] = -1.0;
value[1] = 2.0;
value[2] = -1.0;
for (i = 1; i < n - 1; i++) {
  col[0] = i - 1;
  col[1] = i;
  col[2] = i + 1;
  PetscCall(MatSetValues(A, 1, &i, 3, col, value, INSERT_VALUES));
}
i = n - 1;
col[0] = n - 2;
col[1] = n - 1;
PetscCall(MatSetValues(A, 1, &i, 2, col, value, INSERT_VALUES));
i = 0;
col[0] = 0;
col[1] = 1;
value[0] = 2.0;
value[1] = -1.0;
PetscCall(MatSetValues(A, 1, &i, 2, col, value, INSERT_VALUES));
PetscCall(MatAssemblyBegin(A, MAT_FINAL_ASSEMBLY));
PetscCall(MatAssemblyEnd(A, MAT_FINAL_ASSEMBLY));

/*
 Set exact solution; then compute right-hand-side vector.
*/
PetscCall(VecSet(u, 1.0));
PetscCall(MatMult(A, u, b));

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
 Create the linear solver and set various options
 - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - */
PetscCall(KSPCreate(PETSC_COMM_SELF, &ksp));

/*
 Set operators. Here the matrix that defines the linear system
 also serves as the matrix that defines the preconditioner.
*/
PetscCall(KSPSetOperators(ksp, A, A));

/*
 Set linear solver defaults for this problem (optional).
 - By extracting the KSP and PC contexts from the KSP context,
   we can then directly call any KSP and PC routines to set
   various options.
 - The following four statements are optional; all of these
   parameters could alternatively be specified at runtime via
   KSPSetFromOptions();
*/
PetscCall(KSPGetPC(ksp, &pc));
PetscCall(PCSetType(pc, PCJACOBI));
PetscCall(KSPSetTolerances(ksp, 1.e-5, PETSC_DEFAULT, PETSC_DEFAULT, PETSC_DEFAULT));

/*
 Set runtime options, e.g.,
 -ksp_type <type> -pc_type <type> -ksp_monitor -ksp_rtol <rtol>
 These options will override those specified above as long as
 KSPSetFromOptions() is called _after_ any other customization
 routines.
*/
PetscCall(KSPSetFromOptions(ksp));

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
 Solve the linear system
 - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - */
PetscCall(KSPSolve(ksp, b, x));
/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - 
Check the solution and clean up 
- - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - */
PetscCall(VecAXPY(x, -1.0, u));
PetscCall(VecNorm(x, NORM_2, &norm));
PetscCall(KSPGetIterationNumber(ksp, &its));
PetscCall(PetscPrintf(PETSC_COMM_SELF, "Norm of error %g, Iterations %" PetscInt_ 
...FMT "\n", (double)norm, its));
/* check that KSP automatically handles the fact that the the new non-zero values 
in the matrix are propagated to the KSP solver */
PetscCall(MatShift(A, 2.0));
PetscCall(KSPSolve(ksp, b, x));
/* Free work space. All PETSc objects should be destroyed when they 
are no longer needed. */
PetscCall(KSPDestroy(&ksp));
/* test if prefixes properly propagate to PCMPI objects */
if (PCMPIServerActive) {
PetscCall(KSPCreate(PETSC_COMM_SELF, &ksp));
PetscCall(KSPSetOptionsPrefix(ksp, "prefix_test "));
PetscCall(MatSetOptionsPrefix(A, "prefix_test "));
PetscCall(KSPSetOperators(ksp, A, A));
PetscCall(KSPSetFromOptions(ksp));
PetscCall(KSPSolve(ksp, b, x));
PetscCall(KSPDestroy(&ksp));
}
PetscCall(VecDestroy(&x));
PetscCall(VecDestroy(&u));
PetscCall(VecDestroy(&b));
PetscCall(MatDestroy(&A));
/* Always call PetscFinalize() before exiting a program. This routine 
- finalizes the PETSc libraries as well as MPI 
- provides summary and diagnostic information if certain runtime 
  options are chosen (e.g., -log_view). */
PetscCall(PetscFinalize());
return 0;
}
Include Files

The C/C++ include files for PETSc should be used via statements such as

```c
#include <petscksp.h>
```

where `petscksp.h` is the include file for the linear solver library. Each PETSc program must specify an include file that corresponds to the highest level PETSc objects needed within the program; all of the required lower level include files are automatically included within the higher level files. For example, `petscksp.h` includes `petscmat.h` (matrices), `petscvec.h` (vectors), and `petscsys.h` (base PETSc file). The PETSc include files are located in the directory `$PETSC_DIR/include`. See Modules and Include Files for a discussion of PETSc include files in Fortran programs.

The Options Database

As shown in Simple PETSc Examples, the user can input control data at run time using the options database. In this example the command `PetscOptionsGetInt(NULL,NULL,"-n",&n,&flg)` checks whether the user has provided a command line option to set the value of `n`, the problem dimension. If so, the variable `n` is set accordingly; otherwise, `n` remains unchanged. A complete description of the options database may be found in Runtime Options.

Vectors

One creates a new parallel or sequential vector, `x`, of global dimension `M` with the commands

```c
VecCreate(MPI_Comm comm,Vec *x);
VecSetSizes(Vec x, PetscInt m, PetscInt M);
```

where `comm` denotes the MPI communicator and `m` is the optional local size which may be `PETSC_DECIDE`. The type of storage for the vector may be set with either calls to `VecSetType()` or `VecSetFromOptions()`. Additional vectors of the same type can be formed with

```c
VecDuplicate(Vec old,Vec *new);
```

The commands

```c
VecSet(Vec x,PetscScalar value);
VecSetValues(Vec x,PetscInt n,PetscInt *indices,PetscScalar *values,INSERT_VALUES);
```

respectively set all the components of a vector to a particular scalar value and assign a different value to each component. More detailed information about PETSc vectors, including their basic operations, scattering/gathering, index sets, and distributed arrays, is discussed in Chapter Vectors and Parallel Data.

Note the use of the PETSc variable type `PetscScalar` in this example. The `PetscScalar` is simply defined to be `double` in C/C++ (or correspondingly `double precision` in Fortran) for versions of PETSc that have not been compiled for use with complex numbers. The `PetscScalar` data type enables identical code to be used when the PETSc libraries have been compiled for use with complex numbers. Numbers discusses the use of complex numbers in PETSc programs.
Matrices

Usage of PETSc matrices and vectors is similar. The user can create a new parallel or sequential matrix, $A$, which has $M$ global rows and $N$ global columns, with the routines

```c
MatCreate(MPI_Comm comm, Mat *A);
MatSetSizes(Mat A, PETSC_DECIDE, PETSC_DECIDE, PetscInt M, PetscInt N);
```

where the matrix format can be specified at runtime via the options database. The user could alternatively specify each processes' number of local rows and columns using $m$ and $n$.

```c
MatSetSizes(Mat A, PetscInt m, PetscInt n, PETSC_DETERMINE, PETSC_DETERMINE);
```

Generally one then sets the “type” of the matrix, with, for example,

```c
MatSetType(A, MATAIJ);
```

This causes the matrix $A$ to used the compressed sparse row storage format to store the matrix entries. See MatType for a list of all matrix types. Values can then be set with the command

```c
MatSetValues(Mat A, PetscInt *im, PetscInt *in, PetscScalar *values, INSERT_VALUES);
```

After all elements have been inserted into the matrix, it must be processed with the pair of commands

```c
MatAssemblyBegin(A, MAT_FINAL_ASSEMBLY);
MatAssemblyEnd(A, MAT_FINAL_ASSEMBLY);
```

Matrices discusses various matrix formats as well as the details of some basic matrix manipulation routines.

Linear Solvers

After creating the matrix and vectors that define a linear system, $Ax = b$, the user can then use KSP to solve the system with the following sequence of commands:

```c
KSPCreate(MPI_Comm comm, KSP *ksp);
KSPSetOperators(KSP ksp, Mat Amat, Mat Pmat);
KSPSetFromOptions(KSP ksp);
KSPSolve(KSP ksp, Vec b, Vec x);
KSPDestroy(KSP ksp);
```

The user first creates the KSP context and sets the operators associated with the system (matrix that defines the linear system, $Amat$ and matrix from which the preconditioner is constructed, $Pmat$). The user then sets various options for customized solution, solves the linear system, and finally destroys the KSP context. We emphasize the command KSPSetFromOptions(), which enables the user to customize the linear solution method at runtime by using the options database, which is discussed in Runtime Options. Through this database, the user not only can select an iterative method and preconditioner, but also can prescribe the convergence tolerance, set various monitoring routines, etc. (see, e.g., Profiling Programs).

KSP: Linear System Solvers describes in detail the KSP package, including the PC and KSP packages for preconditioners and Krylov subspace methods.

1.2. Getting Started
Nonlinear Solvers

Most PDE problems of interest are inherently nonlinear. PETSc provides an interface to tackle the nonlinear problems directly called SNES. SNES: Nonlinear Solvers describes the nonlinear solvers in detail. We highly recommend most PETSc users work directly with SNES, rather than using PETSc for the linear problem and writing their own nonlinear solver.

Error Checking

As noted above PETSc functions return a PetscErrorCode, which is an integer indicating whether an error has occurred during the call. Below, we indicate a traceback generated by error detection within a sample PETSc program. The error occurred on line 3618 of the file $PETSC_DIR/src/mat/impls/aij/seq/aij.c and was caused by trying to allocate too large an array in memory. The routine was called in the program ex3.c on line 66. See Error Checking for details regarding error checking when using the PETSc Fortran interface.

When running the debug version of the PETSc libraries, it does a great deal of checking for memory corruption (writing outside of array bounds etc). The macro CHKMEMQ can be called anywhere in the code to check the current status of the memory for corruption. By putting several (or many) of these macros into your code you can usually easily track down in what small segment of your code the corruption has occurred. One can also use Valgrind to track down memory errors; see the FAQ.

For complete error handling, calls to MPI functions should be made with PetscCallMPI(MPI Function(Args)). In the main Fortran program the calls should be PetscCallMPIA(MPI Function(Args)).
PETSc has a small number of C/C++ only macros that do not explicitly return error codes. These are used in the style

```c
XXXBegin(Args);
other code
XXXEnd();
```

and include `PetscOptionsBegin()`, `PetscOptionsEnd()`, `PetscObjectOptionsBegin()`, `PetscOptionsHeadBegin()`, `PetscOptionsHeadEnd()`, `PetscDrawCollectiveBegin()`, `PetscDrawCollectiveEnd()`, `MatPreallocateEnd()`, and `MatPreallocateBegin()`. These should not be checked for error codes. Another class of functions with the `Begin()` and `End()` paradigm including `PetscLogBegin()`, `PetscLogEnd()`, `MatAssemblyBegin()`, and `MatAssemblyEnd()` do return error codes that should be checked.

PETSc also has a set of C/C++ only macros that return an object, or `NULL` if an error has been detected. These include `PETSC_VIEWER_STDOUT_WORLD`, `PETSC_VIEWER_DRAW_WORLD`, `PETSC_VIEWER_STDOUT_(MPI_Comm)`, and `PETSC_VIEWER_DRAW_(MPI_Comm)`.

Finally `PetscObjectComm((PetscObject)x)` returns the communicator associated with the object `x` or `MPI_COMM_NULL` if an error was detected.

### 1.3 Parallel and GPU Programming

Numerical computing today has multiple levels of parallelism (concurrency).

- Low-level, single instruction multiple data (SIMD) parallelism or, somewhat similar, on-GPU parallelism,

- Medium-level, multiple instruction shared memory parallelism (thread parallelism), and

- High-level, distributed memory parallelism

Traditional CPUs support the lower two levels via, for example, Intel AVX-like instructions (CPU SIMD parallelism) and Unix threads, often managed by using OpenMP pragmas (CPU OpenMP parallelism), (or multiple processes). GPUs also support the lower two levels via kernel functions (GPU kernel parallelism) and streams (GPU stream parallelism). Distributed memory parallelism is created by combining multiple CPUs and/or GPUs and using MPI for communication (MPI Parallelism).

In addition there is also concurrency between computations (floating point operations) and data movement (from memory to caches and registers and via MPI between distinct memory nodes).

PETSc provides support for all these levels of parallelism but its strongest support is for MPI-based distributed memory parallelism.

#### 1.3.1 MPI Parallelism

Since PETSc uses the message-passing model for parallel programming and employs MPI for all interprocessor communication, the user is free to employ MPI routines as needed throughout an application code. However, by default the user is shielded from many of the details of message passing within PETSc, since these are hidden within parallel objects, such as vectors, matrices, and solvers. In addition, PETSc provides tools such as generalized vector scatters/gathers to assist in the management of parallel data.

Recall that the user must specify a communicator upon creation of any PETSc object (such as a vector, matrix, or solver) to indicate the processors over which the object is to be distributed. For example, as mentioned above, some commands for matrix, vector, and linear solver creation are:
MatCreate(MPI_Comm comm, Mat *A);
VecCreate(MPI_Comm comm, Vec *x);
KSPCreate(MPI_Comm comm, KSP *ksp);

The creation routines are collective on all processes in the communicator; thus, all processors in the communicator must call the creation routine. In addition, if a sequence of collective routines is being used, they must be called in the same order on each process.

The next example, given below, illustrates the solution of a linear system in parallel. This code, corresponding to KSP Tutorial ex2, handles the two-dimensional Laplacian discretized with finite differences, where the linear system is again solved with KSP. The code performs the same tasks as the sequential version within Simple PETSc Examples. Note that the user interface for initiating the program, creating vectors and matrices, and solving the linear system is exactly the same for the uniprocessor and multiprocessor examples. The primary difference between the examples in Simple PETSc Examples and here is that each processor forms only its local part of the matrix and vectors in the parallel case.

Listing: KSP Tutorial src/ksp/ksp/tutorials/ex2.c

```c
static char help[] = "Solves a linear system in parallel with KSP.\n```
Input parameters include:\n
- view_exact_sol : write exact solution vector to stdout\n
- m <mesh_x> : number of mesh points in x-direction\n
- n <mesh_y> : number of mesh points in y-direction\n
/*
  Include "petscksp.h" so that we can use KSP solvers.
*/
#include <petscksp.h>

int main(int argc, char **args)
{
  Vec x, b, u; /* approx solution, RHS, exact solution */
  Mat A; /* linear system matrix */
  KSP ksp; /* linear solver context */
  PetscReal norm; /* norm of solution error */
  PetscInt i, j, Ii, J, Istart, Iend, m = 8, n = 7, its;
  PetscBool flg;
  PetscScalar v;

  PetscFunctionBeginUser;
  PetscCall(PetscInitialize(&argc, &args, (char *)0, help));
  PetscCall(PetscOptionsGetInt(NULL, NULL, "-m", &m, NULL));
  PetscCall(PetscOptionsGetInt(NULL, NULL, "-n", &n, NULL));
  /* Compute the matrix and right-hand-side vector that define the linear system, Ax = b. */
  /* Create parallel matrix, specifying only its global dimensions. When using MatCreate(), the matrix format can be specified at runtime. Also, the parallel partitioning of the matrix is determined by PETSc at runtime. */
  Performance tuning note: For problems of substantial size, preallocation of matrix memory is crucial for attaining good performance. See the matrix chapter of the users manual for details.

(continues on next page)
PetscCall(MatCreate(PETSC_COMM_WORLD, &A));
PetscCall(MatSetSizes(A, PETSC_DECIDE, PETSC_DECIDE, m * n, m * n));
PetscCall(MatSetFromOptions(A));
PetscCall(MatMPIAIJSetPreallocation(A, 5, NULL, 5, NULL));
PetscCall(MatSeqAIJSetPreallocation(A, 5, NULL));
PetscCall(MatSeqSBAIJSetPreallocation(A, 1, 5, NULL));
PetscCall(MatMPISBAIJSetPreallocation(A, 1, 5, NULL, 5, NULL));
PetscCall(MatMPISELLSetPreallocation(A, 5, NULL, 5, NULL));
PetscCall(MatSeqSELLSetPreallocation(A, 5, NULL));

Currently, all PETSc parallel matrix formats are partitioned by contiguous chunks of rows across the processors. Determine which rows of the matrix are locally owned.

PetscCall(MatGetOwnershipRange(A, &Istart, &Iend));

Set matrix elements for the 2-D, five-point stencil in parallel.
- Each processor needs to insert only elements that it owns locally (but any non-local elements will be sent to the appropriate processor during matrix assembly).
- Always specify global rows and columns of matrix entries.

Note: this uses the less common natural ordering that orders first all the unknowns for $x = h$ then for $x = 2h$ etc; Hence you see $J = I + n$ instead of $J = I + m$ as you might expect. The more standard ordering would first do all variables for $y = h$, then $y = 2h$ etc.

for (Ii = Istart; Ii < Iend; Ii++) {
  v = -1.0;
  i = Ii / n;
  j = Ii - i * n;
  if (i > 0) {
    J = Ii - n;
    PetscCall(MatSetValues(A, 1, &Ii, 1, &J, &v, ADD_VALUES));
  }
  if (i < m - 1) {
    J = Ii + n;
    PetscCall(MatSetValues(A, 1, &Ii, 1, &J, &v, ADD_VALUES));
  }
  if (j > 0) {
    J = Ii - 1;
    PetscCall(MatSetValues(A, 1, &Ii, 1, &J, &v, ADD_VALUES));
  }
  if (j < n - 1) {
    J = Ii + 1;
    PetscCall(MatSetValues(A, 1, &Ii, 1, &J, &v, ADD_VALUES));
  }
  v = 4.0;
  PetscCall(MatSetValues(A, 1, &Ii, 1, &Ii, &v, ADD_VALUES));
}
Assemble matrix, using the 2-step process:
    MatAssemblyBegin(), MatAssemblyEnd()
Computations can be done while messages are in transition
    by placing code between these two statements.
*/
PetscCall(MatAssemblyBegin(A, MAT_FINAL_ASSEMBLY));
PetscCall(MatAssemblyEnd(A, MAT_FINAL_ASSEMBLY));

/* A is symmetric. Set symmetric flag to enable ICC/Cholesky preconditioner */
PetscCall(MatSetOption(A, MAT_SYMMETRIC, PETSC_TRUE));

/* Create parallel vectors.
   - We form 1 vector from scratch and then duplicate as needed.
   - When using VecCreate(), VecSetSizes and VecSetFromOptions()
     in this example, we specify only the
     vector's global dimension; the parallel partitioning is determined
     at runtime.
   - When solving a linear system, the vectors and matrices MUST
     be partitioned accordingly. PETSc automatically generates
     appropriately partitioned matrices and vectors when MatCreate() and
     VecCreate() are used with the same communicator.
   - The user can alternatively specify the local vector and matrix
     dimensions when more sophisticated partitioning is needed
     (replacing the PETSC_DECIDE argument in the VecSetSizes() statement
     below).
*/
PetscCall(VecCreate(PETSC_COMM_WORLD, &u));
PetscCall(VecSetSizes(u, PETSC_DECIDE, m * n));
PetscCall(VecSetFromOptions(u));
PetscCall(VecDuplicate(u, &b));
PetscCall(VecDuplicate(b, &x));

/* Set exact solution; then compute right-hand-side vector.
   By default we use an exact solution of a vector with all
   elements of 1.0;
*/
PetscCall(VecSet(u, 1.0));
PetscCall(MatMult(A, u, b));

/* View the exact solution vector if desired
*/
flg = PETSC_FALSE;
PetcCall(PetscOptionsGetBool(NULL, NULL, "-view exact sol", &flg, NULL));
if (flg) PetscCall(VecView(u, PETSC_VIEWER_STDOUT_WORLD));

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
Create the linear solver and set various options
   - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - */
PetcCall(KSPCreate(PETSC_COMM_WORLD, &ksp));

/* Set operators. Here the matrix that defines the linear system
   also serves as the preconditioning matrix.
PetscCall(KSPSetOperators(ksp, A, A));

/*
 Set linear solver defaults for this problem (optional).
 - By extracting the KSP and PC contexts from the KSP context,
   we can then directly call any KSP and PC routines to set
   various options.
 - The following two statements are optional; all of these
   parameters could alternatively be specified at runtime via
   KSPSetFromOptions(). All of these defaults can be
   overridden at runtime, as indicated below.
 */
PetscCall(KSPSetTolerances(ksp, 1.e-2 / ((m + 1) * (n + 1)), 1.e-50, PETSC_DEFAULT, PETSC_DEFAULT));

/*
 Set runtime options, e.g.,
 - ksp_type <type> -pc_type <type> -ksp_monitor -ksp_rtol <rtol>
 These options will override those specified above as long as
 KSPSetFromOptions() is called _after_ any other customization
 routines.
 */
PetscCall(KSPSetFromOptions(ksp));

/*---------------------------------------------
 Solve the linear system
 ---------------------------------------------*/
PetscCall(KSPSolve(ksp, b, x));

/*---------------------------------------------
 Check the solution and clean up
 ---------------------------------------------*/
PetscCall(VecAXPY(x, -1.0, u));
PetscCall(VecNorm(x, NORM_2, &norm));
PetscCall(KSPGetIterationNumber(ksp, &its));

/*
 Print convergence information. PetscPrintf() produces a single
 print statement from all processes that share a communicator.
 An alternative is PetscFPrintf(), which prints to a file.
 */
PetscCall(PetscPrintf(PETSC_COMM_WORLD, "Norm of error %g iterations %" PetscInt_FMT "\n", (double)norm, its));

/*
 Free work space. All PETSc objects should be destroyed when they
 are no longer needed.
 */
PetscCall(KSPDestroy(&ksp));
PetscCall(VecDestroy(&u));
PetscCall(VecDestroy(&x));
PetscCall(VecDestroy(&b));
PetscCall(MatDestroy(&A));
1.3.2 CPU SIMD parallelism

SIMD parallelism occurs most commonly in the Intel advanced vector extensions (AVX) Wikipedia [https://en.wikipedia.org/wiki/Advanced_Vector_Extensions] families of instructions. It may be automatically used by the optimizing compiler, or in low-level libraries that PETSc uses such as BLAS (see BLIS [https://github.com/flame/blis] or rarely, directly in PETSc C/C++ code, as in MatMult_SeqSELL [https://petsc.org/main/src/mat/impls/sell/seq/sell.c.html#MatMult_SeqSELL].

1.3.3 CPU OpenMP parallelism

OpenMP parallelism is thread parallelism. Multiple threads (independent streams of instructions) process data and perform computations on different parts of memory that is shared (accessible) to all of the threads. The OpenMP model is most-often based on inserting pragmas into code indicating that a series of instructions (often within a loop) can be run in parallel. This is also called a fork-join model of parallelism, since much of the code remains sequential and only the computationally expensive parts in the ‘parallel region’ are parallel. OpenMP thus makes it relatively easy to add some degree of parallelism to a conventional sequential code in a shared memory environment.

POSIX threads (pthreads) is a library that may be called from C/C++. The library contains routines to create, join, and remove threads plus manage communications and synchronizations between threads. Pthreads is rarely used directly in numerical libraries and applications. Sometimes OpenMP is implemented on top of pthreads.

If one adds OpenMP parallelism to an MPI code one must make sure not to over-subscribe the hardware resources. For example, if MPI already has one rank per hardware core then using four OpenMP threads per MPI rank will slow the code down since now one core will need to switch back and forth between four OpenMP threads. There are limited practical advantages to a combined MPI and OpenMP model in PETSc, but it is possible.

For application codes that uses certain external packages including BLAS/LAPACK, SuperLU_DIST, MUMPS, MKL, and SuiteSparse one can build PETSc and these packages to take advantage of OpenMP by using the configure option -with-openmp. The number of OpenMP threads used in the application can be controlled with the PETSc command line option -omp_num_threads <num> or the environmental variable OMP_NUM_THREADS. Running a PETSc program with -omp_view will display the number of threads being used. The default number is often absurdly high for the given hardware so we recommend always setting it appropriately. Users can also put OpenMP pragmas into their own code. However since standard PETSc is not thread-safe, they should not, in general, call PETSc routines from inside the parallel regions.

PETSc’s MPI based linear solvers may be accessed from a sequential or non-MPI OpenMP program, see Using PETSc’s MPI parallel linear solvers from a non-MPI program.
There is an OpenMP thread-safe subset of PETSc that may be configured for using --with-threadsafety (often used along with --with-openmp or --download-concurrencykit). KSP Tutorial ex61f demonstrates how this may be used with OpenMP. In this mode one may have individual OpenMP threads that each manage their own (sequential) PETSc objects (each thread can interact only with its own objects). This is useful when one has many small systems (or sets of ODEs) that must be integrated in an “embarrassingly parallel” fashion on multicore systems.

See also:

1.3.4 GPU kernel parallelism

GPUs offer at least two levels of clearly defined parallelism. Kernel level parallelism is much like SIMD parallelism applied to loops; many different “iterations” of the loop index run on different hardware but in “lock-step” at the same time. PETSc utilizes this parallelism with three similar, but slightly different models:

- CUDA, which is provided by NVIDIA and runs on NVIDIA GPUs
- HIP, provided by AMD, which can, in theory, run on both AMD and NVIDIA GPUs
- and Kokkos, an open-source package that provides a slightly higher level programming model to utilize GPU kernels.

To utilize this one configures PETSc with either --with-cuda or --with-hip and, if they plan to use Kokkos, also --with-kokkos --with-kokkos-kernels.

In the GPU programming model that PETSc uses the GPU memory is distinct from the CPU memory. This means that data that resides on the CPU memory must be copied to the GPU (often this copy is done automatically by the libraries and the user does not need to manage it) if one wishes to use the GPU computational power on it. This memory copy is slow compared to the GPU speed hence it is crucial to minimize these copies. This often translates to trying to do almost all the computation on the GPU and not constantly switching between computations on the CPU and the GPU on the same data.

PETSc utilizes GPUs by providing vector and matrix classes (Vec and Mat) that are specifically written to run fast on the GPU. However, since it is difficult to write an entire PETSc code that runs only on the GPU one can also access and work with (for example, put entries into) the vectors and matrices on the CPU. The vector classes are VECCUDA, MATAIJCUSPARSE, VECKOKKOS, MATAIJKOKKOS, and VECHIP (matrices are not yet supported from PETSc with HIP).

More details on using GPUs from PETSc will follow in this document.

1.3.5 GPU stream parallelism

Incomplete
1.4 Compiling and Running Programs

The output below illustrates compiling and running a PETSc program using MPICH on a macOS laptop. Note that different machines will have compilation commands as determined by the configuration process. See Writing C/C++ or Fortran Applications for a discussion about how to compile your PETSc programs. Users who are experiencing difficulties linking PETSc programs should refer to the FAQ.

```
$ cd $PETSC_DIR/src/ksp/ksp/tutorials
$ make ex2
/Users/patrick/petsc/arch-darwin-double-debug/bin/mpicc -o ex2.o -c -g3 -I/Users/.../include -I/Users/patrick/petsc/arch-darwin-double-debug/include -I/.../include -I/opt/local/include -I`pwd`/ex2.c
/Users/patrick/petsc/arch-darwin-double-debug/bin/mpicc -g3 -o ex2 ex2.o -Wl,-rpath,..
/Users/patrick/petsc/arch-darwin-double-debug/lib -L/Users/patrick/petsc/arch...
_ext.10.5 -Iquadmath -lm -lcblas -lgfortran -lgcc_
/bin/rm -f ex2.o
$ $PETSC_DIR/lib/petsc/bin/petscmpiexec -n 1 ./ex2
Norm of error 0.000156044 iterations 6
$ $PETSC_DIR/lib/petsc/bin/petscmpiexec -n 2 ./ex2
Norm of error 0.000411674 iterations 7
```

1.5 Profiling Programs

The option -log_view activates printing of a performance summary, including times, floating point operation (flop) rates, and message-passing activity. Profiling provides details about profiling, including interpretation of the output data below. This particular example involves the solution of a linear system on one processor using GMRES and ILU. The low floating point operation (flop) rates in this example are due to the fact that the code solved a tiny system. We include this example merely to demonstrate the ease of extracting performance information.

```
$ $PETSC_DIR/lib/petsc/bin/petscmpiexec -n 1 ./ex1 -n 1000 -pc_type ilu -ksp_type gmres -ksp_rtol 1.e-7 -log_view
...
--------------------------------------------------------------------------------------
<table>
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<tr>
<th>Event</th>
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<th>Time (sec)</th>
<th>Flops</th>
<th>Max Ratio</th>
<th>Max Ratio</th>
<th>Max Ratio</th>
<th>Mess</th>
<th>AvgLen</th>
<th>Reduct</th>
<th>%T</th>
</tr>
</thead>
<tbody>
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<td></td>
<td></td>
</tr>
</tbody>
</table>
--------------------------------------------------------------------------------------
| VecMDot                      | 1 1.0 | 3.2830e-06 | 1.0   | 2.00e+03  | 1.0       | 0.0e+00   | 0.0e+00| 0.0e+00| 0.000156044 | 6 |
| -5                            | 0 0   | 0 5 0 0    | 0 0   | 609       |           |           |      |        |        |    |
| VecNorm                       | 3 1.0 | 4.4550e-06 | 1.0   | 6.00e+03  | 1.0       | 0.0e+00   | 0.0e+00| 0.0e+00| 0.000411674 | 7 |
| -14                           | 0 0   | 0 0 0 14   | 0 0   | 1346      |           |           |      |        |        |    |
| VecScale                      | 2 1.0 | 4.0110e-06 | 1.0   | 2.00e+03  | 1.0       | 0.0e+00   | 0.0e+00| 0.0e+00| 0.000156044 | 6 |
| -5                            | 0 0   | 0 5 0 0    | 0 0   | 499       |           |           |      |        |        |    |
| VecCopy                       | 1 1.0 | 3.2280e-06 | 1.0   | 2.00e+03  | 1.0       | 0.0e+00   | 0.0e+00| 0.0e+00| 0.000156044 | 6 |
| -0                            | 0 0   | 0 0 0 2    | 0 0   | 0         |           |           |      |        |        |    |
| VecSet                        | 11 1.0| 2.5537e-05 | 1.0   | 0.00e+00  | 0.0       | 0.0e+00   | 0.0e+00| 0.0e+00| 0.000156044 | 6 |
| -0                            | 0 0   | 0 2 0 0    | 0 0   | 0         |           |           |      |        |        |    |
| VecAXPY                       | 2 1.0 | 2.0930e-06 | 1.0   | 4.00e+03  | 1.0       | 0.0e+00   | 0.0e+00| 0.0e+00| 0.000411674 | 7 |
(continues on next page)
Memory usage is given in bytes:

<table>
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<tr>
<th>Object Type</th>
<th>Creations</th>
<th>Detections</th>
<th>Memory</th>
<th>Descendants' Mem.</th>
</tr>
</thead>
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</tr>
<tr>
<td>Matrix</td>
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<td>2</td>
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<td>0.</td>
</tr>
<tr>
<td>Krylov Solver</td>
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<td>1</td>
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<td>0.</td>
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<tr>
<td>Preconditioner</td>
<td>1</td>
<td>1</td>
<td>1032</td>
<td>0.</td>
</tr>
<tr>
<td>Index Set</td>
<td>3</td>
<td>3</td>
<td>10328</td>
<td>0.</td>
</tr>
<tr>
<td>Viewer</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0.</td>
</tr>
</tbody>
</table>

1.5. Profiling Programs
1.6 Writing C/C++ or Fortran Applications

The examples throughout the library demonstrate the software usage and can serve as templates for developing custom applications. We suggest that new PETSc users examine programs in the directories $PETSC_DIR/src/<library>/tutorials where <library> denotes any of the PETSc libraries (listed in the following section), such as SNES or KSP or TS. The manual pages located at https://petsc.org/release/documentation/ provide links (organized by both routine names and concepts) to the tutorial examples.

To develop an application program that uses PETSc, we suggest the following:

- Download and install PETSc.
- For completely new applications
  1. Make a directory for your source code: for example, mkdir $HOME/application
  2. Change to that directory; for example, cd $HOME/application
  3. Copy an example in the directory that corresponds to the problems of interest into your directory, for example, cp $PETSC_DIR/src/snes/tutorials/ex19.c ex19.c
  4. Select an application build process. The PETSC_DIR (and PETSC_ARCH if the --prefix=directoryname option was not used when configuring PETSc) environmental variable(s) must be set for any of these approaches.
     - make (recommended). Copy $PETSC_DIR/share/petsc/Makefile.user or $PETSC_DIR/share/petsc/Makefile.basic.user to your directory, for example, cp $PETSC_DIR/share/petsc/Makefile.user makefile
       Examine the comments in your makefile
       Makefile.user uses the pkg-config tool and is the recommended approach.
       Use make ex19 to compile your program
     - CMake. Copy $PETSC_DIR/share/petsc/CMakeLists.txt to your directory, for example, cp $PETSC_DIR/share/petsc/CMakeLists.txt CMakeLists.txt
       Edit CMakeLists.txt, read the comments on usage and change the name of application from ex1 to your application executable name.
  5. Run the program, for example, ./ex19
  6. Start to modify the program for developing your application.
- For adding PETSc to an existing application
  1. Start with a working version of your code that you build and run to confirm that it works.
  2. Upgrade your build process. The PETSC_DIR (and PETSC_ARCH if the --prefix=directoryname option was not used when configuring PETSc) environmental variable(s) must be set for any of these approaches.
     - Using make. Update the application makefile to add the appropriate PETSc include directories and libraries.
       * Recommended approach. Examine the comments in $PETSC_DIR/share/petsc/Makefile.user and transfer selected portions of that file to your makefile.
       * Minimalist. Add the line
include ${PETSC_DIR}/lib/petsc/conf/variables
to the bottom of your makefile. This will provide a set of PETSc specific make variables you may use in your makefile. See the comments in the file $PETSC_DIR/share/petsc/Makefile.basic.user for details on the usage.

* Simple, but hands the build process over to PETSc’s control. Add the lines

```c
include ${PETSC_DIR}/lib/petsc/conf/variables
include ${PETSC_DIR}/lib/petsc/conf/rules
```
to the bottom of your makefile. See the comments in the file $PETSC_DIR/share/petsc/Makefile.basic.user for details on the usage. Since PETSc’s rules now control the build process you will likely need to simplify and remove much of your makefile.

* Not recommended since you must change your makefile for each new configuration/computing system. This approach does not require that the environmental variable PETSC_DIR be set when building your application since the information will be hard-wired in your makefile. Run the following command in the PETSc root directory to get the information needed by your makefile:

```bash
$ make getlinklibs getincludedirs getcflags getcxxflags
        getfortranflags getccompiler getfortrancompiler getcxxcompiler
```

All the libraries listed need to be linked into your executable and the include directories and flags need to be passed to the compiler(s). Usually this is done by setting LD-FLAGS=<list of library flags and libraries> and CFLAGS=<list of -I and other flags> and FFLAGS=<list of -I and other flags> etc in your makefile.

– Using CMake. Update the application CMakeLists.txt by examining the code and comments in $PETSC_DIR/share/petsc/CMakeLists.txt

3. Rebuild your application and ensure it still runs correctly.

4. Add a PetscInitialize() near the beginning of your code and PetscFinalize() near the end with appropriate include commands (and use commands in Fortran)

5. Rebuild your application and ensure it still runs correctly.

6. Slowly start utilizing PETSc functionality in your code, ensure that your code continues to build and run correctly.

### 1.7 PETSc’s Object-Oriented Design

Though PETSc has a large API, conceptually it’s rather simple. There are three abstract basic data objects (classes): index sets, `IS`, vectors, `Vec`, and matrices, `Mat`. Plus a larger number of abstract algorithm objects (classes) starting with: preconditioners, `PC`, Krylov solvers, `KSP`, and so forth.

Let `Object` represent any of these objects. Objects are created with

```c
Object obj;
ObjectCreate(MPI_Comm, &obj);
```
The object is empty and little can be done with it. A particular implementation of the class is associated with the object by setting the object’s “type”, where type is merely a string name of an implementation class using

```
Object obj;
ObjectSetType(obj,"Name");
```

Some objects support subclasses which are specializations of the type. These are set with

```
Object obj;
ObjectNameSetType(obj,"SubName");
```

For example, within TS one may do

```
TS obj;
TSCreate(PETSC_COMM_WORLD,&obj);
TSSetType(obj,TSARKIMEX);
TSARKIMEXSetType(obj,TSARKIMEX3);
```

The abstract class TS can embody any ODE/DAE integrator scheme. This example creates an additive Runge-Kutta ODE/DAE IMEX integrator, whose type name is TSARKIMEX, using a 3-order scheme with an L-stable implicit part, whose subtype name is TSARKIMEX3.

In order to allow PETSc objects to be runtime configurable, PETSc objects provide a universal way of selecting types (classes) and subtypes at runtime, from what is referred to as the “options database”. The code above can be replaced with

```
TS obj;
TSCreate(PETSC_COMM_WORLD,&obj);
TSSetFromOptions(obj);
```

now both the type and subtype can be conveniently set from the command line

```
$ ./app -ts_type arkimex -ts_arkimex_type 3
```

The object’s type (implementation class) or subclass can also be changed at any time simply by calling TSSetType() again (though in order to override command line options the call to TSSetType() must be made _after_ TSSetFromOptions()). For example:

```
// (if set) command line options "override" TSSetType()
TSSetType(ts, TSGLLE);
TSSetFromOptions(ts);

// TSSetType() overrides command line options
TSSetFromOptions(ts);
TSSetType(ts, TSGLLE);
```

Since the later call always overrides the earlier call the second form shown is rarely – if ever – used, as it is less flexible than configuring command line settings.

The standard methods on an object are of the general form

```
ObjectSetXXX(obj,...);
ObjectGetXXX(obj,...);
ObjectYYY(obj,...);
```

For example
TSSetRHSFunction(obj,...)

Particular types and subtypes of objects may have their own methods, which are given in the form

ObjectNameSetXXX(obj,...);
ObjectNameGetXXX(obj,...);
ObjectNameYYY(obj,...);

and

ObjectNameSubNameSetXXX(obj,...);
ObjectNameSubNameGetXXX(obj,...);
ObjectNameSubNameYYY(obj,...);

where Name and SubName are the type and subtype names (for example, as above TSARKIMEX and 3. Most “set” operations have options database versions with the same names in lower case, separated by underscores and with the set remove. For example,

KSPGMRESSetRestart(obj,30);  // ignored if the type is not KSPGMRES

can be set at the command line with

$ ./app -ksp_gmres_restart 30

There are a special subset of type-specific methods that are ignored if the type does not match the function name. These are usually setter functions that control some aspect specific to the subtype. For example,

KSPGMRESSetRestart(obj,30);  // ignored if the type is not KSPGMRES

These allow cleaner code since it does not have a multitude of if statements to avoid inactive methods. That is one does not need to write code like

if (type == KSPGMRES) {
    KSPGMRESSetRestart(obj,30);
}

There are many “get” routines that give one temporary access to the internal data of an object. They are used in the style

XXX xxx;
ObjectNameGetXXX(obj,&xxx);
// use xxx
ObjectNameRestoreXXX(obj,&xxx);

Objects obtained with a “get” routine should be returned with a “restore” routine, generally within the same function. Objects obtained with a “create” routine should be freed with a “destroy” routine.

There may be variants of the “get” routines that give more limited access to the obtained object. For example,

const PetscScalar *x;
// specialized variant of VecGetArray()
VecGetArrayRead(vec,&x);
// one can read but not write with x[]
PetscReal y = 2*x[0];
Objects can be displayed (in a large number of ways) with

\[
\text{ObjectView(obj,PetscViewer viewer);} \\
\text{ObjectViewFromOptions(obj,...);}
\]

Where \text{PetscViewer} is an abstract object that can represent standard output, an ASCII or binary file, a graphical window, etc. The second variant allows the user to delay until runtime the decision of what viewer and format to use to view the object or if to view the object at all.

Objects are destroyed with

\[
\text{ObjectDestroy(&obj)}
\]

### 1.7.1 User Callbacks

In many situations the user may also wish to override or provide custom functionality. This is handled via callbacks which the library will call at the appropriate time. The most general callback is provided by

\[
\text{PetscObjecSetCallback(obj,callbackfunction(), void *ctx, callbackdestroy(void *ctx));}
\]

where \text{callbackfunction()} is what is used by the library, \text{ctx} is an optional data-structure (array, struct, PETSc object) that is used by \text{callbackfunction()} and \text{callbackdestroy(void *ctx)} is an optional function that will be called when \text{obj} is destroyed. The use of the \text{callbackdestroy()} allows users to “set and forget” data structures that will not be needed elsewhere but still need to be cleaned up when no longer needed. Here is an example of the use of a full-fledged callback

\[
\text{TS ts;}
\text{TSMonitorLGCtx *ctx;}
\text{TSMonitorLGCtxCreate(..., &ctx)}
\text{TSMonitorSet(ts, TSMonitorLGTimeStep, ctx, (PetscErrorCode(*)(void *, ...))TSMonitorLGCtxDestroy);}
\text{TSSolve(ts);}
\]

Occasionally routines to set callback functions take additional data objects that will be used by the object but are not context data for the function. For example,

\[
\text{SNES obj;}
\text{Vec r;}
\text{void *ctx;}
\text{SNESSetFunction(snes, r, UserApplyFunction(SNES,Vec,Vec,void *ctx), ctx);}
\]

The \text{r} vector is an optional argument provided by the user which will be used as work-space by \text{SNES}. Note that this callback does not provide a way for the user to have the \text{ctx} destroyed when the \text{SNES} object is destroyed, the users must ensure that they free it at an appropriate time. There is no logic to the various ways PETSc accepts callback functions in different places in the code.

See \text{Tao use of PETSc and callbacks} for a cartoon on the use of callbacks in \text{Tao}.
Fig. 1.2: Sample lifetime of a PETSc object
1.8 Directory Structure

We conclude this introduction with an overview of the organization of the PETSc software. The root directory of PETSc contains the following directories:

- **doc** (only in the tarball distribution of PETSc; not the git repository) - All documentation for PETSc. The files `manual.pdf` contains the hyperlinked users manual, suitable for printing or on-screen viewing. Includes the subdirectory `manualpages` (on-line manual pages).

- **lib/petsc/conf** - Base PETSc configuration files that define the standard make variables and rules used by PETSc.

- **include** - All include files for PETSc that are visible to the user.

- **include/petsc/private** - Private PETSc include files that should not need to be used by application programmers.

- **share** - Some small test matrices in data files

- **src** - The source code for all PETSc libraries, which currently includes
  - **vec** - vectors,
    - *is* - index sets,
  - **mat** - matrices,
  - **ksp** - complete linear equations solvers,
    - *ksp* - Krylov subspace accelerators,
    - *pc* - preconditioners,
  - **snes** - nonlinear solvers
  - **ts** - ODE solvers and timestepping,
  - **dm** - data management between meshes and solvers, vectors, and matrices,
  - **sys** - general system-related routines,
    - *logging* - PETSc logging and profiling routines,
    - *classes* - low-level classes
      - **draw** - simple graphics,
      - **viewer** - mechanism for printing and visualizing PETSc objects,
      - **bag** - mechanism for saving and loading from disk user data stored in C structs.
      - **random** - random number generators.

Each PETSc source code library directory has the following subdirectories:

- **tutorials** - Programs designed to teach users about PETSc. These codes can serve as templates for the design of custom applications.

- **tests** - Programs designed for thorough testing of PETSc. As such, these codes are not intended for examination by users.

- **interface** - Provides the abstract base classes for the objects. Code here does not know about particular implementations and does not actually perform operations on the underlying numerical data.
• **impls** - Source code for one or more implementations of the class for particular data structures or algorithms.

• **utils** - Utility routines. Source here may know about the implementations, but ideally will not know about implementations for other components.
2.1 Vectors and Parallel Data

Vectors (denoted by Vec) are used to store discrete PDE solutions, right-hand sides for linear systems, etc. Users can create and manipulate entries in vectors directly with a basic, low-level interface or they can use the PETSc DM objects to connect actions on vectors to the type of discretization and grid that they are working with. These higher level interfaces handle much of the details of the interactions with vectors and hence are preferred in most situations. This chapter is organized as follows:

- **Creating Vectors**
  - User managed
  - DMDA - Creating vectors for structured grids
  - DMStag - Creating vectors for staggered grids
  - DMPLEX - Creating vectors for unstructured grids
  - DMNETWORK - Creating vectors for networks

- Setting vector values
  - For generic vectors
  - DMDA - Setting vector values
  - DMSTAG - Setting vector values
  - DMPLEX - Setting vector values
  - DMNETWORK - Setting vector values

- Basic Vector Operations
- Local/global vectors and communicating between vectors
- Low-level Vector Communication
  - Local to global mappings
  - Global Vectors with locations for ghost values
- Application Orderings
2.1.1 Creating Vectors

PETSc provides many ways to create vectors. The most basic, where the user is responsible for managing the parallel distribution of the vector entries, and a variety of higher-level approaches, based on DM, for classes of problems such as structured grids, staggered grids, unstructured grids, networks, and particles.

The most basic way to create a vector with a local size of \( m \) and a global size of \( M \), is using the commands

```c
VecCreate(MPI_Comm comm, Vec *v);
VecSetSizes(Vec v, PetscInt m, PetscInt M);
VecSetFromOptions(Vec v);
```

which automatically generates the appropriate vector type (sequential or parallel) over all processes in \( \text{comm} \).

The option `-vec_type <type>` can be used in conjunction with `VecSetFromOptions()` to specify the use of a particular type of vector. For example, for NVIDIA GPU CUDA use `cuda`. The GPU based vectors allow one to set values on either the CPU or GPU but do their computations on the GPU.

We emphasize that all processes in \( \text{comm} \) must call the vector creation routines, since these routines are collective on all processes in the communicator. If you are not familiar with MPI communicators, see the discussion in *Writing PETSc Programs* on page . In addition, if a sequence of creation routines is used, they must be called in the same order on each process in the communicator.

Instead of, or before calling `VecSetFromOptions()`, one can call

```c
VecSetType(Vec v, VecType <VECCUDA, VECHIP, VECKOKKOS etc>)
```

One can create vectors whose entries are stored on GPUs using, the short-hand helper routine,

```c
VecCreateMPICUDA(MPI_Comm comm, PetscInt m, PetscInt M, Vec *x);
```

There are short hand creation routines for almost all vector types; we recommend using the more verbose form because it allows selecting CPU or GPU simulations at runtime.

For applications running in parallel that involve multi-dimensional structured grids, unstructured grids, networks, etc it is cumbersome and complicated for users to explicitly manage the needed local and global sizes of the vectors. Hence PETSc provides a powerful abstract object called the DM to help manage the vectors and matrices needed for such applications. Parallel vectors can be created easily with

```c
DMCreateGlobalVector(DM dm, Vec *v)
```

The DM object, see DMDA - *Creating vectors for structured grids*, DMStag - *Creating vectors for staggered grids*, and DMPlex: *Unstructured Grids* for more details on DM for structured grids, staggered structured grids and for unstructured grids, manages creating the correctly sized parallel vectors efficiently. One controls the type of vector that DM creates by calling

```c
DMSetVecType(DM dm, VecType vt)
```

or by calling `DMSetFromOptions(DM dm)` and using the option `-dm_vec_type <standard or cuda or kokkos etc>`
DMDA - Creating vectors for structured grids

Each DM type is suitable for a family of problems. The first of these DMDA are intended for use with logically structured rectangular grids when communication of nonlocal data is needed before certain local computations can occur. PETSc distributed arrays are designed only for the case in which data can be thought of as being stored in a standard multidimensional array; thus, DMDAs are not intended for parallelizing unstructured grid problems, etc.

For example, a typical situation one encounters in solving PDEs in parallel is that, to evaluate a local function, \( f(x) \), each process requires its local portion of the vector \( x \) as well as its ghost points (the bordering portions of the vector that are owned by neighboring processes). Figure Ghost Points for Two Stencil Types on the Seventh Process illustrates the ghost points for the seventh process of a two-dimensional, structured parallel grid. Each box represents a process; the ghost points for the seventh process’s local part of a parallel array are shown in gray.

![Ghost Points for Two Stencil Types on the Seventh Process](image)

The DMDA object contains parallel data layout information and communication information and is used to create vectors and matrices with the proper layout.

One creates a distributed array communication data structure in two dimensions with the command

```c
DMDACreate2d(MPI_Comm comm, DMBoundaryType xperiod, DMBoundaryType yperiod,
     DMDAStencilType st, PetscInt M, PetscInt N, PetscInt m, PetscInt n, PetscInt dof,
     PetscInt s, PetscInt *lx, PetscInt *ly, DM *da);
```

The arguments \( M \) and \( N \) indicate the global numbers of grid points in each direction, while \( m \) and \( n \) denote the process partition in each direction; \( m \times n \) must equal the number of processes in the MPI communicator, \( comm \). Instead of specifying the process layout, one may use PETSC_DECIDE for \( m \) and \( n \) so that PETSc will select the partition. The type of periodicity of the array is specified by \( xperiod \) and \( yperiod \), which can be DM_BOUNDARY_NONE (no periodicity), DM_BOUNDARY_PERIODIC (periodic in that direction), DM_BOUNDARY_TWIST (periodic in that direction, but identified in reverse order), DM_BOUNDARY_GHOSTED, or DM_BOUNDARY_MIRROR. The argument \( dof \) indicates the number of degrees of freedom at each array point, and \( s \) is the stencil width (i.e., the width of the ghost point region). The optional arrays \( \text{lx} \) and \( \text{ly} \) may contain the number of nodes along the x and y axis for each cell, i.e. the dimension of \( \text{lx} \) is \( m \) and the dimension of \( \text{ly} \) is \( n \); alternately, NULL may be passed in.

Two types of distributed array communication data structures can be created, as specified by \( st \). Star-type stencils that radiate outward only in the coordinate directions are indicated by DMDA_STENCIL_STAR, while box-type stencils are specified by DMDA_STENCIL_BOX. For example, for the two-dimensional case, DMDA_STENCIL_STAR with width 1 corresponds to the standard 5-point stencil,
while **DMDA_STENCIL_BOX** with width 1 denotes the standard 9-point stencil. In both instances the ghost points are identical, the only difference being that with star-type stencils certain ghost points are ignored, decreasing substantially the number of messages sent. Note that the **DMDA_STENCIL_STAR** stencils can save interprocess communication in two and three dimensions.

These **DMDA** stencils have nothing directly to do with a specific finite difference stencil one might chose to use for a discretization; they only ensure that the correct values are in place for application of a user-defined finite difference stencil (or any other discretization technique).

The commands for creating distributed array communication data structures in one and three dimensions are analogous:

```
DMDACreate1d(MPI_Comm comm, DMBoundaryType xperiod, PetscInt M, PetscInt w, PetscInt s, ...PetscInt *lc, DM *inra);
DMDACreate3d(MPI_Comm comm, DMBoundaryType xperiod, DMBoundaryType yperiod, ...DMBoundaryType zperiod, DMDAStencilType stencil_type, PetscInt M, PetscInt N, PetscInt w, ...PetscInt m, PetscInt n, PetscInt p, PetscInt w, PetscInt s, PetscInt *lx, PetscInt *ly, ...PetscInt *lz, DM *inra);
```

The routines to create distributed arrays are collective, so that all processes in the communicator **comm** must call the same creation routines in the same order.

**DMStag - Creating vectors for staggered grids**

For structured grids with staggered data (living on elements, faces, edges, and/or vertices), the **DMSTAG** object is available. It behaves much like **DMDA**. See **DMSTAG: Staggered, Structured Grid** for discussion of creating vectors with **DMSTAG**.

**DMPLEX - Creating vectors for unstructured grids**

See **DMplex: Unstructured Grids** for discussion of creating vectors with **DMPLEX**.

**DMNETWORK - Creating vectors for networks**

See **Networks** for discussion of creating vectors with **DMNETWORK**.

One can examine (print out) a vector with the command

```
VecView(Vec x, PetscViewer v);
```

To print the vector to the screen, one can use the viewer **PETSC_VIEWER_STDOUT_WORLD**, which ensures that parallel vectors are printed correctly to **stdout**. To display the vector in an X-window, one can use the default X-windows viewer **PETSC_VIEWER_DRAW_WORLD**, or one can create a viewer with the routine **PetscViewerDrawOpenX()**. A variety of viewers are discussed further in **Viewers: Looking at PETSc Objects**.

To create a new vector of the same format as an existing vector, one uses

```
VecDuplicate(Vec old, Vec *new);
```

To create several new vectors of the same format as an existing vector, one uses

```
VecDuplicateVecs(Vec old, PetscInt n, Vec **new);
```
This routine creates an array of pointers to vectors. The two routines are very useful because they allow one to write library code that does not depend on the particular format of the vectors being used. Instead, the subroutines can automatically correctly create work vectors based on the specified existing vector. As discussed in Duplicating Multiple Vectors, the Fortran interface for VecDuplicateVecs() differs slightly.

When a vector is no longer needed, it should be destroyed with the command

```
VecDestroy(Vec *x);
```

To destroy an array of vectors, use the command

```
VecDestroyVecs(PetscInt n, Vec **vecs);
```

Note that the Fortran interface for VecDestroyVecs() differs slightly, as described in Duplicating Multiple Vectors.

It is also possible to create vectors that use an array provided by the user, rather than having PETSc internally allocate the array space. Such vectors can be created with the routines such as

```
VecCreateSeqWithArray(PETSC_COMM_SELF, PetscInt bs, PetscInt n, PetscScalar *array, Vec *V);
VecCreateMPIWithArray(MPI_Comm comm, PetscInt bs, PetscInt n, PetscInt N, PetscScalar *array, Vec *vv);
VecCreateMPICUDAWithArray(MPI_Comm comm, PetscInt bs, PetscInt n, PetscInt N, PetscScalar *array, Vec *vv);
```

For GPU vectors the array pointer should be a GPU memory location.

Note that here one must provide the value `n`; it cannot be PETSC_DECIDE and the user is responsible for providing enough space in the array; `n*sizeof(PetscScalar)`.

### 2.1.2 Assembling (putting values in) vectors

One can assign a single value to all components of a vector with the command

```
VecSet(Vec x, PetscScalar value);
```

Assigning values to individual components of the vector is more complicated, in order to make it possible to write efficient parallel code. Assigning a set of components on a CPU is a two-step process: one first calls

```
VecSetValues(Vec x, PetscInt n, PetscInt *indices, PetscScalar *values, INSERT_VALUES);
```

any number of times on any or all of the processes. The argument `n` gives the number of components being set in this insertion. The integer array `indices` contains the global component indices, and `values` is the array of values to be inserted. Any process can set any components of the vector; PETSc ensures that they are automatically stored in the correct location. Once all of the values have been inserted with VecSetValues(), one must call

```
VecAssemblyBegin(Vec x);
```

followed by

```
VecAssemblyEnd(Vec x);
```

to perform any needed message passing of nonlocal components. In order to allow the overlap of communication and calculation, the user’s code can perform any series of other actions between these two calls while the messages are in transition.
Example usage of `VecSetValues()` may be found in `src/vec/vec/tutorials/ex2.c` or `src/vec/vec/tutorials/ex2f.F90`.

Often, rather than inserting elements in a vector, one may wish to add values. This process is also done with the command

```c
VecSetValues(Vec x, PetscInt n, PetscInt *indices, PetscScalar *values, ADD_VALUES);
```

Again one must call the assembly routines `VecAssemblyBegin()` and `VecAssemblyEnd()` after all of the values have been added. Note that addition and insertion calls to `VecSetValues()` cannot be mixed. Instead, one must add and insert vector elements in phases, with intervening calls to the assembly routines. This phased assembly procedure overcomes the nondeterministic behavior that would occur if two different processes generated values for the same location, with one process adding while the other is inserting its value. (In this case the addition and insertion actions could be performed in either order, thus resulting in different values at the particular location. Since PETSc does not allow the simultaneous use of `INSERT_VALUES` and `ADD_VALUES` this nondeterministic behavior will not occur in PETSc.)

You can call `VecGetValues()` to pull local values from a vector (but not off-process values).

For vectors obtained with `DMCreateGlobalVector()` on can use `VecSetValuesLocal()` to set values into a global vector but using the local (ghosted) vector indexing of the vector entries.

It is also possible to interact directly with the arrays that the vector values are stored in. The routine `VecGetArray()` returns a pointer to the elements local to the process:

```c
VecGetArray(Vec v, PetscScalar **array);
```

When access to the array is no longer needed, the user should call

```c
VecRestoreArray(Vec v, PetscScalar **array);
```

If the values do not need to be modified, the routines

```c
VecGetArrayRead(Vec v, const PetscScalar **array);
VecRestoreArrayRead(Vec v, const PetscScalar **array);
```

should be used instead.

Listing: SNES Tutorial `src/snes/tutorials/ex1.c`

```c
PetscErrorCode FormFunction1(SNES snes, Vec x, Vec f, void *ctx)
{
    const PetscScalar *xx;
    PetscScalar      *ff;

    PetscFunctionBeginUser;
    /*
    | Get pointers to vector data.
    | - For default PETSc vectors, VecGetArray() returns a pointer to
    |   the data array. Otherwise, the routine is implementation dependent.
    | - You MUST call VecRestoreArray() when you no longer need access to
    |   the array.
    */
    PetscCall(VecGetArrayRead(x, &xx));
    PetscCall(VecGetArray(f, &ff));

    /* Compute function */
    (continues on next page)
```

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ff[0] = xx[0] * xx[0] + xx[0] * xx[1] - 3.0;

/* Restore vectors */
PetscCall(VecRestoreArrayRead(x, &xx));
PetcCall(VecRestoreArray(f, &ff));
PetscFunctionReturn(PETSC_SUCCESS);

Minor differences exist in the Fortran interface for VecGetArray() and VecRestoreArray(), as discussed in Routines that Return Fortran Allocatable Arrays. It is important to note that VecGetArray() and VecRestoreArray() do not copy the vector elements; they merely give users direct access to the vector elements. Thus, these routines require essentially no time to call and can be used efficiently.

For GPU vectors one can access either the values on the CPU as described above or one can call, for example,

VecCUDAGetArray(Vec v, PetscScalar **array);

Listing: SNES Tutorial src/snes/tutorials/ex47cu.cu

PetscCall(VecCUDAGetArrayRead(xlocal, &xarray));
PetcCall(VecCUDAGetArrayWrite(f, &farray));
if (rank) xstartshift = 1;
else xstartshift = 0;
if (rank != size - 1) xendshift = 1;
else xendshift = 0;
PetcCall(VecGetOwnershipRange(f, &fstart, NULL));
PetcCall(VecGetLocalSize(x, &lsize));
// clang-format off
try {
  thrust::for_each(
    thrust::make_zip_iterator(
      thrust::make_tuple(
        thrust::device_ptr<PetscScalar>(farray),
        thrust::device_ptr<const PetscScalar>(xarray + xstartshift),
        thrust::device_ptr<const PetscScalar>(xarray + xstartshift + 1),
        thrust::device_ptr<const PetscScalar>(xarray + xstartshift - 1),
        thrust::counting_iterator<int>(fstart),
        thrust::constant_iterator<int>(Mx),
        thrust::constant_iterator<PetscScalar>(hx))),
    thrust::make_zip_iterator(
      thrust::make_tuple(
        thrust::device_ptr<PetscScalar>(farray + lsize),
        thrust::device_ptr<const PetscScalar>(xarray + lsize - xendshift),
        thrust::device_ptr<const PetscScalar>(xarray + lsize - xendshift + 1),
        thrust::device_ptr<const PetscScalar>(xarray + lsize - xendshift - 1),
        thrust::counting_iterator<int>(fstart + lsize),
        thrust::constant_iterator<int>(Mx),
        thrust::constant_iterator<PetscScalar>(hx))),
    ApplyStencil());
}

or

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```
VecGetArrayAndMemType(Vec v, PetscScalar **array, PetscMemType *mtype);
```

which, in the first case, returns a GPU memory address and in the second case returns either a CPU or GPU memory address depending on the type of the vector. For usage with GPUs one then can launch a GPU kernel function that access the vector’s memory. When computing on GPUs `VecSetValues()` is not used! One always accesses the vector’s arrays and passes them to the GPU code.

It can also be convenient to treat the vectors entries as a Kokkos view. One first creates Kokkos vectors and then calls

```
VecGetKokkosView(Vec v, Kokkos::View<const PetscScalar*, MemorySpace> *kv)
```

to set or access the vectors entries.

Of course in order to provide the correct values to a vector one must know what parts of the vector are owned by each MPI rank. For parallel vectors, either CPU or GPU based, it is possible to determine a process’s local range with the routine

```
VecGetOwnershipRange(Vec vec, PetscInt *start, PetscInt *end);
```

The argument `start` indicates the first component owned by the local process, while `end` specifies one more than the last owned by the local process. This command is useful, for instance, in assembling parallel vectors.

The number of elements stored locally can be accessed with

```
VecGetLocalSize(Vec v, PetscInt *size);
```

The global vector length can be determined by

```
VecGetSize(Vec v, PetscInt *size);
```

**DMDA - Setting vector values**

PETSc provides an easy way to set values into the DMDA vectors and access them using the natural grid indexing. This is done with the routines

```
DMDAVecGetArray(DM da, Vec l, void *array);
... use the array indexing it with 1 or 2 or 3 dimensions ... 
... depending on the dimension of the DMDA ...
DMDAVecRestoreArray(DM da, Vec l, void *array);
DMDAVecGetArrayRead(DM da, Vec l, void *array);
... use the array indexing it with 1 or 2 or 3 dimensions ... 
... depending on the dimension of the DMDA ...
DMDAVecRestoreArrayRead(DM da, Vec l, void *array);
```

where `array` is a multidimensional C array with the same dimension as `da`, and

```
DMDAVecGetArrayDOF(DM da, Vec l, void *array);
... use the array indexing it with 2 or 3 or 4 dimensions ... 
... depending on the dimension of the DMDA ...
DMDAVecRestoreArrayDOF(DM da, Vec l, void *array);
DMDAVecGetArrayDOFRead(DM da, Vec l, void *array);
... use the array indexing it with 2 or 3 or 4 dimensions ... 
... depending on the dimension of the DMDA ...
DMDAVecRestoreArrayDOFRead(DM da, Vec l, void *array);
```
where `array` is a multidimensional C array with one more dimension than `da`. The vector `l` can be either a global vector or a local vector. The `array` is accessed using the usual global indexing on the entire grid, but the user may only refer to the local and ghost entries of this array as all other entries are undefined. For example, for a scalar problem in two dimensions one could use

```c
PetscScalar **f, **u;
...
DMDAVecGetArrayRead(DM da, Vec local,&u);
DMDAVecGetArray(DM da, Vec global,&f);
...
  f[i][j] = u[i][j] - ...
...
DMDAVecRestoreArrayRead(DM da, Vec local,&u);
DMDAVecRestoreArray(DM da, Vec global,&f);
```

Listing: SNES Tutorial src/snes/tutorials/ex3.c

```c
PetscErrorCode FormFunction(SNES snes, Vec x, Vec f, void *ctx)
{
    ApplicationCtx *user = (ApplicationCtx *)ctx;
    DM da = user->da;
    PetscScalar *ff, d;
    const PetscScalar *xx, *FF;
    PetscInt i, M, xs, xm;
    Vec xlocal;

    PetscFunctionBeginUser;
    PetscCall(DMGetLocalVector(da, &xlocal));
    /* Scatter ghost points to local vector, using the 2-step process
       DMGlobalToLocalBegin(), DMGlobalToLocalEnd().
       By placing code between these two statements, computations can
       be done while messages are in transition.
    */
    PetscCall(DMGlobalToLocalBegin(da, x, INSERT_VALUES, xlocal));
    PetscCall(DMGlobalToLocalEnd(da, x, INSERT_VALUES, xlocal));

    /* Get pointers to vector data.
       - The vector xlocal includes ghost point; the vectors x and f do
         NOT include ghost points.
       - Using DMDAVecGetArray() allows accessing the values using global ordering
    */
    PetscCall(DMDAVecGetArrayRead(da, xlocal, (void *)xx));
    PetscCall(DMDAVecGetArray(da, f, &ff));
    PetscCall(DMDAVecGetArrayRead(da, user->F, (void *)FF));

    /* Get local grid boundaries (for 1-dimensional DMDA):
       xs, xm - starting grid index, width of local grid (no ghost points)
    */
    PetscCall(DMGetCorners(da, &xs, NULL, NULL, &xm, NULL, NULL, NULL));
    PetscCall(DMGetInfo(da, NULL, &M, NULL, NULL, NULL, NULL, NULL, NULL, NULL, NULL));

    /* Set function values for boundary points; define local interior grid point range:
       (continues on next page)*/
```
xsi - starting interior grid index
xei - ending interior grid index
*/
if (xs == 0) { /* left boundary */
    ff[0] = xx[0];
    xs++;
    xm--;
}
if (xs + xm == M) { /* right boundary */
    ff[xs + xm - 1] = xx[xs + xm - 1] - 1.0;
    xm--;
}

/*
Compute function over locally owned part of the grid (interior points only)
*/
d = 1.0 / (user->h * user->h);
for (i = xs; i < xs + xm; i++) ff[i] = d * (xx[i - 1] - 2.0 * xx[i] + xx[i + 1]) + u - xx[i] * xx[i] - FF[i];

/*
Restore vectors
*/
PetscCall(DMDAVecRestoreArrayRead(da, xlocal, (void *)&xx));
PetscCall(DMDAVecRestoreArray(da, f, &ff));
PetscCall(DMDAVecRestoreArrayRead(da, user->F, (void *)&FF));
PetscCall(DMRestoreLocalVector(da, &xlocal));
PetscFunctionReturn(PETSC_SUCCESS);

The recommended approach for multi-component PDEs is to declare a struct representing the fields defined at each node of the grid, e.g.

```c
typedef struct {
    PetscScalar u,v,omega,temperature;
} Node;
```

and write the residual evaluation using

```c
Node **f,**u;
DMDAVecGetArray(DM da,Vec local,&u);
DMDAVecGetArray(DM da,Vec global,&f);
...
    f[i][j].omega = ...
...
DMDAVecRestoreArray(DM da,Vec local,&u);
DMDAVecRestoreArray(DM da,Vec global,&f);
```

The DMDAVecGetArray routines are also provided for GPU access with CUDA, HIP, and Kokkos. For example,

```c
DMDAVecGetKokkosOffsetView(DM da,Vec vec,Kokkos::View<const PetscScalar*XX*,
    _MemorySpace> *ov)
```

where *XX* can contain any number of *. This allows one to write very natural Kokkos multi-dimensional parallel for kernels that act on the local portion of DMDA vectors.
PetscErrorCode KokkosFunction(SNES snes, Vec x, Vec r, void *ctx)
{
    ApplicationCtx *user = (ApplicationCtx *)ctx;
    DM da = user->da;
    PetscScalar d;
    PetscInt M;
    Vec xl;
    PetscScalarKokkosOffsetView R;
    ConstPetscScalarKokkosOffsetView X, F;
    PetscFunctionBeginUser;
    PetscCall(DMGetLocalVector(da, &xl));
    PetscCall(DMGlobalToLocal(da, x, INSERT_VALUES, xl));
    d = 1.0 / (user->h * user->h);
    PetscCall(DMDAGetInfo(da, NULL, &M, NULL, NULL, NULL, NULL, NULL, NULL, NULL, NULL, NULL));
    PetscCall(DMDAVecGetKokkosOffsetView(da, xl, &X)); /* read only */
    PetscCall(DMDAVecGetKokkosOffsetViewWrite(da, r, &R)); /* write only */
    PetscCall(DMDAVecGetKokkosOffsetView(da, user->F, &F)); /* read only */
    Kokkos::parallel_for(
        Kokkos::RangePolicy<dMmg::Range>(R.begin(0), R.end(0)),
        KOKKOS_LAMBDA(int i) {
            if (i == 0) R(0) = X(0); /* left boundary */
            else if (i == M - 1) R(i) = X(i) - 1.0; /* right boundary */
            else R(i) = d * (X(i - 1) - 2.0 * X(i) + X(i + 1)) + X(i) * X(i) - F(i); /* interior */
        });
    PetscCall(DMDAVecRestoreKokkosOffsetView(da, xl, &X));
    PetscCall(DMDAVecRestoreKokkosOffsetViewWrite(da, r, &R));
    PetscCall(DMDAVecRestoreKokkosOffsetView(da, user->F, &F));
    PetscCall(DMRestoreLocalVector(da, &xl));
    PetscFunctionReturn(PETSC_SUCCESS);
}

The global indices of the lower left corner of the local portion of vectors obtained from DMDA as well as the local array size can be obtained with the commands

DMDAGetCorners(DM da, PetscInt *x, PetscInt *y, PetscInt *z, PetscInt *m, PetscInt *n, PetscInt *p);
DMDAGetGhostCorners(DM da, PetscInt *x, PetscInt *y, PetscInt *z, PetscInt *m, PetscInt *n, PetscInt *p);

These values can then be used as loop bounds for local function evaluations as demonstrated in the function examples above.

The first version excludes any ghost points, while the second version includes them. The routine DMDAGetGhostCorners() deals with the fact that subarrays along boundaries of the problem domain have ghost points only on their interior edges, but not on their boundary edges.

When either type of stencil is used, DMDA_STENCIL_STAR or DMDA_STENCIL_BOX, the local vectors (with the ghost points) represent rectangular arrays, including the extra corner elements in the DMDA_STENCIL_STAR case. This configuration provides simple access to the elements by employing two- (or three-) dimensional indexing. The only difference between the two cases is that when DMDA_STENCIL_STAR is used, the extra corner components are not scattered between the processes and thus contain undefined...
values that should not be used.

**DMSTAG - Setting vector values**

For structured grids with staggered data (living on elements, faces, edges, and/or vertices), the **DMStag** object is available. It behaves much like **DMDA**; see the **DMSTAG** manual page for more information.

Listing: SNES Tutorial src/dm/impls/stag/tutorials/ex6.c

```c
static PetscErrorCode UpdateVelocity_2d(const Ctx *ctx, Vec velocity, Vec stress, Vec...)
{
    Vec velocity_local, stress_local, buoyancy_local;
    PetscInt ex, ey, startx, starty, nx, ny;
    PetscInt slot_coord_next, slot_coord_element, slot_coord_prev;
    PetscInt slot_vx_left, slot_vy_down, slot_buoyancy_down, slot_buoyancy_left;
    PetscInt slot_txx, slot_tyy, slot_txy_downleft, slot_txy_downright;
    PetscScalar **arr_coord_x, **arr_coord_y;
    PetscScalar ***arr_stress, ***arr_buoyancy;
    PetscScalar ***arr_velocity;

    /* Prepare direct access to buoyancy data */
    PetscCall(DMStagGetLocationSlot(ctx->dm_buoyancy, DMSTAG_LEFT, 0, &slot_buoyancy_left));
    PetscCall(DMStagGetLocationSlot(ctx->dm_buoyancy, DMSTAG_DOWN, 0, &slot_buoyancy_down));
    PetscCall(DMGetLocalVector(ctx->dm_buoyancy, &buoyancy_local));
    PetscCall(DMGlobalToLocal(ctx->dm_buoyancy, buoyancy, INSERT_VALUES, buoyancy_local));
    PetscCall(DMStagVecGetArrayRead(ctx->dm_buoyancy, buoyancy_local, (void *)&arr_buoyancy));

    /* Prepare read-only access to stress data */
    PetscCall(DMStagGetLocationSlot(ctx->dm_stress, DMSTAG_ELEMENT, 0, &slot_txx));
    PetscCall(DMStagGetLocationSlot(ctx->dm_stress, DMSTAG_ELEMENT, 1, &slot_tyy));
    PetscCall(DMStagGetLocationSlot(ctx->dm_stress, DMSTAG_UP_LEFT, 0, &slot_txy_upleft));
    PetscCall(DMGetLocalVector(ctx->dm_stress, &stress_local));
    PetscCall(DMGlobalToLocal(ctx->dm_stress, stress, INSERT_VALUES, stress_local));
    PetscCall(DMStagVecGetArrayRead(ctx->dm_stress, stress_local, (void *)&arr_stress));

    /* Prepare read-write access to velocity data */
    PetscCall(DMStagGetLocationSlot(ctx->dm_velocity, DMSTAG_LEFT, 0, &slot_vx_left));
    PetscCall(DMStagGetLocationSlot(ctx->dm_velocity, DMSTAG_DOWN, 0, &slot_vy_down));
    PetscCall(DMGetLocalVector(ctx->dm_velocity, &velocity_local));
    PetscCall(DMGlobalToLocal(ctx->dm_velocity, velocity, INSERT_VALUES, velocity_local));
    PetscCall(DMStagVecGetArray(ctx->dm_velocity, velocity_local, &arr_velocity));
}
```

(continues on next page)
/* Prepare read-only access to coordinate data */
PetscCall(DMTagGetProductCoordinateLocationSlot(ctx->dm_velocity, DMSTAG_LEFT, &slot_coord_prev));
PetscCall(DMTagGetProductCoordinateLocationSlot(ctx->dm_velocity, DMSTAG_RIGHT, &slot_coord_next));
PetscCall(DMTagGetProductCoordinateLocationSlot(ctx->dm_velocity, DMSTAG_ELEMENT, &slot_coord_element));
PetscCall(DMTagGetProductCoordinateArrays(ctx->dm_velocity, (void *)&arr_coord_x, (void *)&arr_coord_y, NULL));

/* Iterate over interior of the domain, updating the velocities */
PetscCall(DMTagGetCorners(ctx->dm_velocity, &startx, &starty, NULL, &nx, &ny, NULL, NULL, NULL, NULL));
for (ey = starty; ey < starty + ny; ++ey) {
    for (ex = startx; ex < startx + nx; ++ex) {
        /* Update y-velocity */
        if (ey > 0) {
            const PetscScalar dx = arr_coord_x[ex][slot_coord_next] - arr_coord_x[ex][slot_coord_prev];
            const PetscScalar dy = arr_coord_y[ey][slot_coord_element] - arr_coord_y[ey - 1][slot_coord_element];
            const PetscScalar B = arr_buoyancy[ey][ex][slot_buoyancy_down];
            arr_velocity[ey][ex][slot_vy_down] += B * ctx->dt * ((arr_stress[ey][ex][slot_txy_downright] - arr_stress[ey][ex][slot_txy_downleft]) / dx + (arr_stress[ey][ex][slot_tyy] - arr_stress[ey - 1][ex][slot_tyy]) / dy);
        }
        /* Update x-velocity */
    }
}

DMPLEX - Setting vector values

See DMPlex: Unstructured Grids for discussion on setting vector values with DMPLEX.

DMNETWORK - Setting vector values

See Networks for discussion on setting vector values with DMNETWORK.
2.1.3 Basic Vector Operations

Table 2.1: PETSc Vector Operations

<table>
<thead>
<tr>
<th>Function Name</th>
<th>Operation</th>
</tr>
</thead>
<tbody>
<tr>
<td>VecAXPY(Vec y, PetscScalar a, Vec x);</td>
<td>y = y + a * x</td>
</tr>
<tr>
<td>VecAYPX(Vec y, PetscScalar a, Vec x);</td>
<td>x = x + a * y</td>
</tr>
<tr>
<td>VecWAXPY(Vec w, PetscScalar a, Vec x, Vec y);</td>
<td>w = a * x + y</td>
</tr>
<tr>
<td>VecAPBXY(Vec y, PetscScalar a, PetscScalar b, Vec x);</td>
<td>y = a * x + b * y</td>
</tr>
<tr>
<td>VecAXPYPCZ(Vec z, PetscScalar a, PetscScalar b, PetscScalar c, Vec x, Vec y);</td>
<td>z = a * x + b * y + c * z</td>
</tr>
<tr>
<td>VecScale(Vec x, PetscScalar a);</td>
<td>x = a * x</td>
</tr>
<tr>
<td>VecDot(Vec x, Vec y, PetscScalar *r);</td>
<td>r = \bar{x}^T y</td>
</tr>
<tr>
<td>VecTDot(Vec x, Vec y, PetscScalar *r);</td>
<td>r = x' y</td>
</tr>
<tr>
<td>VecNorm(Vec x, NormType type, PetscReal *r);</td>
<td>r =</td>
</tr>
<tr>
<td>VecSum(Vec x, PetscScalar *r);</td>
<td>r = \sum x_i</td>
</tr>
<tr>
<td>VecCopy(Vec x, Vec y);</td>
<td>y = x</td>
</tr>
<tr>
<td>VecSwap(Vec x, Vec y);</td>
<td>y = x while x = y</td>
</tr>
<tr>
<td>VecPointwiseMult(Vec w, Vec x, Vec y);</td>
<td>w_i = x_i * y_i</td>
</tr>
<tr>
<td>VecPointwiseDivide(Vec w, Vec x, Vec y);</td>
<td>w_i = x_i / y_i</td>
</tr>
<tr>
<td>VecMDDot(Vec x, PetscInt n, Vec y[], PetscScalar *r);</td>
<td>r_i = x_i^T * y_i</td>
</tr>
<tr>
<td>VecMDot(Vec x, PetscInt n, Vec y[], PetscScalar *r);</td>
<td>r_i = x_i^T * y_i</td>
</tr>
<tr>
<td>VecMDDot(Vec x, PetscInt n, Vec y[], PetscScalar *r);</td>
<td>r_i = x_i^T * y_i</td>
</tr>
<tr>
<td>VecMax(Vec x, PetscInt *idx, PetscScalar *r);</td>
<td>r = max x_i</td>
</tr>
<tr>
<td>VecMin(Vec x, PetscInt *idx, PetscScalar *r);</td>
<td>r = min x_i</td>
</tr>
<tr>
<td>VecAbs(Vec x);</td>
<td>x_i =</td>
</tr>
<tr>
<td>VecReciprocal(Vec x);</td>
<td>x_i = 1 / x_i</td>
</tr>
<tr>
<td>VecShift(Vec x, PetscScalar s);</td>
<td>x_i = s + x_i</td>
</tr>
<tr>
<td>VecSet(Vec x, PetscScalar alpha);</td>
<td>x_i = alpha</td>
</tr>
</tbody>
</table>

As listed in the table, we have chosen certain basic vector operations to support within the PETSc vector library. These operations were selected because they often arise in application codes. The `NormType` argument to `VecNorm()` is one of `NORM_1`, `NORM_2`, or `NORM_INFINITY`. The 1-norm is \( \sum |x_i| \), the 2-norm is \( (\sum x_i^2)^{1/2} \) and the infinity norm is max \( |x_i| \).

In addition to `VecDot()` and `VecMDot()` and `VecNorm()`, PETSc provides split phase versions of these that allow several independent inner products and/or norms to share the same communication (thus improving parallel efficiency). For example, one may have code such as

```plaintext
VecDot(Vec x, Vec y, PetscScalar *dot);
VecMDot(Vec x, PetscInt nv, Vec y[], PetscScalar *dot);
VecNorm(Vec x, NormType NORM_2, PetscReal *norm2);
VecNorm(Vec x, NormType NORM_1, PetscReal *norm1);
```

This code works fine, but it performs four separate parallel communication operations. Instead, one can write

```plaintext
VecDotBegin(Vec x, Vec y, PetscScalar *dot);
VecMDotBegin(Vec x, PetscInt nv, Vec y[], PetscScalar *dot);
VecNormBegin(Vec x, NormType NORM_2, PetscReal *norm2);
VecNormBegin(Vec x, NormType NORM_1, PetscReal *norm1);
VecDotEnd(Vec x, Vec y, PetscScalar *dot);
VecMDotEnd(Vec x, PetscInt nv, Vec y[], PetscScalar *dot);
```

(continues on next page)
With this code, the communication is delayed until the first call to `VecxxxEnd()` at which a single MPI reduction is used to communicate all the required values. It is required that the calls to the `VecxxxEnd()` are performed in the same order as the calls to the `VecxxxBegin()`; however, if you mistakenly make the calls in the wrong order, PETSc will generate an error informing you of this. There are additional routines `VecTDotBegin()` and `VecTDotEnd()`, `VecMTDotBegin()`, `VecMTDotEnd()`.

For GPU vectors (like CUDA), the numerical computations will, by default, run on the GPU. Any scalar output, like the result of a `VecDot()` are placed in CPU memory.

### 2.1.4 Local/global vectors and communicating between vectors

Many PDE problems require the use of ghost (or halo) values in each MPI rank or even more general parallel communication of vector values. These values are needed in order to perform function evaluation on that rank. The exact structure of the ghost values needed depends on the type of grid being used. DM provides a uniform API for communicating the needed values. We introduce the concept in detail for DMDA.

Each DM object defines the layout of two vectors: a distributed global vector and a local vector that includes room for the appropriate ghost points. The DM object provides information about the size and layout of these vectors, but does not internally allocate any associated storage space for field values. Instead, the user can create vector objects that use the DM layout information with the routines

```c
DMCreateGlobalVector(DM da, Vec *g);
DMCreateLocalVector(DM da, Vec *l);
```

These vectors will generally serve as the building blocks for local and global PDE solutions, etc. If additional vectors with such layout information are needed in a code, they can be obtained by duplicating `l` or `g` via `VecDuplicate()` or `VecDuplicateVecs()`.

We emphasize that a distributed array provides the information needed to communicate the ghost value information between processes. In most cases, several different vectors can share the same communication information (or, in other words, can share a given DM). The design of the DM object makes this easy, as each DM operation may operate on vectors of the appropriate size, as obtained via `DMCreateLocalVector()` and `DMCreateGlobalVector()` or as produced by `VecDuplicate()`.

At certain stages of many applications, there is a need to work on a local portion of the vector that includes the ghost points. This may be done by scattering a global vector into its local parts by using the two-stage commands

```c
DMLocalToGlobalBegin(DM da, Vec g, InsertMode iora, Vec l);
DMLocalToGlobalEnd(DM da, Vec g, InsertMode iora, Vec l);
```

which allow the overlap of communication and computation. Since the global and local vectors, given by `g` and `l`, respectively, must be compatible with the distributed array, `da`, they should be generated by `DMCreateGlobalVector()` and `DMCreateLocalVector()` (or be duplicates of such a vector obtained via `VecDuplicate()`). The `InsertMode` can be either `ADD_VALUES` or `INSERT_VALUES`.

One can scatter the local vectors into the distributed global vector with the command

```c
DMLocalToGlobal(DM da, Vec l, InsertMode mode, Vec g);
```

or the commands
DMLocalToGlobalBegin(DM da, Vec l, InsertMode mode, Vec g);

/* (Computation to overlap with communication) */
DMLocalToGlobalEnd(DM da, Vec l, InsertMode mode, Vec g);

In general this is used with an InsertMode of ADD_VALUES, because if one wishes to insert values into the global vector they should just access the global vector directly and put in the values.

A third type of distributed array scatter is from a local vector (including ghost points that contain irrelevant values) to a local vector with correct ghost point values. This scatter may be done with the commands

DMLocalToLocalBegin(DM da, Vec l1, InsertMode iora, Vec l2);
DMLocalToLocalEnd(DM da, Vec l1, InsertMode iora, Vec l2);

Since both local vectors, l1 and l2, must be compatible with the distributed array, da, they should be generated by DMCreatelocalVector() (or be duplicates of such vectors obtained via VecDuplicate()). The InsertMode can be either ADD_VALUES or INSERT_VALUES.

In most applications the local ghosted vectors are only needed during user “function evaluations”. PETSc provides an easy, lightweight (requiring essentially no CPU time) way to obtain these work vectors and return them when they are no longer needed. This is done with the routines

DMGetLocalVector(DM da, Vec *l);
... use the local vector l ...
DMRestoreLocalVector(DM da, Vec *l);

### 2.1.5 Low-level Vector Communication

Most users of PETSc, who can utilize a DM will not need to utilize the lower-level routines discussed in the rest of this section and should skip ahead to Matrices.

To facilitate creating general vector scatters and gathers used, for example, in updating ghost points for problems for which no DM currently exists PETSc employs the concept of an index set, via the IS class. An index set, which is a generalization of a set of integer indices, is used to define scatters, gathers, and similar operations on vectors and matrices. In fact, much of the underlying code that implements DMGlobalToLocal communication are built on the infrastructure discussed below.

The following command creates an index set based on a list of integers:

ISCreateGeneral(MPI_Comm comm, PetscInt n, PetscInt *indices, PetscCopyMode mode, IS *is);

When mode is PETSC_COPY_VALUES, this routine copies the n indices passed to it by the integer array indices. Thus, the user should be sure to free the integer array indices when it is no longer needed, perhaps directly after the call to ISCreateGeneral(). The communicator, comm, should consist of all processes that will be using the IS.

Another standard index set is defined by a starting point (first) and a stride (step), and can be created with the command

ISCreateStride(MPI_Comm comm, PetscInt n, PetscInt first, PetscInt step, IS *is);

Index sets can be destroyed with the command

ISDestroy(IS &is);

On rare occasions the user may need to access information directly from an index set. Several commands assist in this process:
The function `ISGetIndices()` returns a pointer to a list of the indices in the index set. For certain index sets, this may be a temporary array of indices created specifically for a given routine. Thus, once the user finishes using the array of indices, the routine

```c
ISRestoreIndices(IS is, PetscInt **indices);
```

should be called to ensure that the system can free the space it may have used to generate the list of indices.

A blocked version of the index sets can be created with the command

```c
ISCreateBlock(MPI_Comm comm, PetscInt bs, PetscInt n, PetscInt *indices, PetscCopyMode mode, IS *is);
```

This version is used for defining operations in which each element of the index set refers to a block of `bs` vector entries. Related routines analogous to those described above exist as well, including `ISBlockGetIndices()`, `ISBlockGetSize()`, `ISBlockGetLocalSize()`, `ISGetBlockSize()`.

Most PETSc applications use a particular DM object to manage the details of the communication needed for their grids. In some rare cases however codes need to directly setup their required communication patterns. This is done using PETSc's VecScatter and PetscSF (for more general data than vectors). One can select any subset of the components of a vector to insert or add to any subset of the components of another vector. We refer to these operations as generalized scatters, though they are actually a combination of scatters and gathers.

To copy selected components from one vector to another, one uses the following set of commands:

```c
VecScatterCreate(Vec x, IS ix, Vec y, IS iy, VecScatter *ctx);
VecScatterBegin(VecScatter ctx, Vec x, Vec y, INSERT_VALUES, SCATTER_FORWARD);
VecScatterEnd(VecScatter ctx, Vec x, Vec y, INSERT_VALUES, SCATTER_FORWARD);
VecScatterDestroy(VecScatter *ctx);
```

Here `ix` denotes the index set of the first vector, while `iy` indicates the index set of the destination vector. The vectors can be parallel or sequential. The only requirements are that the number of entries in the index set of the first vector, `ix`, equals the number in the destination index set, `iy`, and that the vectors be long enough to contain all the indices referred to in the index sets. If both `x` and `y` are parallel, their communicator must have the same set of processes, but their process order can be different. The argument `INSERT_VALUES` specifies that the vector elements will be inserted into the specified locations of the destination vector, overwriting any existing values. To add the components, rather than insert them, the user should select the option `ADD_VALUES` instead of `INSERT_VALUES`. One can also use `MAX_VALUES` or `MIN_VALUES` to replace destination with the maximal or minimal of its current value and the scattered values.

To perform a conventional gather operation, the user simply makes the destination index set, `iy`, be a stride index set with a stride of one. Similarly, a conventional scatter can be done with an initial (sending) index set consisting of a stride. The scatter routines are collective operations (i.e. all processes that own a parallel vector must call the scatter routines). When scattering from a parallel vector to sequential vectors, each process has its own sequential vector that receives values from locations as indicated in its own index set. Similarly, in scattering from sequential vectors to a parallel vector, each process has its own sequential vector that makes contributions to the parallel vector.

**Caution:** When `INSERT_VALUES` is used, if two different processes contribute different values to the same component in a parallel vector, either value may end up being inserted. When `ADD_VALUES` is used, the
correct sum is added to the correct location.

In some cases one may wish to “undo” a scatter, that is perform the scatter backwards, switching the roles of the sender and receiver. This is done by using

```c
VecScatterBegin(VecScatter ctx, Vec y, Vec x, INSERT_VALUES, SCATTER_REVERSE);
VecScatterEnd(VecScatter ctx, Vec y, Vec x, INSERT_VALUES, SCATTER_REVERSE);
```

Note that the roles of the first two arguments to these routines must be swapped whenever the SCATTER_REVERSE option is used.

Once a VecScatter object has been created it may be used with any vectors that have the appropriate parallel data layout. That is, one can call VecScatterBegin() and VecScatterEnd() with different vectors than used in the call to VecScatterCreate() as long as they have the same parallel layout (number of elements on each process are the same). Usually, these “different” vectors would have been obtained via calls to VecDuplicate() from the original vectors used in the call to VecScatterCreate().

VecGetValues() can only access local values from the vector. To get off-process values, the user should create a new vector where the components are to be stored, and then perform the appropriate vector scatter. For example, if one desires to obtain the values of the 100th and 200th entries of a parallel vector, p, one could use a code such as that below. In this example, the values of the 100th and 200th components are placed in the array values. In this example each process now has the 100th and 200th component, but obviously each process could gather any elements it needed, or none by creating an index set with no entries.

```c
Vec p, x; /* initial vector, destination vector */
VecScatter scatter; /* scatter context */
IS from, to; /* index sets that define the scatter */
PetscScalar *values;
PetscInt idx_from[] = {100, 200}, idx_to[] = {0, 1};

VecCreateSeq(PETSC_COMM_SELF, 2, &x);
ISCreateGeneral(PETSC_COMM_SELF, 2, idx_from, PETSC_COPY_VALUES, &from);
ISCreateGeneral(PETSC_COMM_SELF, 2, idx_to, PETSC_COPY_VALUES, &to);
VecScatterCreate(p, from, to, &scatter);
VecScatterBegin(scatter, p, x, INSERT_VALUES, SCATTER_FORWARD);
VecScatterEnd(scatter, p, x, INSERT_VALUES, SCATTER_FORWARD);
VecGetArray(x, &values);
ISDestroy(&from);
ISDestroy(&to);
VecScatterDestroy(&scatter);
```

The scatter comprises two stages, in order to allow overlap of communication and computation. The introduction of the VecScatter context allows the communication patterns for the scatter to be computed once and then reused repeatedly. Generally, even setting up the communication for a scatter requires communication; hence, it is best to reuse such information when possible.

Generalized scatters provide a very general method for managing the communication of required ghost values for unstructured grid computations. One scatters the global vector into a local “ghosted” work vector, performs the computation on the local work vectors, and then scatters back into the global solution vector. In the simplest case this may be written as

```c
VecScatterBegin(VecScatter scatter, Vec globalin, Vec localin, InsertMode INSERT_VALUES, ScatterMode SCATTER_FORWARD);
VecScatterEnd(VecScatter scatter, Vec globalin, Vec localin, InsertMode INSERT_VALUES, ScatterMode SCATTER_FORWARD);
/* For example, do local calculations from localin to localout */
...
VecScatterBegin(VecScatter scatter, Vec localout, Vec globalout, InsertMode ADD_VALUES,
```

(continues on next page)
Local to global mappings

When working with a global representation of a vector (usually on a vector obtained with `DMCreateGlobalVector()`) and a local representation of the same vector that includes ghost points required for local computation (obtained with `DMCreateLocalVector()`). PETSc provides routines to help map indices from a local numbering scheme to the PETSc global numbering scheme. This is done via the following routines:

```c
ISLocalToGlobalMappingCreate(MPI_Comm comm, PetscInt bs, PetscInt N, PetscInt* globalnum,
    PetscCopyMode mode, ISLocalToGlobalMapping* ctx);
ISLocalToGlobalMappingApply(ISLocalToGlobalMapping ctx, PetscInt n, PetscInt* in,
    PetscInt* out);
ISLocalToGlobalMappingApplyIS(ISLocalToGlobalMapping ctx, IS isin, IS* isout);
ISLocalToGlobalMappingDestroy(ISLocalToGlobalMapping* ctx);
```

Here \( N \) denotes the number of local indices, \( \text{globalnum} \) contains the global number of each local number, and \( \text{ISLocalToGlobalMapping} \) is the resulting PETSc object that contains the information needed to apply the mapping with either \( \text{ISLocalToGlobalMappingApply()} \) or \( \text{ISLocalToGlobalMappingApplyIS()} \).

Note that the \( \text{ISLocalToGlobalMapping} \) routines serve a different purpose than the AO routines. In the former case they provide a mapping from a local numbering scheme (including ghost points) to a global numbering scheme, while in the latter they provide a mapping between two global numbering schemes. In fact, many applications may use both AO and \( \text{ISLocalToGlobalMapping} \) routines. The AO routines are first used to map from an application global ordering (that has no relationship to parallel processing etc.) to the PETSc ordering scheme (where each process has a contiguous set of indices in the numbering). Then in order to perform function or Jacobian evaluations locally on each process, one works with a local numbering scheme that includes ghost points. The mapping from this local numbering scheme back to the global PETSc numbering can be handled with the \( \text{ISLocalToGlobalMapping} \) routines.

If one is given a list of block indices in a global numbering, the routine

```c
ISGlobalToLocalMappingApplyBlock(ISLocalToGlobalMapping ctx,
    ISGlobalToLocalMappingMode type, PetscInt nin, PetscInt idxin[], PetscInt* nout,
    PetscInt idxout[]);
```

will provide a new list of indices in the local numbering. Again, negative values in \( \text{idxin} \) are left unmapped. But, in addition, if \( \text{type} \) is set to \( \text{IS_GTOLM_MASK} \), then \( \text{nout} \) is set to \( \text{nin} \) and all global values in \( \text{idxin} \) that are not represented in the local to global mapping are replaced by -1. When \( \text{type} \) is set to \( \text{IS_GTOLM_DROP} \), the values in \( \text{idxin} \) that are not represented locally in the mapping are not included in \( \text{idxout} \), so that potentially \( \text{nout} \) is smaller than \( \text{nin} \). One must pass in an array long enough to hold all the indices. One can call \( \text{ISGlobalToLocalMappingApplyBlock()} \) with \( \text{idxout} \) equal to \( \text{NULL} \) to determine the required length (returned in \( \text{nout} \)) and then allocate the required space and call \( \text{ISGlobalToLocalMappingApplyBlock()} \) a second time to set the values.

Often it is convenient to set elements into a vector using the local node numbering rather than the global node numbering (e.g., each process may maintain its own sublist of vertices and elements and number them locally). To set values into a vector with the local numbering, one must first call

---

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and then call

\begin{verbatim}
VecSetValuesLocal(Vec x, PetscInt n, const PetscInt indices[], const PetscScalar *values[], INSERT_VALUES);
\end{verbatim}

Now the indices use the local numbering, rather than the global, meaning the entries lie in \([0, n)\) where \(n\) is the local size of the vector. Global vectors obtained from DM’s already have the global to local mapping provided by the "DM.

To assemble global stiffness matrices, one can use these global indices with MatSetValues() or MatSetValuesStencil(). Alternately, the global node number of each local node, including the ghost nodes, can be obtained by calling

\begin{verbatim}
DMGetLocalToGlobalMapping(DM da, ISLocalToGlobalMapping *map);
\end{verbatim}

followed by

\begin{verbatim}
VecSetLocalToGlobalMapping(Vec v, ISLocalToGlobalMapping map);
MatSetLocalToGlobalMapping(Mat A, ISLocalToGlobalMapping rmapping, ISLocalToGlobalMapping cmapping);
\end{verbatim}

Now entries may be added to the vector and matrix using the local numbering and VecSetValuesLocal() and MatSetValuesLocal().

The example SNES Tutorial ex5 illustrates the use of a distributed array in the solution of a nonlinear problem. The analogous Fortran program is SNES Tutorial ex5f90; see SNES: Nonlinear Solvers for a discussion of the nonlinear solvers.

Global Vectors with locations for ghost values

There are two minor drawbacks to the basic approach described above for unstructured grids:

- the extra memory requirement for the local work vector, \texttt{localin}, which duplicates the memory in \texttt{globalin}, and
- the extra time required to copy the local values from \texttt{localin} to \texttt{globalin}.

An alternative approach is to allocate global vectors with space preallocated for the ghost values; at the local level vector interfaces this may be done with either

\begin{verbatim}
VecCreateGhost(MPI_Comm comm, PetscInt n, PetscInt N, PetscInt nghost, PetscInt *ghosts, Vec *vv)
\end{verbatim}

or

\begin{verbatim}
VecCreateGhostWithArray(MPI_Comm comm, PetscInt n, PetscInt N, PetscInt nghost, PetscInt *ghosts, PetscScalar *array, Vec *vv)
\end{verbatim}

Here \(n\) is the number of local vector entries, \(N\) is the number of global entries (or \texttt{NULL}) and \(nghost\) is the number of ghost entries. The array \texttt{ghosts} is of size \(nghost\) and contains the global vector location for each local ghost location. Using VecDuplicate() or VecDuplicateVecs() on a ghosted vector will generate additional ghosted vectors.

In many ways, a ghosted vector behaves just like any other MPI vector created by VecCreateMPI(). The difference is that the ghosted vector has an additional “local” representation that allows one to access the ghost locations. This is done through the call to
The vector \( l \) is a sequential representation of the parallel vector \( g \) that shares the same array space (and hence numerical values); but allows one to access the “ghost” values past “the end of the” array. Note that one access the entries in \( l \) using the local numbering of elements and ghosts, while they are accessed in \( g \) using the global numbering.

A common usage of a ghosted vector is given by

\begin{verbatim}
VecGhostGetLocalForm(Vec g, Vec *l);
\end{verbatim}

The routines \texttt{VecGhostUpdateBegin()} and \texttt{VecGhostUpdateEnd()} are equivalent to the routines \texttt{VecScatterBegin()} and \texttt{VecScatterEnd()} above except that since they are scattering into the ghost locations, they do not need to copy the local vector values, which are already in place. In addition, the user does not have to allocate the local work vector, since the ghosted vector already has allocated slots to contain the ghost values.

The input arguments \texttt{INSERT\_VALUES} and \texttt{SCATTER\_FORWARD} cause the ghost values to be correctly updated from the appropriate process. The arguments \texttt{ADD\_VALUES} and \texttt{SCATTER\_REVERSE} update the “local” portions of the vector from all the other processes’ ghost values. This would be appropriate, for example, when performing a finite element assembly of a load vector. One can also use \texttt{MAX\_VALUES} or \texttt{MIN\_VALUES} with \texttt{SCATTER\_REVERSE}.

\texttt{DMPLEX} does not yet have support for ghosted vectors sharing memory with the global representation. This is work in progress, if you have interest in this feature please contact the PETSc community members.

\textit{Partitioning} discusses the important topic of partitioning an unstructured grid.

### 2.1.6 Application Orderings

When writing parallel PDE codes, there is extra complexity caused by having multiple ways of indexing (numbering) and ordering objects such as vertices and degrees of freedom. For example, a grid generator or partitioner may renumber the nodes, requiring adjustment of the other data structures that refer to these objects; see Figure \textit{Natural Ordering and PETSc Ordering for a 2D Distributed Array (Four Processes)}. PETSc provides a variety of tools to help to manage the mapping amongst the various numbering systems. The most basic are the \texttt{AO} (application ordering), which enables mapping between different global (cross-process) numbering schemes.

In many applications it is desirable to work with one or more “orderings” (or numberings) of degrees of freedom, cells, nodes, etc. Doing so in a parallel environment is complicated by the fact that each process cannot keep complete lists of the mappings between different orderings. In addition, the orderings used in the PETSc linear algebra routines (often contiguous ranges) may not correspond to the “natural” orderings for the application.

PETSc provides certain utility routines that allow one to deal cleanly and efficiently with the various orderings. To define a new application ordering (called an \texttt{AO} in PETSc), one can call the routine

\begin{verbatim}
VecGhostGetLocalForm(Vec g, Vec *l);
VecGhostUpdateBegin(Vec globalin, InsertMode INSERT\_VALUES, ScatterMode SCATTER\_FORWARD);
VecGhostUpdateEnd(Vec globalin, InsertMode INSERT\_VALUES, ScatterMode SCATTER\_FORWARD);
VecGhostGetLocalForm(Vec globalin, Vec *localin);
VecGhostGetLocalForm(Vec globalout, Vec *localout);
... Do local calculations from localin to localout ...
VecGhostRestoreLocalForm(Vec globalin, Vec *localin);
VecGhostRestoreLocalForm(Vec globalout, Vec *localout);
VecGhostUpdateBegin(Vec globalout, InsertMode ADD\_VALUES, ScatterMode SCATTER\_REVERSE);
VecGhostUpdateEnd(Vec globalout, InsertMode ADD\_VALUES, ScatterMode SCATTER\_REVERSE);
\end{verbatim}
AOCreateBasic(MPI_Comm comm, PetscInt n, const PetscInt apordering[], const PetscInt petscordering[], AO *ao);

The arrays `apordering` and `petscordering`, respectively, contain a list of integers in the application ordering and their corresponding mapped values in the PETSc ordering. Each process can provide whatever subset of the ordering it chooses, but multiple processes should never contribute duplicate values. The argument `n` indicates the number of local contributed values.

For example, consider a vector of length 5, where node 0 in the application ordering corresponds to node 3 in the PETSc ordering. In addition, nodes 1, 2, 3, and 4 of the application ordering correspond, respectively, to nodes 2, 1, 4, and 0 of the PETSc ordering. We can write this correspondence as

\[ \{0, 1, 2, 3, 4\} \rightarrow \{3, 2, 1, 4, 0\}. \]

The user can create the PETSc AO mappings in a number of ways. For example, if using two processes, one could call

```c
AOCreateBasic(PETSC_COMM_WORLD, 2, {0, 3}, {3, 4}, &ao);
```

on the first process and

```c
AOCreateBasic(PETSC_COMM_WORLD, 3, {1, 2, 4}, {2, 1, 0}, &ao);
```

on the other process.

Once the application ordering has been created, it can be used with either of the commands

```c
AOPetscToApplication(AO ao, PetscInt n, PetscInt *indices);
AOApplicationToPetsc(AO ao, PetscInt n, PetscInt *indices);
```

Upon input, the `n`-dimensional array `indices` specifies the indices to be mapped, while upon output, `indices` contains the mapped values. Since we, in general, employ a parallel database for the AO mappings, it is crucial that all processes that called `AOCreateBasic()` also call these routines; these routines cannot be called by just a subset of processes in the MPI communicator that was used in the call to `AOCreateBasic()`.

An alternative routine to create the application ordering, AO, is

```c
AOCreateBasicIS(IS apordering, IS petscordering, AO *ao);
```

where index sets are used instead of integer arrays.

The mapping routines

```c
AOPetscToApplicationIS(AO ao, IS indices);
AOApplicationToPetscIS(AO ao, IS indices);
```

will map index sets (IS objects) between orderings. Both the `AOXxxToYyy()` and `AOXxxToYyyIS()` routines can be used regardless of whether the AO was created with a `AOCreateBasic()` or `AOCreateBasicIS()`.

The AO context should be destroyed with `AODestroy(AO *ao)` and viewed with `AOView(AO ao, PetscViewer viewer)`.

Although we refer to the two orderings as “PETSc” and “application” orderings, the user is free to use them both for application orderings and to maintain relationships among a variety of orderings by employing several AO contexts.

The `AOXxxToxx()` routines allow negative entries in the input integer array. These entries are not mapped; they simply remain unchanged. This functionality enables, for example, mapping neighbor lists that use negative numbers to indicate nonexistent neighbors due to boundary conditions, etc.
Since the global ordering that PETSc uses to manage its parallel vectors (and matrices) does not usually correspond to the “natural” ordering of a two- or three-dimensional array, the DMDA structure provides an application ordering AO (see Application Orderings) that maps between the natural ordering on a rectangular grid and the ordering PETSc uses to parallelize. This ordering context can be obtained with the command

```
DMDAGetAO(DM da, AO *ao);
```

In Figure Natural Ordering and PETSc Ordering for a 2D Distributed Array (Four Processes) we indicate the orderings for a two-dimensional distributed array, divided among four processes.

![Natural Ordering and PETSc Ordering for a 2D Distributed Array (Four Processes)](image)

**2.2 Matrices**

PETSc provides a variety of matrix implementations because no single matrix format is appropriate for all problems. Currently, we support dense storage and compressed sparse row storage (both sequential and parallel versions) for CPU and GPU based matrices, as well as several specialized formats. Additional specialized formats can be easily added.

This chapter describes the basics of using PETSc matrices in general (regardless of the particular format chosen) and discusses tips for efficient use of the several simple uniprocess and parallel matrix types. The use of PETSc matrices involves the following actions: create a particular type of matrix, insert values into it, process the matrix, use the matrix for various computations, and finally destroy the matrix. The application code does not need to know or care about the particular storage formats of the matrices.
2.2.1 Creating matrices

As with vectors, PETSc has APIs that allow the user to specify the exact details of the matrix creation process but also DM based creation routines that handle most of the details automatically for specific families of applications. This is done with

\[
\text{DMCreateMatrix}(\text{DM } \text{dm}, \text{Mat } *A)
\]

The type of matrix created can be controlled with either

\[
\text{DMSetMatType}(\text{DM } \text{dm}, \text{MatType } \text{MATAIJ} \text{ or } \text{MATBAIJ} \text{ or } \text{MATAIJCUSPARSE} \text{ etc})
\]

or with

\[
\text{DMSetFromOptions}(\text{DM } \text{dm})
\]

and the options database option \text{-dm_mat_type <aij or baij or aijcusparse etc>} Matrices can be created for CPU usage, for GPU usage and for usage on both the CPUs and GPUs.

The creation of DM objects is discussed in DMDA - Creating vectors for structured grids, DMPLEX - Creating vectors for unstructured grids, DMNETWORK - Creating vectors for networks.

2.2.2 Low-level matrix creation routines

When using a DM is not practical for a particular application one can create matrices directly using

\[
\text{MatCreate}(\text{MPI_Comm } \text{comm}, \text{Mat } *A) \\
\text{MatSetSizes}(\text{Mat } A, \text{PetscInt } m, \text{PetscInt } n, \text{PetscInt } M, \text{PetscInt } N)
\]

This routine generates a sequential matrix when running one process and a parallel matrix for two or more processes; the particular matrix format is set by the user via options database commands. The user specifies either the global matrix dimensions, given by \text{M} \text{ and } \text{N} or the local dimensions, given by \text{m} \text{ and } \text{n} while PETSc completely controls memory allocation. This routine facilitates switching among various matrix types, for example, to determine the format that is most efficient for a certain application. By default, MatCreate() employs the sparse AIJ format, which is discussed in detail in Sparse Matrices. See the manual pages for further information about available matrix formats.

2.2.3 Assembling (putting values into) matrices

To insert or add entries to a matrix on CPUs, one can call a variant of MatSetValues(), either

\[
\text{MatSetValues}(\text{Mat } A, \text{PetscInt } m, \text{const PetscInt } idxm[], \text{PetscInt } n, \text{const PetscInt } idxn[], \text{const PetscScalar } values[], \text{INSERT_VALUES});
\]

or

\[
\text{MatSetValues}(\text{Mat } A, \text{PetscInt } m, \text{const PetscInt } idxm[], \text{PetscInt } n, \text{const PetscInt } idxn[], \text{const PetscScalar } values[], \text{ADD_VALUES});
\]

This routine inserts or adds a logically dense subblock of dimension \text{m*}n into the matrix. The integer indices \text{idxm} and \text{idxn}, respectively, indicate the global row and column numbers to be inserted. MatSetValues() uses the standard C convention, where the row and column matrix indices begin with zero \textit{regardless of the programming language employed}. The array \text{values} is logically two-dimensional, containing the values that are to be inserted. By default the values are given in row major order, which is the opposite of the
Fortran convention, meaning that the value to be put in row $idxm[i]$ and column $idxn[j]$ is located in $values[i*n+j]$. To allow the insertion of values in column major order, one can call the command

```
MatSetOption(Mat A, MAT_ROW_ORIENTED, PETSC_FALSE);
```

**Warning:** Several of the sparse implementations do not currently support the column-oriented option.

This notation should not be a mystery to anyone. For example, to insert one matrix into another when using MATLAB, one uses the command $A(im,in) = B$; where $im$ and $in$ contain the indices for the rows and columns. This action is identical to the calls above to `MatSetValues()`.

When using the block compressed sparse row matrix format (MATSEQBAIJ or MATMPIBAIJ), one can insert elements more efficiently using the block variant, `MatSetValuesBlocked()` or `MatSetValuesBlockedLocal()`.

The function `MatSetOption()` accepts several other inputs; see the manual page for details.

After the matrix elements have been inserted or added into the matrix, they must be processed (also called “assembled”) before they can be used. The routines for matrix processing are

```
MatAssemblyBegin(Mat A, MAT_FINAL_ASSEMBLY);
MatAssemblyEnd(Mat A, MAT_FINAL_ASSEMBLY);
```

By placing other code between these two calls, the user can perform computations while messages are in transit. Calls to `MatSetValues()` with the INSERT_VALUES and ADD_VALUES options cannot be mixed without intervening calls to the assembly routines. For such intermediate assembly calls the second routine argument typically should be `MAT_FLUSH.getAssembly`, which omits some of the work of the full assembly process. `MAT_FINAL_ASSEMBLY` is required only in the last matrix assembly before a matrix is used.

Even though one may insert values into PETSc matrices without regard to which process eventually stores them, for efficiency reasons we usually recommend generating most entries on the process where they are destined to be stored. To help the application programmer with this task for matrices that are distributed across the processes by ranges, the routine

```
MatGetOwnershipRange(Mat A, PetscInt *first_row, PetscInt *last_row);
```

informs the user that all rows from $first_row$ to $last_row-1$ (since the value returned in $last_row$ is one more than the global index of the last local row) will be stored on the local process.

In the sparse matrix implementations, once the assembly routines have been called, the matrices are compressed and can be used for matrix-vector multiplication, etc. Any space for preallocated nonzeros that was not filled by a call to `MatSetValues()` or a related routine is compressed out by assembling with `MAT_FINAL_ASSEMBLY`. If you intend to use that extra space later, be sure to insert explicit zeros before assembling with `MAT_FINAL_ASSEMBLY` so the space will not be compressed out. Once the matrix has been assembled, inserting new values will be expensive since it will require copies and possible memory allocation.

One may repeatedly assemble matrices that retain the same nonzero pattern (such as within a nonlinear or time-dependent problem). Where possible, data structures and communication information will be reused (instead of regenerated) during successive steps, thereby increasing efficiency. See KSP Tutorial ex5 for a simple example of solving two linear systems that use the same matrix data structure.

For matrices associated with `DMDA` there is a higher-level interface for providing the numerical values based on the concept of stencils. See the manual page of `MatSetValuesStencil()` for usage.

For GPUs the routines `MatSetPreallocationCOO()` and `MatSetValuesCOO()` should be used for efficient matrix assembly instead of `MatSetValues()`.

We now introduce the various families of PETSc matrices. `DMCreateMatrix()` manages the preallocation process (introduced below) automatically so many users do not need to worry about the details of the preallocation process.

### 2.2. Matrices
Matrix and Vector Layouts and Storage Locations

The layout of PETSc matrices across MPI ranks is defined by two things

- the layout of the two compatible vectors in the computation of the matrix-vector product \( y = A \times x \)
- and
- the memory where various parts of the matrix are stored across the MPI ranks.

PETSc vectors always have a contiguous range of vector entries stored on each MPI rank. The first rank has entries from 0 to \( r_{\text{end}1} - 1 \), the next rank has entries from \( r_{\text{end}1} \) to \( r_{\text{end}2} - 1 \), etc. Thus the ownership range on each rank is from \( r_{\text{start}} \) to \( r_{\text{end}} \), these values can be obtained with VecGetOwnershipRange(Vec \( x \), PetscInt * \( r_{\text{start}} \), PetscInt * \( r_{\text{end}} \)). Each PETSc Vec has a PetscLayout object that contains this information.

All PETSc matrices have two PetscLayouts, they define the vector layouts for \( y \) and \( x \) in the product, \( y = A \times x \). Their ownership range information can be obtained with MatGetOwnershipRange(), MatGetOwnershipRangeColumn(), MatGetOwnershipRanges(), and MatGetOwnershipRangesColumn(). Note that MatCreateVecs() provides two vectors that have compatible layouts for the associated vector.

For most PETSc matrices, excluding MATELEMENTAL and MATSCALAPACK, the row ownership range obtained with MatGetOwnershipRange() also defines where the matrix entries are stored; the matrix entries for rows \( r_{\text{start}} \) to \( r_{\text{end}} - 1 \) are stored on the corresponding MPI rank. For other matrices the rank where each matrix entry is stored is more complicated; information about the storage locations can be obtained with MatGetOwnershipIS(). Note that for most PETSc matrices the values returned by MatGetOwnershipIS() are the same as those returned by MatGetOwnershipRange() and MatGetOwnershipRangeColumn().

The PETSc object PetscLayout contains the ownership information that is provided by VecGetOwnershipRange() and with MatGetOwnershipRange(), MatGetOwnershipRangeColumn(). Each vector has one layout, which can be obtained with VecGetLayout() and MatGetLayouts(). Layouts support the routines PetscLayoutGetLocalSize(), PetscLayoutGetSize(), PetscLayoutGetBlockSize(), PetscLayoutGetRanges(), PetscLayoutCompare() as well as a variety of creation routines. These are used by the Vec and Mat and so are rarely needed directly. Finally PetscSplitOwnership() is a utility routine that does the same splitting of ownership ranges as PetscLayout.

Sparse Matrices

The default matrix representation within PETSc is the general sparse AIJ format (also called the compressed sparse row format, CSR). This section discusses tips for efficiently using this matrix format for large-scale applications. Additional formats (such as block compressed row and block symmetric storage, which are generally much more efficient for problems with multiple degrees of freedom per node) are discussed below. Beginning users need not concern themselves initially with such details and may wish to proceed directly to Basic Matrix Operations. However, when an application code progresses to the point of tuning for efficiency and/or generating timing results, it is crucial to read this information.
Sequential AIJ Sparse Matrices

In the PETSc AIJ matrix formats, we store the nonzero elements by rows, along with an array of corresponding column numbers and an array of pointers to the beginning of each row. Note that the diagonal matrix entries are stored with the rest of the nonzeros (not separately).

To create a sequential AIJ sparse matrix, \(A\), with \(m\) rows and \(n\) columns, one uses the command

\[
\text{MatCreateSeqAIJ(PETSC_COMM_SELF, PetscInt } m, \text{PetscInt } n, \text{PetscInt } nz, \text{PetscInt } *nnz, \text{Mat } *A); \]

where \(nz\) or \(nnz\) can be used to preallocate matrix memory, as discussed below. The user can set \(nz=0\) and \(nnz=NULL\) for PETSc to control all matrix memory allocation.

The sequential and parallel AIJ matrix storage formats by default employ i-nodes (identical nodes) when possible. We search for consecutive rows with the same nonzero structure, thereby reusing matrix information for increased efficiency. Related options database keys are \(-\text{mat_no_inode}\) (do not use i-nodes) and \(-\text{mat_inode_limit } <\text{limit}>\) (set i-node limit (max limit=5)). Note that problems with a single degree of freedom per grid node will automatically not use i-nodes.

The internal data representation for the AIJ formats employs zero-based indexing.

Preallocation of Memory for Sequential AIJ Sparse Matrices

The dynamic process of allocating new memory and copying from the old storage to the new is intrinsically very expensive. Thus, to obtain good performance when assembling an AIJ matrix, it is crucial to preallocate the memory needed for the sparse matrix. The user has two choices for preallocating matrix memory via \(\text{MatCreateSeqAIJ()}\).

One can use the scalar \(nz\) to specify the expected number of nonzeros for each row. This is generally fine if the number of nonzeros per row is roughly the same throughout the matrix (or as a quick and easy first step for preallocation). If one underestimates the actual number of nonzeros in a given row, then during the assembly process PETSc will automatically allocate additional needed space. However, this extra memory allocation can slow the computation.

If different rows have very different numbers of nonzeros, one should attempt to indicate (nearly) the exact number of elements intended for the various rows with the optional array, \(nnz\) of length \(m\), where \(m\) is the number of rows, for example

\[
\text{PetscInt } nnz[m]; \nnz[0] = \text{nonzeros in row 0} \nnz[1] = \text{nonzeros in row 1} \nnz[m-1] = \text{nonzeros in row m-1} \]

In this case, the assembly process will require no additional memory allocations if the \(nnz\) estimates are correct. If, however, the \(nnz\) estimates are incorrect, PETSc will automatically obtain the additional needed space, at a slight loss of efficiency.

Using the array \(nnz\) to preallocate memory is especially important for efficient matrix assembly if the number of nonzeros varies considerably among the rows. One can generally set \(nnz\) either by knowing in advance the problem structure (e.g., the stencil for finite difference problems on a structured grid) or by precomputing the information by using a segment of code similar to that for the regular matrix assembly. The overhead of determining the \(nnz\) array will be quite small compared with the overhead of the inherently expensive \(\text{malloc}\)s and moves of data that are needed for dynamic allocation during matrix assembly. Always guess high if an exact value is not known (extra space is cheaper than too little).
Thus, when assembling a sparse matrix with very different numbers of nonzeros in various rows, one could proceed as follows for finite difference methods:

1. Allocate integer array \( \text{nnz} \).
2. Loop over grid, counting the expected number of nonzeros for the row(s) associated with the various grid points.
3. Create the sparse matrix via \texttt{MatCreateSeqAIJ() or alternative}. 
4. Loop over the grid, generating matrix entries and inserting in matrix via \texttt{MatSetValues()}. 

For (vertex-based) finite element type calculations, an analogous procedure is as follows:

1. Allocate integer array \( \text{nnz} \).
2. Loop over vertices, computing the number of neighbor vertices, which determines the number of nonzeros for the corresponding matrix row(s).
3. Create the sparse matrix via \texttt{MatCreateSeqAIJ()} or alternative.
4. Loop over elements, generating matrix entries and inserting in matrix via \texttt{MatSetValues()}. 

The \texttt{-info} option causes the routines \texttt{MatAssemblyBegin()} and \texttt{MatAssemblyEnd()} to print information about the success of the preallocation. Consider the following example for the \texttt{MATSEQAIJ} matrix format:

\begin{verbatim}
MatAssemblyEnd_SeqAIJ:Matrix size 10 X 10; storage space:20 unneeded, 100 used
MatAssemblyEnd_SeqAIJ:Number of mallocs during MatSetValues is 0
\end{verbatim}

The first line indicates that the user preallocated 120 spaces but only 100 were used. The second line indicates that the user preallocated enough space so that PETSc did not have to internally allocate additional space (an expensive operation). In the next example the user did not preallocate sufficient space, as indicated by the fact that the number of mallocs is very large (bad for efficiency):

\begin{verbatim}
MatAssemblyEnd_SeqAIJ:Matrix size 10 X 10; storage space:47 unneeded, 1000 used
MatAssemblyEnd_SeqAIJ:Number of mallocs during MatSetValues is 40000
\end{verbatim}

Although at first glance such procedures for determining the matrix structure in advance may seem unusual, they are actually very efficient because they alleviate the need for dynamic construction of the matrix data structure, which can be very expensive.

**Parallel AIJ Sparse Matrices**

Parallel sparse matrices with the AIJ format can be created with the command

\begin{verbatim}
MatCreateAIJ(MPI_Comm comm,PetscInt m,PetscInt n,PetscInt M,PetscInt N,PetscInt d_nz, PetscInt *d_nnz, PetscInt o_nz,PetscInt *o_nnz,Mat *A);
\end{verbatim}

\( A \) is the newly created matrix, while the arguments \( m, M, \) and \( N, \) indicate the number of local rows and the number of global rows and columns, respectively. In the PETSc partitioning scheme, all the matrix columns are local and \( n \) is the number of columns corresponding to the local part of a parallel vector. Either the local or global parameters can be replaced with \texttt{PETSC_DECIDE}, so that PETSc will determine them. The matrix is stored with a fixed number of rows on each process, given by \( m \), or determined by PETSc if \( m \) is \texttt{PETSC_DECIDE}.

If \texttt{PETSC_DECIDE} is not used for the arguments \( m \) and \( n \), then the user must ensure that they are chosen to be compatible with the vectors. To do this, one first considers the matrix-vector product \( y = Ax \). The \( m \) that is used in the matrix creation routine \texttt{MatCreateAIJ()} must match the local size used in the vector
creation routine VecCreateMPI() for \( y \). Likewise, the \( n \) used must match that used as the local size in VecCreateMPI() for \( x \).

The user must set \( d_{nz}=0, o_{nz}=0, d_{nnz}={\text{NULL}}, \) and \( o_{nnz}={\text{NULL}} \) for PETSc to control dynamic allocation of matrix memory space. Analogous to \( nz \) and \( nnz \) for the routine MatCreateSeqAIJ(), these arguments optionally specify nonzero information for the diagonal (\( d_{nz} \) and \( d_{nnz} \)) and off-diagonal (\( o_{nz} \) and \( o_{nnz} \)) parts of the matrix. For a square global matrix, we define each process’s diagonal portion to be its local rows and the corresponding columns (a square submatrix); each process’s off-diagonal portion encompasses the remainder of the local matrix (a rectangular submatrix). The rank in the MPI communicator determines the absolute ordering of the blocks. That is, the process with rank 0 in the communicator given to MatCreateAIJ() contains the top rows of the matrix; the \( i^{th} \) process in that communicator contains the \( i^{th} \) block of the matrix.

### Preallocation of Memory for Parallel AIJ Sparse Matrices

As discussed above, preallocation of memory is critical for achieving good performance during matrix assembly, as this reduces the number of allocations and copies required. We present an example for three processes to indicate how this may be done for the MATMPIAIJ matrix format. Consider the 8 by 8 matrix, which is partitioned by default with three rows on the first process, three on the second and two on the third.

\[
\begin{pmatrix}
1 & 2 & 0 & | & 0 & 3 & 0 & | & 0 & 4 \\
0 & 5 & 6 & | & 7 & 0 & 0 & | & 8 & 0 \\
9 & 0 & 10 & | & 11 & 0 & 0 & | & 12 & 0 \\
13 & 0 & 14 & | & 15 & 16 & 17 & | & 0 & 0 \\
0 & 18 & 0 & | & 19 & 20 & 21 & | & 0 & 0 \\
0 & 0 & 0 & | & 22 & 23 & 0 & | & 24 & 0 \\
25 & 26 & 27 & | & 0 & 0 & 28 & | & 29 & 0 \\
30 & 0 & 0 & | & 31 & 32 & 33 & | & 0 & 34
\end{pmatrix}
\]

The “diagonal” submatrix, \( d \), on the first process is given by

\[
\begin{pmatrix}
1 & 2 & 0 \\
0 & 5 & 6 \\
9 & 0 & 10
\end{pmatrix},
\]

while the “off-diagonal” submatrix, \( o \), matrix is given by

\[
\begin{pmatrix}
0 & 0 & 0 & 4 \\
7 & 0 & 0 & 8 & 0 \\
11 & 0 & 0 & 12 & 0
\end{pmatrix}.
\]

For the first process one could set \( d_{nz} \) to 2 (since each row has 2 nonzeros) or, alternatively, set \( d_{nnz} \) to \{2, 2, 2\}. The \( o_{nz} \) could be set to 2 since each row of the \( o \) matrix has 2 nonzeros, or \( o_{nnz} \) could be set to \{2, 2, 2\}.

For the second process the \( d \) submatrix is given by

\[
\begin{pmatrix}
15 & 16 & 17 \\
19 & 20 & 21 \\
22 & 23 & 0
\end{pmatrix}.
\]

Thus, one could set \( d_{nz} \) to 3, since the maximum number of nonzeros in each row is 3, or alternatively one could set \( d_{nnz} \) to \{3, 3, 2\}, thereby indicating that the first two rows will have 3 nonzeros while the third

2.2. Matrices
has 2. The corresponding o submatrix for the second process is

\[
\begin{pmatrix}
13 & 0 & 14 & 0 & 0 \\
0 & 18 & 0 & 0 & 0 \\
0 & 0 & 0 & 24 & 0
\end{pmatrix}
\]

so that one could set \( o_{nz} \) to 2 or \( o_{nnz} \) to \( \{2,1\} \).

Note that the user never directly works with the d and o submatrices, except when preallocating storage space as indicated above. Also, the user need not preallocate exactly the correct amount of space; as long as a sufficiently close estimate is given, the high efficiency for matrix assembly will remain.

As described above, the option -info will print information about the success of preallocation during matrix assembly. For the MATMPIAIJ and MATMPIBAIJ formats, PETSc will also list the number of elements owned by on each process that were generated on a different process. For example, the statements

\[
\begin{array}{l}
\text{MatAssemblyBegin\_MPIAIJ: Stash has 10 entries, uses 0 mallocs} \\
\text{MatAssemblyBegin\_MPIAIJ: Stash has 3 entries, uses 0 mallocs} \\
\text{MatAssemblyBegin\_MPIAIJ: Stash has 5 entries, uses 0 mallocs}
\end{array}
\]

indicate that very few values have been generated on different processes. On the other hand, the statements

\[
\begin{array}{l}
\text{MatAssemblyBegin\_MPIAIJ: Stash has 100000 entries, uses 100 mallocs} \\
\text{MatAssemblyBegin\_MPIAIJ: Stash has 77777 entries, uses 70 mallocs}
\end{array}
\]

indicate that many values have been generated on the “wrong” processes. This situation can be very inefficient, since the transfer of values to the “correct” process is generally expensive. By using the command MatGetOwnershipRange() in application codes, the user should be able to generate most entries on the owning process.

*Note:* It is fine to generate some entries on the “wrong” process. Often this can lead to cleaner, simpler, less buggy codes. One should never make code overly complicated in order to generate all values locally. Rather, one should organize the code in such a way that most values are generated locally.

The routine MatCreateAIJCUSPARSE() allows one to create GPU based matrices for NVIDIA systems. MatCreateAIJKokkos() can create matrices for use with CPU, OpenMP, NVIDIA, AMD, or Intel based GPU systems.

It is sometimes difficult to compute the required preallocation information efficiently, hence PETSc provides a special MatType, MATPREALLOCATOR that helps make computing this information more straightforward. One first creates a matrix of this type and then, using the same code that one would use to actually compute the matrices numerical values, calls MatSetValues() for this matrix, without needing to provide any preallocation information (one need not provide the matrix numerical values). Once this is complete one uses MatPreallocatorPreallocate() to provide the accumulated preallocation information to the actual matrix one will use for the computations. We hope to simplify this process in the future, allowing the removal of MATPREALLOCATOR, instead simply allowing the use of its efficient insertion process automatically during the first assembly of any matrix type directly without requiring the detailed preallocation information.

See doc\_matrix for a table of the matrix types available in PETSc.
Limited-Memory Variable Metric (LMVM) Matrices

Variable metric methods, also known as quasi-Newton methods, are frequently used for root finding problems and approximate Jacobian matrices or their inverses via sequential nonlinear updates based on the secant condition. The limited-memory variants do not store the full explicit Jacobian, and instead compute forward products and inverse applications based on a fixed number of stored update vectors.

Table 2.2: PETSc LMVM matrix implementations.

<table>
<thead>
<tr>
<th>Method</th>
<th>PETSc Type</th>
<th>Name</th>
<th>Property</th>
</tr>
</thead>
<tbody>
<tr>
<td>“Good” Broyden [ref-Gri12]</td>
<td>MATLMVMBrdn</td>
<td>lmvmbrdn</td>
<td>Square</td>
</tr>
<tr>
<td>“Bad” Broyden [ref-Gri12]</td>
<td>MATLMVMBad-Brdn</td>
<td>lmvmbadbrdn</td>
<td>Square</td>
</tr>
<tr>
<td>Symmetric Rank-1 [ref-NW06]</td>
<td>MATLMVMSR1</td>
<td>lmvmsr1</td>
<td>Symmetric</td>
</tr>
<tr>
<td>Davidon-Fletcher-Powell (DFP) [ref-NW06]</td>
<td>MATLMVMDFP</td>
<td>lmvmdfp</td>
<td>SPD</td>
</tr>
<tr>
<td>Broyden-Fletcher-Goldfarb-Shanno (BFGS) [ref-NW06]</td>
<td>MATLMVMBFGS</td>
<td>lmvmbfgs</td>
<td>SPD</td>
</tr>
<tr>
<td>Restricted Broyden Family [ref-EM17]</td>
<td>MATLMVMSym-Brdn</td>
<td>lmvmsymbrdn</td>
<td>SPD</td>
</tr>
<tr>
<td>Restricted Broyden Family (full-memory diagonal)</td>
<td>MATLMVMDiag-Brdn</td>
<td>lmvmdiagbrdn</td>
<td>SPD</td>
</tr>
</tbody>
</table>

PETSc implements seven different LMVM matrices listed in the table above. They can be created using the MatCreate() and MatSetType() workflow, and share a number of common interface functions. We will review the most important ones below:

- **MatLMVMAllocate(Mat B, Vec X, Vec F)** – Creates the internal data structures necessary to store nonlinear updates and compute forward/inverse applications. The X vector defines the solution space while the F defines the function space for the history of updates.

- **MatLMVMUpdate(Mat B, Vec X, Vec F)** – Applies a nonlinear update to the approximate Jacobian such that \( s_k = x_k - x_{k-1} \) and \( y_k = f(x_k) - f(x_{k-1}) \), where \( k \) is the index for the update.

- **MatLMVMReset(Mat B, PetscBool destructive)** – Flushes the accumulated nonlinear updates and resets the matrix to the initial state. If destructive = PETSC_TRUE, the reset also destroys the internal data structures and necessitates another allocation call before the matrix can be updated and used for products and solves.

- **MatLMVMSetJ0(Mat B, Mat J0)** – Defines the initial Jacobian to apply the updates to. If no initial Jacobian is provided, the updates are applied to an identity matrix.

LMVM matrices can be applied to vectors in forward mode via MatMult() or MatMultAdd(), and in inverse mode via MatSolve(). They also support MatCreateVecs(), MatDuplicate() and MatCopy() operations.

Restricted Broyden Family, DFP and BFGS methods additionally implement special Jacobian initialization and scaling options available via -mat_lmvm_scale_type <none, scalar, diagonal>. We describe these choices below:

- **none** – Sets the initial Jacobian to be equal to the identity matrix. No extra computations are required when obtaining the search direction or updating the approximation. However, the number of function evaluations required to converge the Newton solution is typically much larger than what is required when using other initializations.

- **scalar** – Defines the initial Jacobian as a scalar multiple of the identity matrix. The scalar value \( \sigma \)
is chosen by solving the one dimensional optimization problem

$$\min_\sigma \| \sigma^\alpha Y - \sigma^{\alpha-1} S \|^2_F,$$

where $S$ and $Y$ are the matrices whose columns contain a subset of update vectors $s_k$ and $y_k$, and $\alpha \in [0, 1]$ is defined by the user via `-mat_lmvm_alpha` and has a different default value for each LMVM implementation (e.g.: default $\alpha = 1$ for BFGS produces the well-known $y_k^T s_k / y_k^T y_k$ scalar initialization). The number of updates to be used in the $S$ and $Y$ matrices is 1 by default (i.e.: the latest update only) and can be changed via `-mat_lmvm_sigma_hist`. This technique is inspired by Gilbert and Lemarechal [ref-GL89].

- **diagonal** – Uses a full-memory restricted Broyden update formula to construct a diagonal matrix for the Jacobian initialization. Although the full-memory formula is utilized, the actual memory footprint is restricted to only the vector representing the diagonal and some additional work vectors used in its construction. The diagonal terms are also re-scaled with every update as suggested in [ref-GL89]. This initialization requires the most computational effort of the available choices but typically results in a significant reduction in the number of function evaluations taken to compute a solution.

Note that the user-provided initial Jacobian via `MatLMVMSetJ0()` overrides and disables all built-in initialization methods.

### Dense Matrices

PETSc provides both sequential and parallel dense matrix formats, where each process stores its entries in a column-major array in the usual Fortran style. To create a sequential, dense PETSc matrix, $A$ of dimensions $m$ by $n$, the user should call

```c
MatCreateSeqDense(PETSC_COMM_SELF,PetscInt m,PetscInt n,PetscScalar *data,Mat *A);
```

The variable `data` enables the user to optionally provide the location of the data for matrix storage (intended for Fortran users who wish to allocate their own storage space). Most users should merely set `data` to NULL for PETSc to control matrix memory allocation. To create a parallel, dense matrix, $A$, the user should call

```c
MatCreateDense(MPI_Comm comm,PetscInt m,PetscInt n,PetscInt M,PetscInt N,PetscScalar *data,Mat *A);
```

The arguments $m$, $n$, $M$, and $N$, indicate the number of local rows and columns and the number of global rows and columns, respectively. Either the local or global parameters can be replaced with `PETSC_DECIDE`, so that PETSc will determine them. The matrix is stored with a fixed number of rows on each process, given by $m$, or determined by PETSc if $m$ is `PETSC_DECIDE`.

PETSc does not provide parallel dense direct solvers, instead interfacing to external packages that provide these solvers. Our focus is on sparse iterative solvers.

### Block Matrices

Block matrices arise when coupling variables with different meaning, especially when solving problems with constraints (e.g. incompressible flow) and “multi-physics” problems. Usually the number of blocks is small and each block is partitioned in parallel. We illustrate for a $3 \times 3$ system with components labeled $a, b, c$. With some numbering of unknowns, the matrix could be written as

$$
\begin{pmatrix}
A_{aa} & A_{ab} & A_{ac} \\
A_{ba} & A_{bb} & A_{bc} \\
A_{ca} & A_{cb} & A_{cc}
\end{pmatrix}.
$$
There are two fundamentally different ways that this matrix could be stored, as a single assembled sparse matrix where entries from all blocks are merged together (“monolithic”), or as separate assembled matrices for each block (“nested”). These formats have different performance characteristics depending on the operation being performed. In particular, many preconditioners require a monolithic format, but some that are very effective for solving block systems (see Solving Block Matrices) are more efficient when a nested format is used. In order to stay flexible, we would like to be able to use the same code to assemble block matrices in both monolithic and nested formats. Additionally, for software maintainability and testing, especially in a multi-physics context where different groups might be responsible for assembling each of the blocks, it is desirable to be able to use exactly the same code to assemble a single block independently as to assemble it as part of a larger system. To do this, we introduce the four spaces shown in Fig. 2.3.

- The monolithic global space is the space in which the Krylov and Newton solvers operate, with collective semantics across the entire block system.
- The split global space splits the blocks apart, but each split still has collective semantics.
- The split local space adds ghost points and separates the blocks. Operations in this space can be performed with no parallel communication. This is often the most natural, and certainly the most powerful, space for matrix assembly code.
- The monolithic local space can be thought of as adding ghost points to the monolithic global space, but it is often more natural to use it simply as a concatenation of split local spaces on each process. It is not common to explicitly manipulate vectors or matrices in this space (at least not during assembly), but it is a useful for declaring which part of a matrix is being assembled.

![Fig. 2.3: The relationship between spaces used for coupled assembly.](image)

The key to format-independent assembly is the function

**2.2. Matrices**
MatGetLocalSubMatrix(Mat A, IS isrow, IS iscol, Mat *submat);

which provides a “view” submat into a matrix A that operates in the monolithic global space. The submat transforms from the split local space defined by iscol to the split local space defined by isrow. The index sets specify the parts of the monolithic local space that submat should operate on. If a nested matrix format is used, then MatGetLocalSubMatrix() finds the nested block and returns it without making any copies. In this case, submat is fully functional and has a parallel communicator. If a monolithic matrix format is used, then MatGetLocalSubMatrix() returns a proxy matrix on PETSC_COMM_SELF that does not provide values or implement MatMult(), but does implement MatSetValuesLocal() and, if isrow, iscol have a constant block size, MatSetValuesBlockedLocal(). Note that although submat may not be a fully functional matrix and the caller does not even know a priori which communicator it will reside on, it always implements the local assembly functions (which are not collective). The index sets isrow, iscol can be obtained using DMCompositeGetLocalISs() if DMCOMPOSITE is being used. DMCOMPOSITE can also be used to create matrices, in which case the MATNEST format can be specified using -prefix_dm_mat_type nest and MATAIJ can be specified using -prefix_dm_mat_type aij. See SNES Tutorial ex28 for a simple example using this interface.

### 2.2.4 Basic Matrix Operations

Table 2.2 summarizes basic PETSc matrix operations. We briefly discuss a few of these routines in more detail below.

The parallel matrix can multiply a vector with n local entries, returning a vector with m local entries. That is, to form the product

\[
\text{MatMult(Mat A, Vec x, Vec y);}
\]

the vectors x and y should be generated with

\[
\text{VecCreateMPI(MPI_Comm comm, n, N, &x);} \\
\text{VecCreateMPI(MPI_Comm comm, m, M, &y);} \\
\]

By default, if the user lets PETSc decide the number of components to be stored locally (by passing in PETSC_DECIDE as the second argument to VecCreateMPI() or using VecCreate()), vectors and matrices of the same dimension are automatically compatible for parallel matrix-vector operations.

Along with the matrix-vector multiplication routine, there is a version for the transpose of the matrix,

\[
\text{MatMultTranspose(Mat A, Vec x, Vec y);} \\
\]

There are also versions that add the result to another vector:

\[
\text{MatMultAdd(Mat A, Vec x, Vec y, Vec w);} \\
\text{MatMultTransposeAdd(Mat A, Vec x, Vec y, Vec w);} \\
\]

These routines, respectively, produce \( w = A \times x + y \) and \( w = A^T \times x + y \). In C it is legal for the vectors y and w to be identical. In Fortran, this situation is forbidden by the language standard, but we allow it anyway.

One can print a matrix (sequential or parallel) to the screen with the command

\[
\text{MatView(Mat mat, PETSC_VIEWER_STDOUT_WORLD);} \\
\]

Other viewers can be used as well. For instance, one can draw the nonzero structure of the matrix into the default X-window with the command.
MatView(Mat mat, PETSC_VIEWER_DRAW_WORLD);

Also one can use

MatView(Mat mat, PetscViewer viewer);

where viewer was obtained with PetscViewerDrawOpen(). Additional viewers and options are given in the MatView() man page and Viewers: Looking at PETSc Objects.

Table 2.3: PETSc Matrix Operations

<table>
<thead>
<tr>
<th>Function Name</th>
<th>Operation</th>
</tr>
</thead>
<tbody>
<tr>
<td>MatAXPY(Mat Y, PetscScalar a, Mat X, MatStructure s);</td>
<td>$Y = Y + a \times X$</td>
</tr>
<tr>
<td>MatAYPX(Mat Y, PetscScalar a, Mat X, MatStructure s);</td>
<td>$Y = a \times Y + X$</td>
</tr>
<tr>
<td>MatMult(Mat A, Vec x, Vec y);</td>
<td>$y = A \times x$</td>
</tr>
<tr>
<td>MatMultAdd(Mat A, Vec x, Vec y, Vec z);</td>
<td>$z = y + A \times x$</td>
</tr>
<tr>
<td>MatMultTranspose(Mat A, Vec x, Vec y);</td>
<td>$y = A^T \times x$</td>
</tr>
<tr>
<td>MatMultTransposeAdd(Mat A, Vec x, Vec y, Vec z);</td>
<td>$z = y + A^T \times x$</td>
</tr>
<tr>
<td>MatNorm(Mat A, NormType type, PetscReal *r);</td>
<td>$r = A_{\text{type}}$</td>
</tr>
<tr>
<td>MatDiagonalScale(Mat A, Vec l, Vec r);</td>
<td>$A = \text{diag}(l) \times A \times \text{diag}(r)$</td>
</tr>
<tr>
<td>MatScale(Mat A, PetscScalar a)</td>
<td>$A = a \times A$</td>
</tr>
<tr>
<td>MatConvert(Mat A, MatType type, Mat *B);</td>
<td>$B = A$</td>
</tr>
<tr>
<td>MatCopy(Mat A, Mat B, MatStructure s);</td>
<td>$B = A$</td>
</tr>
<tr>
<td>MatGetDiagonal(Mat A, Vec x);</td>
<td>$x = \text{diag}(A)$</td>
</tr>
<tr>
<td>MatTranspose(Mat A, MatReuse, Mat * B);</td>
<td>$B = A^T$</td>
</tr>
<tr>
<td>MatZeroEntries(Mat A);</td>
<td>$A = 0$</td>
</tr>
<tr>
<td>MatShift(Mat Y, PetscScalar a);</td>
<td>$Y = Y + a \times I$</td>
</tr>
</tbody>
</table>

Table 2.4: Values of MatStructure

<table>
<thead>
<tr>
<th>Name</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>SAME_NONZERO_PATTERN</td>
<td>the matrices have an identical nonzero pattern</td>
</tr>
<tr>
<td>DIFFERENT_NONZERO_PATTERN</td>
<td>the matrices may have a different nonzero pattern</td>
</tr>
<tr>
<td>SUBSET_NONZERO_PATTERN</td>
<td>the second matrix has a subset of the nonzeros in the first matrix</td>
</tr>
<tr>
<td>UNKNOWN_NONZERO_PATTERN</td>
<td>there is nothing known about the relation between the nonzero patterns of the two matrices</td>
</tr>
</tbody>
</table>

The NormType argument to MatNorm() is one of NORM_1, NORM_INFINITY, and NORM_FROBENIUS.

### 2.2.5 Matrix-Free Matrices

Some people like to use matrix-free methods, which do not require explicit storage of the matrix, for the numerical solution of partial differential equations. To support matrix-free methods in PETSc, one can use the following command to create a Mat structure without ever actually generating the matrix:

```c
MatCreateShell(MPI_Comm comm, PetscInt m, PetscInt n, PetscInt M, PetscInt N, void *ctx, ...Mat *mat);
```

Here M and N are the global matrix dimensions (rows and columns), m and n are the local matrix dimensions, and ctx is a pointer to data needed by any user-defined shell matrix operations; the manual page has...
additional details about these parameters. Most matrix-free algorithms require only the application of the linear operator to a vector. To provide this action, the user must write a routine with the calling sequence

```c
UserMult(Mat mat, Vec x, Vec y);
```

and then associate it with the matrix, `mat`, by using the command

```c
MatShellSetOperation(Mat mat, MatOperation MATOP_MULTI, (void (*)(void)) PetscErrorCode
                   (*UserMult)(Mat, Vec, Vec));
```

Here `MATOP_MULTI` is the name of the operation for matrix-vector multiplication. Within each user-defined routine (such as `UserMult()`), the user should call `MatShellGetContext()` to obtain the user-defined context, `ctx`, that was set by `MatCreateShell()`. This shell matrix can be used with the iterative linear equation solvers discussed in the following chapters.

The routine `MatShellSetOperation()` can be used to set any other matrix operations as well. The file `$PETSC_DIR/include/petscmat.h` (source) provides a complete list of matrix operations, which have the form `MATOP_<OPERATION>`, where `<OPERATION>` is the name (in all capital letters) of the user interface routine (for example, `MatMult()` → `MATOP_MULTI`). All user-provided functions have the same calling sequence as the usual interface routines, since the user-defined functions are intended to be accessed through the same interface, e.g., `MatMult(Mat, Vec, Vec) → UserMult(Mat, Vec, Vec)`. The final argument for `MatShellSetOperation()` needs to be cast to a `void *`, since the final argument could (depending on the `MatOperation`) be a variety of different functions.

Note that `MatShellSetOperation()` can also be used as a “backdoor” means of introducing user-defined changes in matrix operations for other storage formats (for example, to override the default LU factorization routine supplied within PETSc for the `MATSEQAIJ` format). However, we urge anyone who introduces such changes to use caution, since it would be very easy to accidentally create a bug in the new routine that could affect other routines as well.

See also *Matrix-Free Methods* for details on one set of helpful utilities for using the matrix-free approach for nonlinear solvers.

### 2.2.6 Transposes of Matrices

PETSc provides several ways to work with transposes of matrix.

```c
MatTranspose(Mat A, MatReuse MAT_INITIAL_MATRIX or MAT_INPLACE_MATRIX or MAT_REUSE_MATRIX, Mat *B)
```

will either do an in-place or out-of-place matrix explicit formation of the matrix transpose. After it has been called with `MAT_INPLACE_MATRIX` it may be called again with `MAT_REUSE_MATRIX` and it will recompute the transpose if the A matrix has changed. Internally it keeps track of whether the nonzero pattern of A has not changed so will reuse the symbolic transpose when possible for efficiency.

```c
MatTransposeSymbolic(Mat A, Mat *B)
```

only does the symbolic transpose on the matrix. After it is called `MatTranspose()` may be called with `MAT_REUSE_MATRIX` to compute the numerical transpose.

Occasionally one may already have a B matrix with the needed sparsity pattern to store the transpose and wants to reuse that space instead of creating a new matrix by calling `MatTranspose(A,"MAT_INITIAL_MATRIX",&B)` but they cannot just call `MatTranspose(A,"MAT_REUSE_MATRIX",&B)` so instead they can call `MatTransposeSetPrecursor(A,B)` and then call `MatTranspose(A,"MAT_REUSE_MATRIX",&B)`. This routine just provides to B the meta-data it needs to compute the numerical factorization efficiently.
The routine `MatCreateTranspose(A,&B)` provides a surrogate matrix B that behaviors like the transpose of A without forming the transpose explicitly. For example, `MatMult(B,x,y)` will compute the matrix-vector product of A transpose times x.

### 2.2.7 Other Matrix Operations

In many iterative calculations (for instance, in a nonlinear equations solver), it is important for efficiency purposes to reuse the nonzero structure of a matrix, rather than determining it anew every time the matrix is generated. To retain a given matrix but reinitialize its contents, one can employ

```
MatZeroEntries(Mat A);
```

This routine will zero the matrix entries in the data structure but keep all the data that indicates where the nonzeros are located. In this way a new matrix assembly will be much less expensive, since no memory allocations or copies will be needed. Of course, one can also explicitly set selected matrix elements to zero by calling `MatSetValues()`.

By default, if new entries are made in locations where no nonzeros previously existed, space will be allocated for the new entries. To prevent the allocation of additional memory and simply discard those new entries, one can use the option

```
MatSetOption(Mat A, MAT_NEW_NONZERO_LOCATIONS, PETSC_FALSE);
```

Once the matrix has been assembled, one can factor it numerically without repeating the ordering or the symbolic factorization. This option can save some computational time, although it does require that the factorization is not done in-place.

In the numerical solution of elliptic partial differential equations, it can be cumbersome to deal with Dirichlet boundary conditions. In particular, one would like to assemble the matrix without regard to boundary conditions and then at the end apply the Dirichlet boundary conditions. In numerical analysis classes this process is usually presented as moving the known boundary conditions to the right-hand side and then solving a smaller linear system for the interior unknowns. Unfortunately, implementing this requires extracting a large submatrix from the original matrix and creating its corresponding data structures. This process can be expensive in terms of both time and memory.

One simple way to deal with this difficulty is to replace those rows in the matrix associated with known boundary conditions, by rows of the identity matrix (or some scaling of it). This action can be done with the command

```
MatZeroRows(Mat A, PetscInt numRows, PetscInt rows[], PetscScalar diag_value, Vec x, Vec b),
```

or equivalently,

```
MatZeroRowsIS(Mat A, IS rows, PetscScalar diag_value, Vec x, Vec b);
```

For sparse matrices this removes the data structures for certain rows of the matrix. If the pointer `diag_value` is NULL, it even removes the diagonal entry. If the pointer is not null, it uses that given value at the pointer location in the diagonal entry of the eliminated rows.

One nice feature of this approach is that when solving a nonlinear problem such that at each iteration the Dirichlet boundary conditions are in the same positions and the matrix retains the same nonzero structure, the user can call `MatZeroRows()` in the first iteration. Then, before generating the matrix in the second iteration the user should call
MatSetOption(Mat A, MAT_NEW_NONZERO_LOCATIONS, PETSC_FALSE);

From that point, no new values will be inserted into those (boundary) rows of the matrix.

The functions MatZeroRowsLocal() and MatZeroRowsLocalIS() can also be used if for each process one provides the Dirichlet locations in the local numbering of the matrix. A drawback of MatZeroRows() is that it destroys the symmetry of a matrix. Thus one can use

MatZeroRowsColumns(Mat A, PetscInt numRows, PetscInt rows[], PetscScalar diag_value, Vec x, Vec b),

or equivalently,

MatZeroRowsColumnsIS(Mat A, IS rows, PetscScalar diag_value, Vec x, Vec b);

Note that with all of these for a given assembled matrix it can be only called once to update the x and b vector. It cannot be used if one wishes to solve multiple right hand side problems for the same matrix since the matrix entries needed for updating the b vector are removed in its first use.

Once the zeroed rows are removed the new matrix has possibly many rows with only a diagonal entry affecting the parallel load balancing. The PCREDISTRIBUTE preconditioner removes all the zeroed rows (and associated columns and adjusts the right hand side based on the removed columns) and then rebalances the resulting rows of smaller matrix across the processes. Thus one can use MatZeroRows() to set the Dirichlet points and then solve with the preconditioner PCREDISTRIBUTE. Note if the original matrix was symmetric the smaller solved matrix will also be symmetric.

Another matrix routine of interest is

MatConvert(Mat mat, MatType newtype, Mat *M)

which converts the matrix mat to new matrix, M, that has either the same or different format. Set newtype to MATSAME to copy the matrix, keeping the same matrix format. See $PETSC_DIR/include/petscmat.h (source) for other available matrix types; standard ones are MATSEQDENSE, MATSEQAIJ, MATMPIAIJ, MATSEQBAIJ and MATMPIBAIJ.

In certain applications it may be necessary for application codes to directly access elements of a matrix. This may be done by using the the command (for local rows only)

MatGetRow(Mat A, PetscInt row, PetscInt *ncols, const PetscInt (*cols)[], const PetscScalar (*vals)[]);

The argument ncols returns the number of nonzeros in that row, while cols and vals returns the column indices (with indices starting at zero) and values in the row. If only the column indices are needed (and not the corresponding matrix elements), one can use NULL for the vals argument. Similarly, one can use NULL for the cols argument. The user can only examine the values extracted with MatGetRow(); the values cannot be altered. To change the matrix entries, one must use MatSetValues().

Once the user has finished using a row, he or she must call

MatRestoreRow(Mat A, PetscInt row, PetscInt *ncols, PetscInt **cols, PetscScalar **vals);

to free any space that was allocated during the call to MatGetRow().
2.2.8 Symbolic and Numeric Stages in Sparse Matrix Operations

Many sparse matrix operations can be optimized by dividing the computation into two stages: a symbolic stage that creates any required data structures and does all the computations that do not require the matrices’ numerical values followed by one or more uses of a numerical stage that use the symbolically computed information. Examples of such operations include `MatTranspose()`, `MatCreateSubMatrices()`, `MatCholeskyFactorSymbolic()`, and `MatCholeskyFactorNumeric()`. PETSc uses two different API's to take advantage of these optimizations.

The first approach explicitly divides the computation in the API. This approach is used, for example, with `MatCholeskyFactorSymbolic()`, `MatCholeskyFactorNumeric()`. The caller can take advantage of their knowledge of changes in the nonzero structure of the sparse matrices to call the appropriate routines as needed. In fact, they can use `MatGetNonzeroState()` to determine if a new symbolic computation is needed. The drawback of this approach is that the caller of these routines has to manage the creation of new matrices when the nonzero structure changes.

The second approach, as exemplified by `MatTranspose()`, does not expose the two stages explicit in the API, instead a flag, `MatReuse` is passed through the API to indicate if a symbolic data structure is already available or needs to be computed. Thus `MatTranspose(A,MAT_INITIAL_MATRIX,&B)` is called first, then `MatTranspose(A,MAT_REUSE_MATRIX,&B)` can be called repeatedly with new numerical values in the A matrix. In theory, if the nonzero structure of A changes, the symbolic computations for B could be redone automatically inside the same B matrix when there is a change in the nonzero state of the A matrix. In practice, in PETSc, the `MAT_REUSE_MATRIX` for most PETSc routines only works if the nonzero structure does not change and the code may crash otherwise. The advantage of this approach (when the nonzero structure changes are handled correctly) is that the calling code does not need to keep track of the nonzero state of the matrices; everything “just works”. However, the caller must still know when it is the first call to the routine so the flag `MAT_INITIAL_MATRIX` is being used. If the underlying implementation language supported detecting a yet to be initialized variable at run time, the `MatReuse` flag would not be need.

PETSc uses two approaches because the same programming problem was solved with two different ways during PETSc’s early development. A better model would combine both approaches; an explicit separation of the stages and a unified operation that internally utilized the two stages appropriately and also handled changes to the nonzero structure. Code could be simplified in many places with this approach, in most places the use of the unified API would replace the use of the separate stages.

See Extracting Submatrices and Matrix-Matrix Products.

2.2.9 Graph Operations

PETSc has four families of graph operations that treat sparse Mat as representing graphs.

<table>
<thead>
<tr>
<th>Operation</th>
<th>Type</th>
<th>Available methods</th>
<th>User guide</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ordering to reduce fill</td>
<td>N/A</td>
<td>MatOrderingType</td>
<td>Matrix Factorization</td>
</tr>
<tr>
<td>Partitioning for parallelism</td>
<td>MatPartitioning</td>
<td>MatPartitioningType</td>
<td>Partitioning</td>
</tr>
<tr>
<td>Coloring for parallelism or Jacobians</td>
<td>MatColoring</td>
<td>MatColoringType</td>
<td>Finite Difference Jacobian Approximations</td>
</tr>
<tr>
<td>Coarsening for multigrid</td>
<td>MatCoarsen</td>
<td>MatCoarsenType</td>
<td>Algebraic Multigrid (AMG) Preconditioners</td>
</tr>
</tbody>
</table>
2.2.10 Partitioning

For almost all unstructured grid computation, the distribution of portions of the grid across the process’s work load and memory can have a very large impact on performance. In most PDE calculations the grid partitioning and distribution across the processes can (and should) be done in a “pre-processing” step before the numerical computations. However, this does not mean it need be done in a separate, sequential program; rather, it should be done before one sets up the parallel grid data structures in the actual program. PETSc provides an interface to the ParMETIS (developed by George Karypis; see the PETSc installation instructions for directions on installing PETSc to use ParMETIS) to allow the partitioning to be done in parallel. PETSc does not currently provide directly support for dynamic repartitioning, load balancing by migrating matrix entries between processes, etc. For problems that require mesh refinement, PETSc uses the “rebuild the data structure” approach, as opposed to the “maintain dynamic data structures that support the insertion/deletion of additional vector and matrix rows and columns entries” approach.

Partitioning in PETSc is organized around the MatPartitioning object. One first creates a parallel matrix that contains the connectivity information about the grid (or other graph-type object) that is to be partitioned. This is done with the command

```c
MatCreateMPIAdj(MPI_Comm comm, int mlocal, PetscInt n, const PetscInt ia[], const PetscInt ja[], PetscInt *weights, Mat *Adj);
```

The argument mlocal indicates the number of rows of the graph being provided by the given process, n is the total number of columns; equal to the sum of all the mlocal. The arguments ia and ja are the row pointers and column pointers for the given rows; these are the usual format for parallel compressed sparse row storage, using indices starting at 0, not 1.

![Diagram of a simple unstructured grid](image)

Fig. 2.4: Numbering on Simple Unstructured Grid

This, of course, assumes that one has already distributed the grid (graph) information among the processes. The details of this initial distribution is not important; it could be simply determined by assigning to the first process the first \( n_0 \) nodes from a file, the second process the next \( n_1 \) nodes, etc.

For example, we demonstrate the form of the ia and ja for a triangular grid where we

1. partition by element (triangle)
   - Process 0: mlocal = 2, n = 4, ja = {2, 3, 3}, ia = {0, 2, 3}
• Process 1: \( m_{local} = 2, n = 4, ja = \{0, 0, 1\}, ia = \{0, 1, 3\} \)

Note that elements are not connected to themselves and we only indicate edge connections (in some contexts single vertex connections between elements may also be included). We use a space above to denote the transition between rows in the matrix; and

\((2)\) partition by vertex.

• Process 0: \( m_{local} = 3, n = 6, ja = \{3, 4, 5, 3, 4, 5\}, ia = \{0, 2, 4, 7\} \)

• Process 1: \( m_{local} = 3, n = 6, ja = \{0, 2, 4, 0, 1, 2, 3, 5, 1, 2, 4\}, ia = \{0, 3, 8, 11\} \).

Once the connectivity matrix has been created the following code will generate the renumbering required for the new partition

\[
\begin{align*}
\text{MatPartitioningCreate(MPI Comm comm,MatPartitioning *part);} \\
\text{MatPartitioningSetAdjacency(MatPartitioning part,Mat Adj);} \\
\text{MatPartitioningSetFromOptions(MatPartitioning part);} \\
\text{MatPartitioningApply(MatPartitioning part,IS *is);} \\
\text{MatPartitioningDestroy(MatPartitioning *part);} \\
\text{ISPartitioningToNumbering(IS is,IS *isg);} \\
\end{align*}
\]

The resulting \( isg \) contains for each local node the new global number of that node. The resulting \( is \) contains the new process number that each local node has been assigned to.

Now that a new numbering of the nodes has been determined, one must renumber all the nodes and migrate the grid information to the correct process. The command

\[
\text{AOCreateBasicIS(isg,NULL,&ao);}
\]

generates, see Application Orderings, an \( AO \) object that can be used in conjunction with the \( is \) and \( isg \) to move the relevant grid information to the correct process and renumber the nodes etc. In this context, the new ordering is the “application” ordering so \( AO\text{PetscToApplication()} \) converts old global indices to new global indices and \( AO\text{ApplicationToPetsc()} \) converts new global indices back to old global indices.

PETSc does not currently provide tools that completely manage the migration and node renumbering, since it will be dependent on the particular data structure you use to store the grid information and the type of grid information that you need for your application. We do plan to include more support for this in the future, but designing the appropriate general user interface and providing a scalable implementation that can be used for a wide variety of different grids requires a great deal of time.

See Finite Difference Jacobian Approximations and Matrix Factorization for discussions on performing graph coloring and computing graph reorderings to reduce fill in sparse matrix factorizations.

### 2.3 KSP: Linear System Solvers

The KSP object is the heart of PETSc, because it provides uniform and efficient access to all of the package’s linear system solvers, including parallel and sequential, direct and iterative. KSP is intended for solving systems of the form

\[
Ax = b, \tag{2.1}
\]

where \( A \) denotes the matrix representation of a linear operator, \( b \) is the right-hand-side vector, and \( x \) is the solution vector. KSP uses the same calling sequence for both direct and iterative solution of a linear system. In addition, particular solution techniques and their associated options can be selected at runtime.
The combination of a Krylov subspace method and a preconditioner is at the center of most modern numerical codes for the iterative solution of linear systems. Many textbooks (e.g. [FGN92] [vdV03], or [Saa03]) provide an overview of the theory of such methods. The KSP package, discussed in Krylov Methods, provides many popular Krylov subspace iterative methods; the PC module, described in Preconditioners, includes a variety of preconditioners.

2.3.1 Using KSP

To solve a linear system with KSP, one must first create a solver context with the command

```
KSPCreate(MPI_Comm comm,KSP *ksp);
```

Here comm is the MPI communicator and ksp is the newly formed solver context. Before actually solving a linear system with KSP, the user must call the following routine to set the matrices associated with the linear system:

```
KSPSetOperators(KSP ksp,Mat Amat,Mat Pmat);
```

The argument Amat, representing the matrix that defines the linear system, is a symbolic placeholder for any kind of matrix or operator. In particular, KSP does support matrix-free methods. The routine MatCreateShell() in Matrix-Free Matrices provides further information regarding matrix-free methods. Typically, the matrix from which the preconditioner is to be constructed, Pmat, is the same as the matrix that defines the linear system, Amat; however, occasionally these matrices differ (for instance, when a preconditioning matrix is obtained from a lower order method than that employed to form the linear system matrix).

Much of the power of KSP can be accessed through the single routine

```
KSPSetFromOptions(KSP ksp);
```

This routine accepts the option -help as well as any of the KSP and PC options discussed below. To solve a linear system, one sets the right hand size and solution vectors using the command

```
KSPSolve(KSP ksp,Vec b,Vec x);
```

where b and x respectively denote the right-hand-side and solution vectors. On return, the iteration number at which the iterative process stopped can be obtained using

```
KSPGetIterationNumber(KSP ksp, PetscInt *its);
```

Note that this does not state that the method converged at this iteration: it can also have reached the maximum number of iterations, or have diverged.

Convergence Tests gives more details regarding convergence testing. Note that multiple linear solves can be performed by the same KSP context. Once the KSP context is no longer needed, it should be destroyed with the command

```
KSPDestroy(KSP *ksp);
```

The above procedure is sufficient for general use of the KSP package. One additional step is required for users who wish to customize certain preconditioners (e.g., see Block Jacobi and Overlapping Additive Schwarz Preconditioners) or to log certain performance data using the PETSc profiling facilities (as discussed in Profiling). In this case, the user can optionally explicitly call

```
KSPSetUp(KSP ksp);
```
before calling `KSPSolve()` to perform any setup required for the linear solvers. The explicit call of this routine enables the separate monitoring of any computations performed during the set up phase, such as incomplete factorization for the ILU preconditioner.

The default solver within KSP is restarted GMRES, preconditioned for the uniprocess case with ILU(0), and for the multiprocess case with the block Jacobi method (with one block per process, each of which is solved with ILU(0)). A variety of other solvers and options are also available. To allow application programmers to set any of the preconditioner or Krylov subspace options directly within the code, we provide routines that extract the PC and KSP contexts,

```c
KSPGetPC(KSP ksp, PC *pc);
```

The application programmer can then directly call any of the PC or KSP routines to modify the corresponding default options.

To solve a linear system with a direct solver (currently supported by PETSc for sequential matrices, and by several external solvers through PETSc interfaces, see Using External Linear Solvers) one may use the options `-ksp_type preonly` (or the equivalent `-ksp_type none`) `-pc_type lu` (see below).

By default, if a direct solver is used, the factorization is not done in-place. This approach prevents the user from the unexpected surprise of having a corrupted matrix after a linear solve. The routine `PCFactorSetUseInPlace()`, discussed below, causes factorization to be done in-place.

### 2.3.2 Solving Successive Linear Systems

When solving multiple linear systems of the same size with the same method, several options are available. To solve successive linear systems having the same preconditioner matrix (i.e., the same data structure with exactly the same matrix elements) but different right-hand-side vectors, the user should simply call `KSPSolve()` multiple times. The preconditioner setup operations (e.g., factorization for ILU) will be done during the first call to `KSPSolve()` only; such operations will not be repeated for successive solves.

To solve successive linear systems that have different preconditioner matrices (i.e., the matrix elements and/or the matrix data structure change), the user must call `KSPSetOperators()` and `KSPSolve()` for each solve.

### 2.3.3 Krylov Methods

The Krylov subspace methods accept a number of options, many of which are discussed below. First, to set the Krylov subspace method that is to be used, one calls the command

```c
KSPSetType(KSP ksp, KSPType method);
```

The type can be one of `KSPRICHARDSON`, `KSPCHEBYSHEV`, `KSPCG`, `KSPGMRES`, `KSPTCQMR`, `KSPBCGS`, `KSPCGS`, `KSPTFQMR`, `KSPCR`, `KSPLSQR`, `KSPBICG`, `KSPPREONLY` (or the equivalent `KSPNONE`), or others; see `KSP Objects` or the `KSPType` man page for more. The KSP method can also be set with the options database command `-ksp_type`, followed by one of the options `richardson`, `chebyshev`, `cg`, `gmres`, `tcqmr`, `bcgs`, `cgs`, `tfqmr`, `cr`, `lsqr`, `bicg`, `preonly` (or the equivalent `none`), or others (see `KSP Objects` or the `KSPType` man page). There are method-specific options. For instance, for the Richardson, Chebyshev, and GMRES methods:

```c
KSPRICHARDSONSetScale(KSP ksp, PetscReal scale);
KSPCHEBYSHEVSetEigenvalues(KSP ksp, PetscReal emax, PetscReal emin);
KSPGMRESRestart(KSP ksp, PetscInt max_steps);
```
The default parameter values are `scale=1.0`, `emax=0.01`, `emin=100.0`, and `max_steps=30`. The GMRES restart and Richardson damping factor can also be set with the options `-ksp_gmres_restart <n>` and `-ksp_richardson_scale <factor>.

The default technique for orthogonalization of the Krylov vectors in GMRES is the unmodified (classical) Gram-Schmidt method, which can be set with

```c
KSPGMRESSetOrthogonalization(KSP ksp,KSPGMRESClassicalGramSchmidtOrthogonalization);
```

or the options database command `-ksp_gmres_classicalgramschmidt`. By default this will not use iterative refinement to improve the stability of the orthogonalization. This can be changed with the option

```c
KSPGMRESSetCGSRefinementType(KSP ksp,KSPGMRESCGSRefinementType type)
```

or via the options database with

```bash
-ksp_gmres_cgs_refinement_type <refine_never,refine_ifneeded,refine_always>
```

The values for `KSPGMRESCGSRefinementType()` are `KSP_GMRES_CGS_REFINE_NEVER`, `KSP_GMRES_CGS_REFINE_IFNEEDED` and `KSP_GMRES_CGS_REFINE_ALWAYS`.

One can also use modified Gram-Schmidt, by using the orthogonalization routine `KSPGMRESModifiedGramSchmidtOrthogonalization()` or by using the command line option `-ksp_gmres_modifiedgramschmidt`.

For the conjugate gradient method with complex numbers, there are two slightly different algorithms depending on whether the matrix is Hermitian symmetric or truly symmetric (the default is to assume that it is Hermitian symmetric). To indicate that it is symmetric, one uses the command

```c
KSPCGSetType(ksp,KSP_CG_SYMMETRIC);
```

Note that this option is not valid for all matrices.

Some `KSP` types do not support preconditioning. For instance, the CGLS algorithm does not involve a preconditioner; any preconditioner set to work with the `KSP` object is ignored if `KSPCGLS` was selected.

By default, `KSP` assumes an initial guess of zero by zeroing the initial value for the solution vector that is given; this zeroing is done at the call to `KSPSolve()`. To use a nonzero initial guess, the user must call

```c
KSPSetInitialGuessNonzero(KSP ksp,PetscBool flg);
```

### Preconditioning within KSP

Since the rate of convergence of Krylov projection methods for a particular linear system is strongly dependent on its spectrum, preconditioning is typically used to alter the spectrum and hence accelerate the convergence rate of iterative techniques. Preconditioning can be applied to the system (2.1) by

\[
(M_L^{-1} A M_R^{-1}) (M_R x) = M_L^{-1} b,
\]

where $M_L$ and $M_R$ indicate preconditioning matrices (or, matrices from which the preconditioner is to be constructed). If $M_L = I$ in (2.2), right preconditioning results, and the residual of (2.1),

\[
r \equiv b - A x = b - A M_R^{-1} M_R x,
\]

is preserved. In contrast, the residual is altered for left ($M_R = I$) and symmetric preconditioning, as given by

\[
r_L \equiv M_L^{-1} b - M_L^{-1} A x = M_L^{-1} r.
\]
By default, most KSP implementations use left preconditioning. Some more naturally use other options, though. For instance, KSPQCG defaults to use symmetric preconditioning and KSPFGMRES uses right preconditioning by default. Right preconditioning can be activated for some methods by using the options database command -ksp_pc_side right or calling the routine

```c
KSPSetPCSide(ksp,PC_RIGHT);
```

Attempting to use right preconditioning for a method that does not currently support it results in an error message of the form

```
KSPSetUp_Richardson:No right preconditioning for KSPRICHARDSON
```

We summarize the defaults for the residuals used in KSP convergence monitoring within KSP Objects. Details regarding specific convergence tests and monitoring routines are presented in the following sections. The preconditioned residual is used by default for convergence testing of all left-preconditioned KSP methods. For the conjugate gradient, Richardson, and Chebyshev methods the true residual can be used by the options database command -ksp_norm_type unpreconditioned or by calling the routine

```c
KSPSetNormType(ksp,KSP_NORM_UNPRECONDITIONED);
```

**Table 2.5: KSP Objects**

<table>
<thead>
<tr>
<th>Method</th>
<th>KSPT ype</th>
<th>Options Database Name</th>
</tr>
</thead>
<tbody>
<tr>
<td>Richardson</td>
<td>KSPRICHARDSON</td>
<td>richardson</td>
</tr>
<tr>
<td>Chebyshev</td>
<td>KSPCHEBYSHEV</td>
<td>chebyshev</td>
</tr>
<tr>
<td>Conjugate Gradient [HS52]</td>
<td>KSPCG</td>
<td>cg</td>
</tr>
<tr>
<td>Pipelined Conjugate Gradients [GV14]</td>
<td>KSPPIPECG</td>
<td>pipexcg</td>
</tr>
<tr>
<td>Pipelined Conjugate Gradients (Gropp)</td>
<td>KSPGROPPCG</td>
<td>groppcg</td>
</tr>
<tr>
<td>Pipelined Conjugate Gradients with Residual Replacement</td>
<td>KSPPIPECGRR</td>
<td>pipecgrr</td>
</tr>
<tr>
<td>Conjugate Gradients for the Normal Equations</td>
<td>KSPCGNE</td>
<td>cgne</td>
</tr>
<tr>
<td>Flexible Conjugate Gradients [Not00]</td>
<td>KSPFCG</td>
<td>fcg</td>
</tr>
<tr>
<td>Pipelined, Flexible Conjugate Gradients [SSM16]</td>
<td>KSPPIPEFCG</td>
<td>pipef cg</td>
</tr>
<tr>
<td>Conjugate Gradients for Least Squares</td>
<td>KSPCGLS</td>
<td>cgls</td>
</tr>
<tr>
<td>Conjugate Gradients with Constraint (1)</td>
<td>KSPNASH</td>
<td>nash</td>
</tr>
<tr>
<td>Conjugate Gradients with Constraint (2)</td>
<td>KSPSTCG</td>
<td>stcg</td>
</tr>
<tr>
<td>Conjugate Gradients with Constraint (3)</td>
<td>KSPGLTR</td>
<td>gltr</td>
</tr>
<tr>
<td>Conjugate Gradients with Constraint (4)</td>
<td>KSPQCG</td>
<td>qcg</td>
</tr>
<tr>
<td>BiConjugate Gradient</td>
<td>KSPBICG</td>
<td>bicg</td>
</tr>
<tr>
<td>BiCGSTAB [vandVorst92]</td>
<td>KSPBCGS</td>
<td>bcs</td>
</tr>
<tr>
<td>Improved BiCGSTAB</td>
<td>KSPIBCGS</td>
<td>ibcgs</td>
</tr>
<tr>
<td>QMRCGSTAB [CGS+94]</td>
<td>KSPQMRCS</td>
<td>qmrccs</td>
</tr>
<tr>
<td>Flexible BiCGSTAB</td>
<td>KSPFBCGS</td>
<td>fbcs</td>
</tr>
<tr>
<td>Flexible BiCGSTAB (variant)</td>
<td>KSPFBCGSR</td>
<td>fbcsr</td>
</tr>
<tr>
<td>Enhanced BiCGSTAB(L)</td>
<td>KSPBGCGL</td>
<td>bgcs</td>
</tr>
<tr>
<td>Minimal Residual Method [PS75]</td>
<td>KSPMINRES</td>
<td>minres</td>
</tr>
<tr>
<td>Generalized Minimal Residual [SS86]</td>
<td>KSPGMRRES</td>
<td>gmres</td>
</tr>
<tr>
<td>Flexible Generalized Minimal Residual [Saa93]</td>
<td>KSPFGMRRES</td>
<td>fgmres</td>
</tr>
<tr>
<td>Deflated Generalized Minimal Residual</td>
<td>KSPDGMRES</td>
<td>dgmres</td>
</tr>
<tr>
<td>Pipelined Generalized Minimal Residual [GAMV13]</td>
<td>KSPPGMRRES</td>
<td>pgmres</td>
</tr>
<tr>
<td>Pipelined, Flexible Generalized Minimal Residual [SSM16]</td>
<td>KSPPIPEFGMRRES</td>
<td>pipexfgmres</td>
</tr>
<tr>
<td>Generalized Minimal Residual with Accelerated Restart</td>
<td>KSPLMRRES</td>
<td>lgmres</td>
</tr>
<tr>
<td>Conjugate Residual [EES83]</td>
<td>KSPCR</td>
<td>cr</td>
</tr>
<tr>
<td>Generalized Conjugate Residual</td>
<td>KSPGCR</td>
<td>gcr</td>
</tr>
</tbody>
</table>

continues on next page

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Note: the bi-conjugate gradient method requires application of both the matrix and its transpose plus the preconditioner and its transpose. Currently not all matrices and preconditioners provide this support and thus the KSPBICG cannot always be used.

Note: PETSc implements the FETI-DP (Finite Element Tearing and Interconnecting Dual-Primal) method as an implementation of KSP since it recasts the original problem into a constrained minimization one with Lagrange multipliers. The only matrix type supported is MATIS. Support for saddle point problems is provided. See the man page for KSPFETIDP for further details.

**Convergence Tests**

The default convergence test, KSPConvergedDefault(), is based on the $l_2$-norm of the residual. Convergence (or divergence) is decided by three quantities: the decrease of the residual norm relative to the norm of the right hand side, $rtol$, the absolute size of the residual norm, $atol$, and the relative increase in the residual, $dtol$. Convergence is detected at iteration $k$ if

$$\|r_k\|_2 < \max(rtol \ast \|b\|_2, atol),$$

where $r_k = b - Ax_k$. Divergence is detected if

$$\|r_k\|_2 > dtol \ast \|b\|_2.$$

These parameters, as well as the maximum number of allowable iterations, can be set with the routine

```c
KSPSetTolerances(KSP ksp, PetscReal rtol, PetscReal atol, PetscReal dtol, PetscInt maxits);
```

The user can retain the default value of any of these parameters by specifying PETSC_DEFAULT as the corresponding tolerance; the defaults are $rtol=1e-5$, $atol=1e-50$, $dtol=1e5$, and $maxits=1e4$. These parameters can also be set from the options database with the commands -ksp_rtol <rtol>, -ksp_atol <atol>, -ksp_divtol <dtol>, and -ksp_max_it <its>.

In addition to providing an interface to a simple convergence test, KSP allows the application programmer the flexibility to provide customized convergence-testing routines. The user can specify a customized routine with the command

```c
KSPSetConvergenceTest(KSP ksp, PetscErrorCode (*test)(KSP ksp, PetscInt it, PetscReal rnorm, KSPConvergedReason *reason, void *ctx), void *ctx, PetscErrorCode (*destroy)(void *ctx));
```
The final routine argument, `ctx`, is an optional context for private data for the user-defined convergence routine, `test`. Other `test` routine arguments are the iteration number, `it`, and the residual’s $l_2$ norm, `rnorm`. The routine for detecting convergence, `test`, should set `reason` to positive for convergence, 0 for no convergence, and negative for failure to converge. A full list of possible values is given in the `KSPConvergedReason` manual page. You can use `KSPGetConvergedReason()` after `KSPSolve()` to see why convergence/divergence was detected.

**Convergence Monitoring**

By default, the Krylov solvers run silently without displaying information about the iterations. The user can indicate that the norms of the residuals should be displayed by using `-ksp_monitor` within the options database. To display the residual norms in a graphical window (running under X Windows), one should use `-ksp_monitor draw::draw_lg`. Application programmers can also provide their own routines to perform the monitoring by using the command

```
KSPMonitorSet(KSP ksp,PetscErrorCode (*mon)(KSP ksp,PetscInt it,PetscReal rnorm,void *,--*ctx),void *ctx,PetscErrorCode (*mondestroy)(void **));
```

The final routine argument, `ctx`, is an optional context for private data for the user-defined monitoring routine, `mon`. Other `mon` routine arguments are the iteration number (`it`) and the residual’s $l_2$ norm (`rnorm`). A helpful routine within user-defined monitors is `PetscObjectGetComm((PetscObject)ksp,MPI_Comm *comm)`, which returns in `comm` the MPI communicator for the KSP context. See *Writing PETSc Programs* for more discussion of the use of MPI communicators within PETSc.

Several monitoring routines are supplied with PETSc, including

```
KSPMonitorResidual(KSP,PetscInt,PetscReal, void *);
KSPMonitorSingularValue(KSP,PetscInt,PetscReal, void *);
KSPMonitorTrueResidual(KSP,PetscInt,PetscReal, void *);
```

The default monitor simply prints an estimate of the $l_2$-norm of the residual at each iteration. The routine `KSPMonitorSingularValue()` is appropriate only for use with the conjugate gradient method or GMRES, since it prints estimates of the extreme singular values of the preconditioned operator at each iteration. Since `KSPMonitorTrueResidual()` prints the true residual at each iteration by actually computing the residual using the formula $r = b - Ax$, the routine is slow and should be used only for testing or convergence studies, not for timing. These monitors may be accessed with the command line options `-ksp_monitor`, `-ksp_monitor_singular_value`, and `-ksp_monitor_true_residual`.

To employ the default graphical monitor, one should use the command `-ksp_monitor draw::draw_lg`. One can cancel hardwired monitoring routines for KSP at runtime with `-ksp_monitor_cancel`.

Unless the Krylov method converges so that the residual norm is small, say $10^{-10}$, many of the final digits printed with the `-ksp_monitor` option are meaningless. Worse, they are different on different machines; due to different round-off rules used by, say, the IBM RS6000 and the Sun SPARC. This makes testing between different machines difficult. The option `-ksp_monitor_short` causes PETSc to print fewer of the digits of the residual norm as it gets smaller; thus on most of the machines it will always print the same numbers making cross system testing easier.
Understanding the Operator’s Spectrum

Since the convergence of Krylov subspace methods depends strongly on the spectrum (eigenvalues) of the preconditioned operator, PETSc has specific routines for eigenvalue approximation via the Arnoldi or Lanczos iteration. First, before the linear solve one must call

\[
\text{KSPSetComputeEigenvalues(ksp,PETSC_TRUE);} \\
\]

Then after the \textit{KSP} solve one calls

\[
\text{KSPComputeEigenvalues(KSP \; \text{ksp}, \text{PetscInt} \; n, \text{PetscReal} \; *\text{realpart}, \text{PetscReal} \; *\text{complexpart}, \ldots \text{PetscInt} \; *\text{neig});} \\
\]

Here, \( n \) is the size of the two arrays and the eigenvalues are inserted into those two arrays. \( \text{neig} \) is the number of eigenvalues computed; this number depends on the size of the Krylov space generated during the linear system solution, for GMRES it is never larger than the restart parameter. There is an additional routine

\[
\text{KSPComputeEigenvaluesExplicitly(KSP \; \text{ksp}, \text{PetscInt} \; n, \text{PetscReal} \; *\text{realpart}, \text{PetscReal} \; *\text{complexpart});} \\
\]

that is useful only for very small problems. It explicitly computes the full representation of the preconditioned operator and calls LAPACK to compute its eigenvalues. It should be only used for matrices of size up to a couple hundred. The \text{PetscDrawSP*()} routines are very useful for drawing scatter plots of the eigenvalues.

The eigenvalues may also be computed and displayed graphically with the options data base commands \text{-ksp_view_eigenvalues draw} and \text{-ksp_view_eigenvalues_explicit draw}. Or they can be dumped to the screen in ASCII text via \text{-ksp_view_eigenvalues} and \text{-ksp_view_eigenvalues_explicit}.

Flexible Krylov Methods

Standard Krylov methods require that the preconditioner be a linear operator, thus, for example, a standard \text{KSP} method cannot use a \text{KSP} in its preconditioner, as is common in the Block-Jacobi method \text{PCBJACOBI}, for example. Flexible Krylov methods are a subset of methods that allow (with modest additional requirements on memory) the preconditioner to be nonlinear. For example, they can be used with the \text{PCKSP} preconditioner. The flexible \text{KSP} methods have the label “Flexible” in \text{KSP Objects}.

One can use \text{KSPMonitorDynamicTolerance()} to control the tolerances used by inner \text{KSP} solvers in \text{PCKSP}, \text{PCBJACOBI}, and \text{PCDEFLATION}.

In addition to supporting \text{PCKSP}, the flexible methods support \text{KSP*SetModifyPC()}, for example, \text{KSPFGMRESSetModifyPC()}, these functions allow the user to provide a callback function that changes the preconditioner at each Krylov iteration. Its calling sequence is as follows.

\[
\text{PetscErrorCode \; f(KSP \; \text{ksp}, \text{PetscInt} \; \text{total_its}, \text{PetscInt} \; \text{its_since_restart}, \text{PetscReal} \; \text{res}\_\text{norm}, \text{void} \; *\text{ctx});} \\
\]
Pipelined Krylov Methods

Standard Krylov methods have one or more global reductions resulting from the computations of inner products or norms in each iteration. These reductions need to block until all MPI ranks have received the results. For a large number of MPI ranks (this number is machine dependent but can be above 10,000 ranks) this synchronization is very time consuming and can significantly slow the computation. Pipelined Krylov methods overlap the reduction operations with local computations (generally the application of the matrix-vector products and preconditioners) thus effectively “hiding” the time of the reductions. In addition, they may reduce the number of global synchronizations by rearranging the computations in a way that some of them can be collapsed, e.g., two or more calls to `MPI_Allreduce()` may be combined into one call. The pipeline KSP methods have the label “Pipeline” in *KSP Objects*.

Special configuration of MPI may be necessary for reductions to make asynchronous progress, which is important for performance of pipelined methods. See *doc_faq_pipelined* for details.

Other KSP Options

To obtain the solution vector and right hand side from a KSP context, one uses

```c
KSPGetSolution(KSP ksp, Vec *x);
KSPGetRhs(KSP ksp, Vec *rhs);
```

During the iterative process the solution may not yet have been calculated or it may be stored in a different location. To access the approximate solution during the iterative process, one uses the command

```c
KSPBuildSolution(KSP ksp, Vec w, Vec *v);
```

where the solution is returned in `v`. The user can optionally provide a vector in `w` as the location to store the vector; however, if `w` is `NULL`, space allocated by PETSc in the KSP context is used. One should not destroy this vector. For certain KSP methods (e.g., GMRES), the construction of the solution is expensive, while for many others it doesn’t even require a vector copy.

Access to the residual is done in a similar way with the command

```c
KSPBuildResidual(KSP ksp, Vec t, Vec w, Vec *v);
```

Again, for GMRES and certain other methods this is an expensive operation.

2.3.4 Preconditioners

As discussed in *Preconditioning within KSP*, Krylov subspace methods are typically used in conjunction with a preconditioner. To employ a particular preconditioning method, the user can either select it from the options database using input of the form `-pc_type <methodname>` or set the method with the command

```c
PCSetType(PC pc, PCTYPE method);
```

In *PETSc Preconditioners (partial list)* we summarize the basic preconditioning methods supported in PETSc. See the PCTYPE manual page for a complete list. The PCSHELL preconditioner uses a specific, application-provided preconditioner. The direct preconditioner, PCLU, is, in fact, a direct solver for the linear system that uses LU factorization. PCLU is included as a preconditioner so that PETSc has a consistent interface among direct and iterative linear solvers.
Table 2.6: PETSc Preconditioners (partial list)

<table>
<thead>
<tr>
<th>Method</th>
<th>PCTYPE</th>
<th>Options Database Name</th>
</tr>
</thead>
<tbody>
<tr>
<td>Jacobi</td>
<td>PCJACOBI</td>
<td>jacobi</td>
</tr>
<tr>
<td>Block Jacobi</td>
<td>PCBJACOBI</td>
<td>bjacobi</td>
</tr>
<tr>
<td>SOR (and SSOR)</td>
<td>PCSOR</td>
<td>sor</td>
</tr>
<tr>
<td>SOR with Eisenstat trick</td>
<td>PCEISENSTAT</td>
<td>eisenstat</td>
</tr>
<tr>
<td>Incomplete Cholesky</td>
<td>PCICC</td>
<td>icc</td>
</tr>
<tr>
<td>Incomplete LU</td>
<td>PCILU</td>
<td>ilu</td>
</tr>
<tr>
<td>Additive Schwarz</td>
<td>PCASM</td>
<td>asm</td>
</tr>
<tr>
<td>Generalized Additive Schwarz</td>
<td>PCGASM</td>
<td>gasm</td>
</tr>
<tr>
<td>Algebraic Multigrid</td>
<td>PCGAMG</td>
<td>gamg</td>
</tr>
<tr>
<td>Balancing Domain Decomposition by Constraints</td>
<td>PCBDDC</td>
<td>bddc</td>
</tr>
<tr>
<td>Linear solver</td>
<td>PCKSP</td>
<td>ksp</td>
</tr>
<tr>
<td>Combination of preconditioners</td>
<td>PCCOMPOSITE</td>
<td>composite</td>
</tr>
<tr>
<td>LU</td>
<td>PCLU</td>
<td>lu</td>
</tr>
<tr>
<td>Cholesky</td>
<td>PCCHOLESKY</td>
<td>cholesky</td>
</tr>
<tr>
<td>No preconditioning</td>
<td>PCNONE</td>
<td>none</td>
</tr>
<tr>
<td>Shell for user-defined PC</td>
<td>PCSHELL</td>
<td>shell</td>
</tr>
</tbody>
</table>

Each preconditioner may have associated with it a set of options, which can be set with routines and options database commands provided for this purpose. Such routine names and commands are all of the form `PC<TYPE><Option>` and `-pc_<type>_<option> [value]`. A complete list can be found by consulting the `PCTYPE` manual page; we discuss just a few in the sections below.

### ILU and ICC Preconditioners

Some of the options for ILU preconditioner are

```c
PCFactorSetLevels(PC pc,PetscInt levels);
PCFactorSetReuseOrdering(PC pc,PetscBool flag);
PCFactorSetDropTolerance(PC pc,PetscReal dt,PetscReal dtcol,PetscInt dtcount);
PCFactorSetReuseFill(PC pc,PetscBool flag);
PCFactorSetUseInPlace(PC pc,PetscBool flg);
PCFactorSetAllowDiagonalFill(PC pc,PetscBool flg);
```

When repeatedly solving linear systems with the same KSP context, one can reuse some information computed during the first linear solve. In particular, `PCFactorSetReuseOrdering()` causes the ordering (for example, set with `-pc_factor_mat_ordering_type order`) computed in the first factorization to be reused for later factorizations. `PCFactorSetUseInPlace()` is often used with `PCASM` or `PCBJACOBI` when zero fill is used, since it reuses the matrix space to store the incomplete factorization it saves memory and copying time. Note that in-place factorization is not appropriate with any ordering besides natural and cannot be used with the drop tolerance factorization. These options may be set in the database with

- `-pc_factor_levels <levels>`
- `-pc_factor_reuse_ordering`
- `-pc_factor_reuse_fill`
- `-pc_factor_in_place`
- `-pc_factor_nonzeros_along_diagonal`
- `-pc_factor_diagonal_fill`
See *Memory Allocation for Sparse Matrix Factorization* for information on preallocation of memory for anticipated fill during factorization. By alleviating the considerable overhead for dynamic memory allocation, such tuning can significantly enhance performance.

PETSc supports incomplete factorization preconditioners for several matrix types for sequential matrices (for example **MATSEQAIJ**, **MATSEQBAIJ**, and **MATSEQSBAIJ**).

### SOR and SSOR Preconditioners

PETSc provides only a sequential SOR preconditioner; it can only be used with sequential matrices or as the subblock preconditioner when using block Jacobi or ASM preconditioning (see below).

The options for SOR preconditioning with **PCSOR** are

```c
PCSORSetOmega(PC pc, PetscReal omega);
PCSORSetIterations(PC pc, PetscInt its, PetscInt lits);
PCSORSetSymmetric(PC pc, MatSORType type);
```

The first of these commands sets the relaxation factor for successive over (under) relaxation. The second command sets the number of inner iterations **its** and local iterations **lits** (the number of smoothing sweeps on a process before doing a ghost point update from the other processes) to use between steps of the Krylov space method. The total number of SOR sweeps is given by **its***lits**. The third command sets the kind of SOR sweep, where the argument **type** can be one of **SOR_FORWARD_SWEEP**, **SOR_BACKWARD_SWEEP** or **SOR_SYMMETRIC_SWEEP**, the default being **SOR_FORWARD_SWEEP**. Setting the type to be **SOR_SYMMETRIC_SWEEP** produces the SSOR method. In addition, each process can locally and independently perform the specified variant of SOR with the types **SOR_LOCAL_FORWARD_SWEEP**, **SOR_LOCAL_BACKWARD_SWEEP**, and **SOR_LOCAL_SYMMETRIC_SWEEP**. These variants can also be set with the options `-pc_sor_omega <omega>`, `-pc_sor_its <its>`, `-pc_sor_lits <lits>`, `-pc_sor_backward`, `-pc_sor_symmetric`, `-pc_sor_local_forward`, `-pc_sor_local_backward`, and `-pc_sor_local_symmetric`.

The Eisenstat trick [Eis81] for SSOR preconditioning can be employed with the method **PCEISENSTAT** (**-pc_type eisenstat**). By using both left and right preconditioning of the linear system, this variant of SSOR requires about half of the floating-point operations for conventional SSOR. The option `-pc_eisenstat_no_diagonal_scaling` (or the routine **PCEisenstatSetNoDiagonalScaling()**) turns off diagonal scaling in conjunction with Eisenstat SSOR method, while the option `-pc_eisenstat_omega <omega>` (or the routine **PCEisenstatSetOmega(PC pc, PetscReal omega)**) sets the SSOR relaxation coefficient, **omega**, as discussed above.

### LU Factorization

The LU preconditioner provides several options. The first, given by the command

```c
PCFactorSetUseInPlace(PC pc, PetscBool flg);
```

causes the factorization to be performed in-place and hence destroys the original matrix. The options database variant of this command is **-pc_factor_in_place**. Another direct preconditioner option is selecting the ordering of equations with the command **-pc_factor_mat_ordering_type <ordering>**. The possible orderings are

- **MATORDERINGNATURAL** - Natural
- **MATORDERINGND** - Nested Dissection
- **MATORDERING1WD** - One-way Dissection
- **MATORDERINGRCM** - Reverse Cuthill-McKee

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- **MATORDERINGQMD** - Quotient Minimum Degree

These orderings can also be set through the options database by specifying one of the following:

- pc_factor_mat_ordering_type natural, or nd, or lwd, or rcm, or qmd. In addition, see MatGe-
tOrdering(), discussed in Matrix Factorization.

The sparse LU factorization provided in PETSc does not perform pivoting for numerical stability (since they are designed to preserve nonzero structure), and thus occasionally an LU factorization will fail with a zero pivot when, in fact, the matrix is non-singular. The option -pc_factor_nonzeros_along_diagonal <tol> will often help eliminate the zero pivot, by preprocessing the column ordering to remove small values from the diagonal. Here, tol is an optional tolerance to decide if a value is nonzero; by default it is 1.e-10.

In addition, Memory Allocation for Sparse Matrix Factorization provides information on preallocation of memory for anticipated fill during factorization. Such tuning can significantly enhance performance, since it eliminates the considerable overhead for dynamic memory allocation.

### Block Jacobi and Overlapping Additive Schwarz Preconditioners

The block Jacobi and overlapping additive Schwarz methods in PETSc are supported in parallel; however, only the uniprocess version of the block Gauss-Seidel method is currently in place. By default, the PETSc implementations of these methods employ ILU(0) factorization on each individual block (that is, the default solver on each subblock is PCType=PCILU, KSPTYPE=KSPPREONLY (or equivalently KSPTYPE=KSPNONE); the user can set alternative linear solvers via the options -sub_ksp_type and -sub_pc_type. In fact, all of the KSP and PC options can be applied to the subproblems by inserting the prefix -sub_ at the beginning of the option name. These options database commands set the particular options for all of the blocks within the global problem. In addition, the routines

```c
PCBJacobiGetSubKSP(PC pc,PetscInt *n_local,PetscInt *first_local,KSP **subksp);
PCASMGetSubKSP(PC pc,PetscInt *n_local,PetscInt *first_local,KSP **subksp);
```

extract the KSP context for each local block. The argument n_local is the number of blocks on the calling process, and first_local indicates the global number of the first block on the process. The blocks are numbered successively by processes from zero through bg - 1, where bg is the number of global blocks. The array of KSP contexts for the local blocks is given by subksp. This mechanism enables the user to set different solvers for the various blocks. To set the appropriate data structures, the user must explicitly call KSPSetUp() before calling PCBJacobiGetSubKSP() or PCASMGetSubKSP(). For further details, see KSP Tutorial ex7 or KSP Tutorial ex8.

The block Jacobi, block Gauss-Seidel, and additive Schwarz preconditioners allow the user to set the number of blocks into which the problem is divided. The options database commands to set this value are -pc_bjacobi_blocks n and -pc_bgs_blocks n, and, within a program, the corresponding routines are

```c
PCBJacobiSetTotalBlocks(PC pc,PetscInt blocks,PetscInt *size);
PCASMSetTotalSubdomains(PC pc,PetscInt n,IS *is,IS *islocal);
PCASMSetType(PC pc,PCASMType type);
```

The optional argument size is an array indicating the size of each block. Currently, for certain parallel matrix formats, only a single block per process is supported. However, the MATMPIAIJ and MATMPIBAIJ formats support the use of general blocks as long as no blocks are shared among processes. The is argument contains the index sets that define the subdomains.

The object PCASMType is one of PC_ASM_BASIC, PC_ASM_INTERPOLATE, PC_ASM_RESTRICT, or PC_ASM_NONE and may also be set with the options database -pc_asm_type [basic, interpolate, restrict, none]. The type PC_ASM_BASIC (or -pc_asm_type basic) corresponds to the standard additive Schwarz method that uses the full restriction and interpolation operators. The type
PC_ASM_RESTRICT (or -pc_asm_type restrict) uses a full restriction operator, but during the interpolation process ignores the off-process values. Similarly, PC_ASM_INTERPOLATE (or -pc_asm_type interpolate) uses a limited restriction process in conjunction with a full interpolation, while PC_ASM_NONE (or -pc_asm_type none) ignores off-process values for both restriction and interpolation. The ASM types with limited restriction or interpolation were suggested by Xiao-Chuan Cai and Marcus Sarkis [CS99].

PC_ASM_RESTRICT is the PETSc default, as it saves substantial communication and for many problems has the added benefit of requiring fewer iterations for convergence than the standard additive Schwarz method.

The user can also set the number of blocks and sizes on a per-process basis with the commands

```c
PCBJacobiSetLocalBlocks(PC pc, PetscInt blocks, PetscInt *size);
PCASMSetLocalSubdomains(PC pc, PetscInt N, IS *is, IS *islocal);
```

For the ASM preconditioner one can use the following command to set the overlap to compute in constructing the subdomains.

```c
PCASMSetOverlap(PC pc, PetscInt overlap);
```

The overlap defaults to 1, so if one desires that no additional overlap be computed beyond what may have been set with a call to PCASMSetTotalSubdomains() or PCASMSetLocalSubdomains(), then overlap must be set to be 0. In particular, if one does not explicitly set the subdomains in an application code, then all overlap would be computed internally by PETSc, and using an overlap of 0 would result in an ASM variant that is equivalent to the block Jacobi preconditioner. Note that one can define initial index sets is with any overlap via PCASMSetTotalSubdomains() or PCASMSetLocalSubdomains(); the routine PCASMSetOverlap() merely allows PETSc to extend that overlap further if desired.

PCGASM is an experimental generalization of PCASM that allows the user to specify subdomains that span multiple MPI ranks. This can be useful for problems where small subdomains result in poor convergence. To be effective, the multirank subproblems must be solved using a sufficient strong subsolver, such as LU, for which SuperLU_DIST or a similar parallel direct solver could be used; other choices may include a multigrid solver on the subdomains.

The interface for PCGASM is similar to that of PCASM. In particular, PCGASMTypedef is one of PC_GASM_BASIC, PC_GASM_INTERPOLATE, PC_GASM_RESTRICT, PC_GASM_NONE. These options have the same meaning as with PCASM and may also be set with the options database -pc_gasm_type [basic, interpolate, restrict, none].

Unlike PCASM, however, PCGASM allows the user to define subdomains that span multiple MPI ranks. The simplest way to do this is using a call to PCGASMSetTotalSubdomains(PC pc, PetscInt N) with the total number of subdomains N that is smaller than the MPI communicator size. In this case PCGASM will coalesce size/N consecutive single-rank subdomains into a single multi-rank subdomain. The single-rank subdomains contain the degrees of freedom corresponding to the locally-owned rows of the PCGASM preconditioning matrix – these are the subdomains PCASM and PCGASM use by default.

Each of the multirank subdomain subproblems is defined on the subcommunicator that contains the coalesced PCGASM ranks. In general this might not result in a very good subproblem if the single-rank problems corresponding to the coalesced ranks are not very strongly connected. In the future this will be addressed with a hierarchical partitioner that generates well-connected coarse subdomains first before subpartitioning them into the single-rank subdomains.

In the meantime the user can provide his or her own multi-rank subdomains by calling PCGASMSetsSubdomains(PC, IS[], IS[]) where each of the IS objects on the list defines the inner (without the overlap) or the outer (including the overlap) subdomain on the subcommunicator of the IS object. A helper subroutine PCGASMCreateSubdomains2D() is similar to PCASM’s but is capable of constructing multi-rank subdomains that can be then used with PCGASMSetsSubdomains(). An alternative way of creating multi-rank subdomains is by using the underlying DM object, if it is capable of generating such decompositions via DMCreateDomainDecomposition(). Ordinarily the decomposition specified
by the user via `PCGASMSetSubdomains()` takes precedence, unless `PCGASMSetUseDMSubdomains()` instructs `PCGASM` to prefer DM-created decompositions.

Currently there is no support for increasing the overlap of multi-rank subdomains via `PCGASMSetOverlap()` – this functionality works only for subdomains that fit within a single MPI rank, exactly as in `PCASM`.

Examples of the described `PCGASM` usage can be found in KSP Tutorial ex62. In particular, `runex62_superlu_dist` illustrates the use of `SuperLU_DIST` as the subdomain solver on coalesced multi-rank subdomains. The `runex62_2D_*` examples illustrate the use of `PCGASMCreateSubdomains2D()`.

### Algebraic Multigrid (AMG) Preconditioners

PETSc has a native algebraic multigrid preconditioner `PCGAMG` – `gamg` – and interfaces to three external AMG packages: `hypre`, `ML` and `AMGx` (CUDA platforms only), that can be downloaded in the configuration phase (eg, `-download-hypre`) and used by specifying that command line parameter (eg, `-pc_type hypre`). `Hypre` is relatively monolithic in that a PETSc matrix is converted into a hypre matrix and then `hypre` is called to do the entire solve. `ML` is more modular in that PETSc only has `ML` generate the coarse grid spaces (columns of the prolongation operator), which is core of an AMG method, and then constructs a `PCMG` with Galerkin coarse grid operator construction. `PCGAMG` is designed from the beginning to be modular, to allow for new components to be added easily and also populates a multigrid preconditioner `PCMG` so generic multigrid parameters are used (see Multigrid Preconditioners). PETSc provides a fully supported (smoothed) aggregation AMG, but supports the addition of new methods (`-pc_type gamg -pc_gamg_type agg` or `PCSetType(pc,PCGAMG)` and `PCGAMGSetType(pc,PCGAMMAGG)`). Examples of extension are a reference implementations of a classical AMG method (`-pc_gamg_type classical`), a (2D) hybrid geometric AMG method (`-pc_gamg_type geo`) that are not supported. A 2.5D AMG method `DofColumns [ISG15]` supports 2D coarsenings extruded in the third dimension. `PCGAMG` does require the use of `MATAIJ` matrices. For instance, `MATBAIJ` matrices are not supported. One can use `MATAIJ` instead of `MATBAIJ` without changing any code other than the constructor (or the `mat_type` from the command line). For instance, `MatSetValuesBlocked` works with `MATAIJ` matrices.

`PCGAMG` provides unsmoothed aggregation (`-pc_gamg_agg_nsmooths 0`) and smoothed aggregation (`-pc_gamg_agg_nsmooths 1` or `PCGAMGSetNsmooths(pc,1)`). Smoothed aggregation (SA) is recommended for symmetric positive definite systems. Unsmoothed aggregation can be useful for asymmetric problems and problems where highest eigen estimates are problematic. If poor convergence rates are observed using the smoothed version one can test unsmoothed aggregation.

#### Eigenvalue estimates:

The parameters for the KSP eigen estimator, used for SA, can be set with `-pc_gamg_esteig_ksp_max_it` and `-pc_gamg_esteig_ksp_type`. For example CG generally converges to the highest eigenvalue fast than GMRES (the default for KSP) if your problem is symmetric positive definite. One can specify CG with `-pc_gamg_esteig_ksp_type cg`. The default for `-pc_gamg_esteig_ksp_max_it` is 10, which we have found is pretty safe with a (default) safety factor of 1.1. One can specify the range of real eigenvalues, in the same way that one can for Chebyshev KSP solvers (smoothers), with `-pc_gamg_eigenvalues <emin,emax>`. GAMG sets the MG smoother type to chebyshev by default. By default, GAMG uses its own estimate, if it has one, for Chebyshev smoothers if the smoother uses Jacobi preconditioning. This can be overridden with `-pc_gamg_use_sa_esteig true,false`.

AMG methods requires knowledge of the number of degrees of freedom per vertex, the default is one (a scalar problem). Vector problems like elasticity should set the block size of the matrix appropriately with `-mat_block_size bs` or `MatSetBlockSize(mat,bs)`. Equations must be ordered in “vertex-major” ordering (e.g., $x_1, y_1, z_1, x_2, y_2, ...$).

#### Near null space:

Smoothed aggregation requires an explicit representation of the (near) null space of the operator for optimal performance. One can provide an orthonormal set of null space vectors with
MatSetNearNullSpace(). The vector of all ones is the default, for each variable given by the block size (e.g., the translational rigid body modes). For elasticity, where rotational rigid body modes are required to complete the null space you can use MatNullSpaceCreateRigidBody() to create the null space vectors and then MatSetNearNullSpace().

Coarse grid data model: The GAMG framework provides for reducing the number of active processes on coarse grids to reduce communication costs when there is not enough parallelism to keep relative communication costs down. Most AMG solver reduce to just one active process on the coarsest grid (the PETSc MG framework also supports redundantly solving the coarse grid on all processes to potentially reduce communication costs), although this forcing to one process can be overridden if one wishes to use a parallel coarse grid solver. GAMG generalizes this by reducing the active number of processes on other coarse grids as well. GAMG will select the number of active processors by fitting the desired number of equation per process (set with -pc_gamg_process_eq_limit <50>,) at each level given that size of each level. If \( P_i < P \) processes are desired on a level \( i \) then the first \( P_i \) ranks are populated with the grid and the remaining are empty on that grid. One can, and probably should, repartition the coarse grids with -pc_gamg_repartition <true>, otherwise an integer process reduction factor (\( q \)) is selected and the equations on the first \( q \) processes are moved to process 0, and so on. As mentioned multigrid generally coarsens the problem until it is small enough to be solved with an exact solver (eg, LU or SVD) in a relatively small time. GAMG will stop coarsening when the number of equation on a grid falls below at threshold give by -pc_gamg_coarse_eq_limit <50>.

Coarse grid parameters: There are several options to provide parameters to the coarsening algorithm and parallel data layout. Run a code that uses PCGAMG with -help to get full listing of GAMG parameters with short parameter descriptions. The rate of coarsening is critical in AMG performance – too slow coarsening will result in an overly expensive solver per iteration and too fast coarsening will result in decrease in the convergence rate. -pc_gamg_threshold <-1> and -pc_gamg_aggressive_coarsening <false> are the primary parameters that control coarsening rates, which is very important for AMG performance. A greedy maximal independent set (MIS) algorithm is used in coarsening. Squaring the graph implements so called MIS-2, the root vertex in an aggregate is more than two edges away from another root vertex, instead of more than one in MIS. The threshold parameter sets a normalized threshold for which edges are removed from the MIS graph, thereby coarsening slower. Zero will keep all non-zero edges, a negative number will keep zero edges, a positive number will drop small edges. Typical finite threshold values are in the range of 0.01 – 0.05. There are additional parameters for changing the weights on coarse grids.

The parallel MIS algorithms requires symmetric weights/matrix. Thus PCGAMG will automatically make the graph symmetric if it is not symmetric. Since this has additional cost users should indicate the symmetry of the matrices they provide by calling MatSetOption\`(mat,` `MAT_SYMMETRIC,` `PETSC_TRUE`` (or PETSC_FALSE)) or MatSetOption\`(mat, ` `MAT_STRUCTURALLY_SYMMETRIC,` `PETSC_TRUE`` (or PETSC_FALSE)) . If they know that the matrix will always have symmetry, despite future changes to the matrix (with, for example, MatSetValue()) then they should also call MatSetOption\`(mat, ` `MAT_SYMMETRY_ETERNAL,` `PETSC_TRUE`` (or PETSC_FALSE)) or MatSetOption\`(mat, ` `MAT_STRUCTURAL_SYMMETRY_ETERNAL,` `PETSC_TRUE`` (or PETSC_FALSE)). Using this information allows the algorithm to skip the unnecessary computations.

Trouble shooting algebraic multigrid methods: If PCGAMG, ML, AMGx or hypre does not perform well the first thing to try is one of the other methods. Often the default parameters or just the strengths of different algorithms can fix performance problems or provide useful information to guide further debugging. There are several sources of poor performance of AMG solvers and often special purpose methods must be developed to achieve the full potential of multigrid. To name just a few sources of performance degradation that may not be fixed with parameters in PETSc currently: non-elliptic operators, curl/curl operators, highly stretched grids or highly anisotropic problems, large jumps in material coefficients with complex geometry (AMG is particularly well suited to jumps in coefficients but it is not a perfect solution), highly incompressible elasticity, not to mention ill-posed problems, and many others. For Grad-Div and Curl-Curl operators, you may want to try the Auxiliary-space Maxwell Solver (AMS, -pc_type hypre -pc_hypre_type ams) or the Auxiliary-space Divergence Solver (ADS, -pc_type hypre -pc_hypre_type ads) solvers. These solvers need some additional information on the underlying mesh; specifically, AMS needs the discrete
gradient operator, which can be specified via \texttt{PCHYPRESetDiscreteGradient()}. In addition to the
discrete gradient, ADS also needs the specification of the discrete curl operator, which can be set using
\texttt{PCHYPRESetDiscreteCurl()}. 

**I am converging slowly, what do I do?** AMG methods are sensitive to coarsening rates and methods;
for GAMG use \texttt{-pc_gamg_threshold <x> } or \texttt{PCGAMGSetThreshold()} to regulate coarsening rates,
higher values decrease coarsening rate. Squaring the graph is the second mechanism for increasing coarsening
rate. Use \texttt{-pc_gamg_aggressive_coarsening <N>}, or \texttt{PCGAMGSetAggressiveLevels(pc,N)}, to
aggressively coarsen (MIS-2) the graph on the finest N levels. A high threshold \( (e.g., x = 0.08) \)
will result in an expensive but potentially powerful preconditioner, and a low threshold \( (e.g., x = 0.0) \)
will result in faster coarsening, fewer levels, cheaper solves, and generally worse convergence rates.

One can run with \texttt{-info :pc} and grep for \texttt{PCGAMG} to get statistics on each level, which can be used to
see if you are coarsening at an appropriate rate. With smoothed aggregation you generally want to coarse
at about a rate of 3:1 in each dimension. Coarsening too slow will result in large numbers of non-zeros
per row on coarse grids (this is reported). The number of non-zeros can go up very high, say about 300
(times the degrees-of-freedom per vertex) on a 3D hex mesh. One can also look at the grid complexity,
which is also reported (the ratio of the total number of matrix entries for all levels to the number of matrix
entries on the fine level). Grid complexity should be well under 2.0 and preferably around 1.3 or lower.
If convergence is poor and the Galerkin coarse grid construction is much smaller than the time for each solve
then one can safely decrease the coarsening rate. \texttt{-pc_gamg_threshold} \(-1.0\) is the simplest and most
robust option, and is recommended if poor convergence rates are observed, at least until the source of the
problem is discovered. In conclusion, if convergence is slow then decreasing the coarsening rate (increasing
the threshold) should be tried.

**A note on Chebyshev smoothers.** Chebyshev solvers are attractive as multigrid smoothers because they
can target a specific interval of the spectrum which is the purpose of a smoother. The spectral bounds for
Chebyshev solvers are simple to compute because they rely on the highest eigenvalue of your (diagonally
preconditioned) operator, which is conceptually simple to compute. However, if this highest eigenvalue
estimate is not accurate (too low) then the solvers can fail with an indefinite preconditioner message.
One can run with \texttt{-info} and grep for \texttt{PCGAMG} to get these estimates or use \texttt{-ksp_view}.
These highest
eigenvalues are generally between 1.5-3.0. For symmetric positive definite systems CG is a better eigenvalue
estimator \texttt{-mg_levels_esteig_ksp_type cg}. Indefinite matrix messages are often caused by bad Eigen
estimates. Explicitly damped Jacobi or Krylov smoothers can provide an alternative to Chebyshev and \texttt{hypre}
has alternative smoothers.

**Now am I solving alright, can I expect better?** If you find that you are getting nearly one digit in
reduction of the residual per iteration and are using a modest number of point smoothing steps \( (e.g., 1-4
iterations of SOR) \), then you may be fairly close to textbook multigrid efficiency. Although you also need
to check the setup costs. This can be determined by running with \texttt{-log_view} and check that the time for
the Galerkin coarse grid construction \( \texttt{MatPtAP()} \) is not \( (\text{much}) \) more than the time spent in each solve
\( \texttt{KSPSolve()} \). If the \texttt{MatPtAP()} time is too large then one can increase the coarsening rate by decreasing
the threshold and using aggressive coarsening \( \texttt{-pc_gamg_aggressive_coarsening <N>} \), squares the
graph on the finest N levels). Likewise if your \texttt{MatPtAP()} time is small and your convergence rate is not
ideal then you could decrease the coarsening rate.

PETSc’s AMG solver is constructed as a framework for developers to easily add AMG capabilities, like a new
AMG methods or an AMG component like a matrix triple product. Contact us directly if you are interested
in contributing.

It is possible but not recommended to use algebraic multigrid as a “standalone” solver, that is not accelerating
it with a Krylov method. Use a \texttt{KSPType} of \texttt{KSPRICHARDSON} (or equivalently \texttt{-ksp_type richardson}) to
achieve this. Using \texttt{KSPPREONLY} will not work since it only applies a single cycle of multigrid.
Adaptive Interpolation

Interpolation transfers a function from the coarse space to the fine space. We would like this process to be accurate for the functions resolved by the coarse grid, in particular the approximate solution computed there. By default, we create these matrices using local interpolation of the fine grid dual basis functions in the coarse basis. However, an adaptive procedure can optimize the coefficients of the interpolator to reproduce pairs of coarse/fine functions which should approximate the lowest modes of the generalized eigenproblem

\[ Ax = \lambda M x \]

where \( A \) is the system matrix and \( M \) is the smoother. Note that for defect-correction MG, the interpolated solution from the coarse space need not be as accurate as the fine solution, for the same reason that updates in iterative refinement can be less accurate. However, in FAS or in the final interpolation step for each level of Full Multigrid, we must have interpolation as accurate as the fine solution since we are moving the entire solution itself.

Injection should accurately transfer the fine solution to the coarse grid. Accuracy here means that the action of a coarse dual function on either should produce approximately the same result. In the structured grid case, this means that we just use the same values on coarse points. This can result in aliasing.

Restriction is intended to transfer the fine residual to the coarse space. Here we use averaging (often the transpose of the interpolation operation) to damp out the fine space contributions. Thus, it is less accurate than injection, but avoids aliasing of the high modes.

For a multigrid cycle, the interpolator \( P \) is intended to accurately reproduce “smooth” functions from the coarse space in the fine space, keeping the energy of the interpolant about the same. For the Laplacian on a structured mesh, it is easy to determine what these low-frequency functions are. They are the Fourier modes. However an arbitrary operator \( A \) will have different coarse modes that we want to resolve accurately on the fine grid, so that our coarse solve produces a good guess for the fine problem. How do we make sure that our interpolator \( P \) can do this?

We first must decide what we mean by accurate interpolation of some functions. Suppose we know the continuum function \( f \) that we care about, and we are only interested in a finite element description of discrete functions. Then the coarse function representing \( f \) is given by

\[ f^C = \sum_i f_i^C \phi_i^C, \]

and similarly the fine grid form is

\[ f^F = \sum_i f_i^F \phi_i^F. \]

Now we would like the interpolant of the coarse representer to the fine grid to be as close as possible to the fine representer in a least squares sense, meaning we want to solve the minimization problem

\[ \min_P \| f^F - Pf^C \|_2 \]

Now we can express \( P \) as a matrix by looking at the matrix elements \( P_{ij} = \phi_i^F P \phi_j^C \). Then we have

\[ \phi_i^F f^F - \phi_i^F Pf^C = \sum_j P_{ij} f_j^C \]

so that our discrete optimization problem is

\[ \min_{P_{ij}} \| f_i^F - \sum_j P_{ij} f_j^C \|_2 \]
and we will treat each row of the interpolator as a separate optimization problem. We could allow an arbitrary
sparsity pattern, or try to determine adaptively, as is done in sparse approximate inverse preconditioning.
However, we know the supports of the basis functions in finite elements, and thus the naive sparsity pattern
from local interpolation can be used.

We note here that the BAMG framework of Brannick et al. [BBKL11] does not use fine and coarse functions
spaces, but rather a fine point/coarse point division which we will not employ here. Our general PETSc
routine should work for both since the input would be the checking set (fine basis coefficients or fine space
points) and the approximation set (coarse basis coefficients in the support or coarse points in the sparsity
pattern).

We can easily solve the above problem using QR factorization. However, there are many smooth functions
from the coarse space that we want interpolated accurately, and a single \( f \) would not constrain the values
\( P_{ij} \) well. Therefore, we will use several functions \( \{f_k\} \) in our minimization,

\[
\min_{P_{ij}} \sum_k w_k \| f_i^F, k - \sum_j P_{ij} f_j^C, k \|_2
\]

\[
= \min_{P_{ij}} \sum_k \| \sqrt{w_k} f_i^F, k - \sqrt{w_k} \sum_j P_{ij} f_j^C, k \|_2
\]

\[
= \min_{P_{ij}} \| W^{1/2} f_i^F - W^{1/2} f_j^C p_i \|_2
\]

where

\[
W = \begin{pmatrix}
w_0 \\
\vdots \\
w_K
\end{pmatrix}
\]

\[
f_i^F = \begin{pmatrix}
f_i^F, 0 \\
\vdots \\
f_i^F, K
\end{pmatrix}
\]

\[
f_i^C = \begin{pmatrix}
f_i^C, 0 \\
\vdots \\
f_i^C, K
\end{pmatrix}
\]

\[
p_i = \begin{pmatrix}
P_{i0} \\
\vdots \\
P_{inn}
\end{pmatrix}
\]

or alternatively

\[
k_k = w_k
\]

\[
[f_i^F]_k = f_i^F, k
\]

\[
[f_i^C]_j = f_j^C, k
\]

\[
[p_i]_j = P_{ij}
\]

We thus have a standard least-squares problem

\[
\min_{P_{ij}} \| b - Ax \|_2
\]

where

\[
A = W^{1/2} f_i^C
\]

\[
b = W^{1/2} f_i^F
\]

\[
x = p_i
\]
which can be solved using LAPACK.

We will typically perform this optimization on a multigrid level \( l \) when the change in eigenvalue from level \( l + 1 \) is relatively large, meaning

\[
\frac{|\lambda_l - \lambda_{l+1}|}{|\lambda_l|}
\]

This indicates that the generalized eigenvector associated with that eigenvalue was not adequately represented by \( P^l_{l+1} \), and the interpolator should be recomputed.

### Balancing Domain Decomposition by Constraints

PETSc provides the Balancing Domain Decomposition by Constraints (PCBDDC) method for preconditioning parallel finite element problems stored in unassembled format (see MATIS). PCBDDC is a 2-level non-overlapping domain decomposition method which can be easily adapted to different problems and discretizations by means of few user customizations. The application of the preconditioner to a vector consists in the static condensation of the residual at the interior of the subdomains by means of local Dirichlet solves, followed by an additive combination of Neumann local corrections and the solution of a global coupled coarse problem. Command line options for the underlying KSP objects are prefixed by `-pc_bddc_dirichlet`, `-pc_bddc_neumann`, and `-pc_bddc_coarse` respectively.

The current implementation supports any kind of linear system, and assumes a one-to-one mapping between subdomains and MPI processes. Complex numbers are supported as well. For non-symmetric problems, use the runtime option `-pc_bddc_symmetric 0`.

Unlike conventional non-overlapping methods that iterates just on the degrees of freedom at the interface between subdomain, PCBDDC iterates on the whole set of degrees of freedom, allowing the use of approximate subdomain solvers. When using approximate solvers, the command line switches `-pc_bddc_dirichlet_approximate` and/or `-pc_bddc_neumann_approximate` should be used to inform PCBDDC. If any of the local problems is singular, the nullspace of the local operator should be attached to the local matrix via `MatSetNullSpace()`.

At the basis of the method there’s the analysis of the connected components of the interface for the detection of vertices, edges and faces equivalence classes. Additional information on the degrees of freedom can be supplied to PCBDDC by using the following functions:

- PCBDDCSetDofsSplitting()
- PCBDDCSetLocalAdjacencyGraph()
- PCBDDCSetPrimalVerticesLocalIS()
- PCBDDCSetNeumannBoundaries()
- PCBDDCSetDirichletBoundaries()
- PCBDDCSetNeumannBoundariesLocal()
- PCBDDCSetDirichletBoundariesLocal()

Crucial for the convergence of the iterative process is the specification of the primal constraints to be imposed at the interface between subdomains. PCBDDC uses by default vertex continuities and edge arithmetic averages, which are enough for the three-dimensional Poisson problem with constant coefficients. The user can switch on and off the usage of vertices, edges or face constraints by using the command line switches `-pc_bddc_use_vertices`, `-pc_bddc_use_edges`, `-pc_bddc_use_faces`. A customization of the constraints is available by attaching a `MatNullSpace` object to the preconditioning matrix via `MatSetNearNullSpace()`. The vectors of the `MatNullSpace` object should represent the constraints in the form of quadrature rules; quadrature rules for different classes of the interface can be listed in the same vector. The number of vectors of the `MatNullSpace` object corresponds to the maximum number of constraints.

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that can be imposed for each class. Once all the quadrature rules for a given interface class have been extracted, an SVD operation is performed to retain the non-singular modes. As an example, the rigid body modes represent an effective choice for elasticity, even in the almost incompressible case. For particular problems, e.g., edge-based discretization with Nedelec elements, a user defined change of basis of the degrees of freedom can be beneficial for PCBDDC; use PCBDDCSetChangeOfBasisMat() to customize the change of basis.

The PCBDDC method is usually robust with respect to jumps in the material parameters aligned with the interface; for PDEs with more than one material parameter you may also consider to use the so-called deluxe scaling, available via the command line switch -pc_bddc_use_deluxe_scaling. Other scalings are available, see PCISSetSubdomainScalingFactor(), PCISSetSubdomainDiagonalScaling() or PCISSetUseStiffnessScaling(). However, the convergence properties of the PCBDDC method degrades in presence of large jumps in the material coefficients not aligned with the interface; for such cases, PETSc has the capability of adaptively computing the primal constraints. Adaptive selection of constraints could be requested by specifying a threshold value at command line by using -pc_bddc_adaptive_threshold x. Valid values for the threshold x ranges from 1 to infinity, with smaller values corresponding to more robust preconditioners. For SPD problems in 2D, or in 3D with only face degrees of freedom (like in the case of Raviart-Thomas or Brezzi-Douglas-Marini elements), such a threshold is a very accurate estimator of the condition number of the resulting preconditioned operator. Since the adaptive selection of constraints for PCBDDC methods is still an active topic of research, its implementation is currently limited to SPD problems; moreover, because the technique requires the explicit knowledge of the local Schur complements, it needs the external package MUMPS.

When solving problems decomposed in thousands of subdomains or more, the solution of the PCBDDC coarse problem could become a bottleneck; in order to overcome this issue, the user could either consider to solve the parallel coarse problem on a subset of the communicator associated with PCBDDC by using the command line switch -pc_bddc_coarse_redistribute, or instead use a multilevel approach. The latter can be requested by specifying the number of requested level at command line (-pc_bddc_levels) or by using PCBDDCSetLevels(). An additional parameter (see PCBDDCSetCoarseningRatio()) controls the number of subdomains that will be generated at the next level; the larger the coarsening ratio, the lower the number of coarser subdomains.

For further details, see the example KSP Tutorial ex59 and the online documentation for PCBDDC.

Shell Preconditioners

The shell preconditioner simply uses an application-provided routine to implement the preconditioner. To set this routine, one uses the command

```c
PCShellSetApply(PC pc,PetscErrorCode (*apply)(PC,Vec,Vec));
```

Often a preconditioner needs access to an application-provided data structured. For this, one should use

```c
PCShellSetContext(PC pc,void *ctx);
```

to set this data structure and

```c
PCShellGetContext(PC pc,void *ctx);
```

to retrieve it in apply. The three routine arguments of apply() are the PC, the input vector, and the output vector, respectively.

For a preconditioner that requires some sort of “setup” before being used, that requires a new setup every time the operator is changed, one can provide a routine that is called every time the operator is changed (usually via KSPSetOperators()).
PCShellSetSetUp(PC pc, PetscErrorCode (*setup)(PC));

The argument to the setup routine is the same PC object which can be used to obtain the operators with PCGetOperators() and the application-provided data structure that was set with PCShellSetContext().

Combining Preconditioners

The PC type PCCOMPOSITE allows one to form new preconditioners by combining already-defined preconditioners and solvers. Combining preconditioners usually requires some experimentation to find a combination of preconditioners that works better than any single method. It is a tricky business and is not recommended until your application code is complete and running and you are trying to improve performance. In many cases using a single preconditioner is better than a combination; an exception is the multigrid/multilevel preconditioners (solvers) that are always combinations of some sort, see Multigrid Preconditioners.

Let $B_1$ and $B_2$ represent the application of two preconditioners of type type1 and type2. The preconditioner $B = B_1 + B_2$ can be obtained with

```c
PCSetType(pc, PCCOMPOSITE);
PCCompositeAddPCType(pc, type1);
PCCompositeAddPCType(pc, type2);
```

Any number of preconditioners may added in this way.

This way of combining preconditioners is called additive, since the actions of the preconditioners are added together. This is the default behavior. An alternative can be set with the option

```c
PCCompositeSetType(pc, PC_COMPOSITE_MULTIPLICATIVE);
```

In this form the new residual is updated after the application of each preconditioner and the next preconditioner applied to the next residual. For example, with two composed preconditioners: $B_1$ and $B_2$: $y = Bx$ is obtained from

\[
\begin{align*}
y &= B_1x \\
w_1 &= x - Ay \\
y &= y + B_2w_1
\end{align*}
\]

Loosely, this corresponds to a Gauss-Seidel iteration, while additive corresponds to a Jacobi iteration.

Under most circumstances, the multiplicative form requires one-half the number of iterations as the additive form; however, the multiplicative form does require the application of $A$ inside the preconditioner.

In the multiplicative version, the calculation of the residual inside the preconditioner can be done in two ways: using the original linear system matrix or using the matrix used to build the preconditioners $B_1$, $B_2$, etc. By default it uses the “preconditioner matrix”, to use the Amat matrix use the option

```c
PCSetUseAmat(PC pc);
```

The individual preconditioners can be accessed (in order to set options) via

```c
PCCompositeGetPC(PC pc, PetscInt count, PC *subpc);
```

For example, to set the first sub preconditioners to use ILU(1)

```c
PC subpc;
PCCompositeGetPC(pc, 0, &subpc);
PCFactorSetFill(subpc, 1);
```
One can also change the operator that is used to construct a particular PC in the composite PC call
PCSetOperators() on the obtained PC.

These various options can also be set via the options database. For example, -pc_type composite
-pc_composite_pcs jacobi,ilu causes the composite preconditioner to be used with two precondition-
ers: Jacobi and ILU. The option -pc_composite_type multiplicative initiates the multiplicative version
of the algorithm, while -pc_composite_type additive the additive version. Using the Amat matrix
is obtained with the option -pc_use_amat. One sets options for the sub-preconditioners with the extra pre-
fix -sub_N_ where N is the number of the sub-preconditioner. For example, -sub_0_pc_ifactor_fill 0.

PETSc also allows a preconditioner to be a complete linear solver. This is achieved with the PCKSP type.

```c
PCSetType(PC pc,PCKSP PCKSP);
PCKSPGetKSP(pc,&ksp);
/* set any KSP/PC options */
```

From the command line one can use 5 iterations of biCG-stab with ILU(0) preconditioning as the precondi-
tioner with -pc_type ksp -ksp_pc_type ilu -ksp_ksp_max_it 5 -ksp_ksp_type bcgs.

By default the inner KSP solver uses the outer preconditioner matrix, Pmat, as the matrix to be solved in
the linear system; to use the matrix that defines the linear system, Amat use the option

```c
PCSetUseAmat(PC pc);
```
or at the command line with -pc_use_amat.

Naturally, one can use a PCKSP preconditioner inside a composite preconditioner. For example, -pc_type composite
-pc_composite_pcs ilu,ksp -sub_1_pc_type jacobi -sub_1_ksp_max_it 10 uses two preconditioners: ILU(0) and 10 iterations of GMRES with Jacobi preconditioning. However, it is
not clear whether one would ever wish to do such a thing.

### Multigrid Preconditioners

A large suite of routines is available for using geometric multigrid as a preconditioner\(^2\). In the PC framework,
the user is required to provide the coarse grid solver, smoothers, restriction and interpolation operators, and
code to calculate residuals. The PC package allows these components to be encapsulated within a PETSc-
compliant preconditioner. We fully support both matrix-free and matrix-based multigrid solvers.

A multigrid preconditioner is created with the four commands

```c
KSPCreate(MPI_Comm comm,KSP *ksp);
KSPGetPC(KSP ksp,PC *pc);
PCSetType(PC pc,PCMG);
PCMGSetLevels(pc,PetscInt levels,MPI_Comm *comms);
```

A large number of parameters affect the multigrid behavior. The command

```c
PCMGSetType(PC pc,PCMGType mode);
```

indicates which form of multigrid to apply [SBjorstadG96].

For standard V or W-cycle multigrids, one sets the mode to be PC_MG_MULTIPLICATIVE; for the additive form
(which in certain cases reduces to the BPX method, or additive multilevel Schwarz, or multilevel
diagonal scaling), one uses PC_MG_ADDITIVE as the mode. For a variant of full multigrid, one can use
PC_MG_FULL, and for the Kaskade algorithm PC_MG_KASKADE. For the multiplicative and full multigrid
options, one can use a W-cycle by calling

\(^2\) See Algebraic Multigrid (AMG) Preconditioners for information on using algebraic multigrid.
with a value of `PC_MG_CYCLE_W` for `ctype`. The commands above can also be set from the options database. The option names are `-pc_mg_type` [multiplicative, additive, full, kaskade], and `-pc_mg_cycle_type <ctype>`.

The user can control the amount of smoothing by configuring the solvers on the levels. By default, the up and down smoothers are identical. If separate configuration of up and down smooths is required, it can be requested with the option `-pc_mg_distinct_smoothup` or the routine

```c
PCMGSetDistinctSmoothUp(PC pc);
```

The multigrid routines, which determine the solvers and interpolation/restriction operators that are used, are mandatory. To set the coarse grid solver, one must call

```c
PCMGGetCoarseSolve(PC pc,KSP *ksp);
```

and set the appropriate options in `ksp`. Similarly, the smoothers are controlled by first calling

```c
PCMGGetSmoother(PC pc,PetscInt level,KSP *ksp);
```

and then setting the various options in the `ksp`. For example,

```c
PCMGGetSmoother(pc,1,&ksp);
KSPSetOperators(ksp,A1,A1);
```

sets the matrix that defines the smoother on level 1 of the multigrid. While

```c
PCMGGetSmoother(pc,1,&ksp);
KSPGetPC(ksp,&pc);
PCSetType(pc,PCSOR);
```

sets SOR as the smoother to use on level 1.

To use a different pre- or postsmoother, one should call the following routines instead.

```c
PCMGGetSmootherUp(PC pc,PetscInt level,KSP *upksp);
PCMGGetSmootherDown(PC pc,PetscInt level,KSP *downksp);
```

Use

```c
PCMGSetInterpolation(PC pc,PetscInt level,Mat P);
```

and

```c
PCMGSetRestriction(PC pc,PetscInt level,Mat R);
```

to define the intergrid transfer operations. If only one of these is set, its transpose will be used for the other.

It is possible for these interpolation operations to be matrix-free (see `Matrix-Free Matrices`); One should then make sure that these operations are defined for the (matrix-free) matrices passed in. Note that this system is arranged so that if the interpolation is the transpose of the restriction, you can pass the same `mat` argument to both `PCMGSetRestriction()` and `PCMGSetInterpolation()`.

On each level except the coarsest, one must also set the routine to compute the residual. The following command suffices:
The `residual()` function normally does not need to be set if one's operator is stored in `Mat` format. In certain circumstances, where it is much cheaper to calculate the residual directly, rather than through the usual formula $b - Ax$, the user may wish to provide an alternative.

Finally, the user may provide three work vectors for each level (except on the finest, where only the residual work vector is required). The work vectors are set with the commands

```c
PCMGSetRhs(PC pc, PetscInt level, Vec b);
PCMGSetX(PC pc, PetscInt level, Vec x);
PCMGSetR(PC pc, PetscInt level, Vec r);
```

The `PC` references these vectors, so you should call `VecDestroy()` when you are finished with them. If any of these vectors are not provided, the preconditioner will allocate them.

One can control the `KSP` and `PC` options used on the various levels (as well as the coarse grid) using the prefix `mg_levels_` (mg_coarse_ for the coarse grid). For example, `-mg_levels_ksp_type cg` will cause the CG method to be used as the Krylov method for each level. Or `-mg_levels_pc_type ilu -mg_levels_pc_factor_levels 2` will cause the ILU preconditioner to be used on each level with two levels of fill in the incomplete factorization.

### 2.3.5 Solving Block Matrices

Block matrices represent an important class of problems in numerical linear algebra and offer the possibility of far more efficient iterative solvers than just treating the entire matrix as black box. In this section we use the common linear algebra definition of block matrices where matrices are divided in a small, problem-size independent (two, three or so) number of very large blocks. These blocks arise naturally from the underlying physics or discretization of the problem, for example, the velocity and pressure. Under a certain numbering of unknowns the matrix can be written as

\[
\begin{pmatrix}
  A_{00} & A_{01} & A_{02} & A_{03} \\
  A_{10} & A_{11} & A_{12} & A_{13} \\
  A_{20} & A_{21} & A_{22} & A_{23} \\
  A_{30} & A_{31} & A_{32} & A_{33}
\end{pmatrix},
\]

where each $A_{ij}$ is an entire block. On a parallel computer the matrices are not explicitly stored this way. Instead, each process will own some of the rows of $A_{0*}$, $A_{1*}$, etc. On a process, the blocks may be stored one block followed by another

\[
\begin{pmatrix}
  A_{0000} & A_{0001} & A_{0002} & \cdots & A_{0100} & A_{0102} & \cdots \\
  A_{0010} & A_{0011} & A_{0012} & \cdots & A_{0110} & A_{0112} & \cdots \\
  A_{0020} & A_{0021} & A_{0022} & \cdots & A_{0120} & A_{0122} & \cdots \\
  \vdots & \vdots & \vdots & \ddots & \vdots & \vdots & \ddots \\
  A_{1000} & A_{1001} & A_{1002} & \cdots & A_{1100} & A_{1102} & \cdots \\
  A_{1010} & A_{1011} & A_{1012} & \cdots & A_{1110} & A_{1112} & \cdots \\
  \vdots & \vdots & \vdots & \ddots & \vdots & \vdots & \ddots
\end{pmatrix}
\]

or interlaced, for example with two blocks

\[
\begin{pmatrix}
  A_{0000} & A_{0100} & A_{0001} & A_{0101} & \cdots \\
  A_{1000} & A_{1100} & A_{1001} & A_{1101} & \cdots \\
  \vdots & \vdots & \vdots & \vdots & \ddots
\end{pmatrix}
\]

or

\[
\begin{pmatrix}
  A_{0000} & A_{0100} & A_{0001} & A_{0101} & \cdots \\
  A_{1000} & A_{1100} & A_{1001} & A_{1101} & \cdots \\
  \vdots & \vdots & \vdots & \vdots & \ddots
\end{pmatrix}
\]
Note that for interlaced storage the number of rows/columns of each block must be the same size. Matrices obtained with `DMCreateMatrix()` where the DM is a DMDA are always stored interlaced. Block matrices can also be stored using the MATNEST format which holds separate assembled blocks. Each of these nested matrices is itself distributed in parallel. It is more efficient to use MATNEST with the methods described in this section because there are fewer copies and better formats (e.g. MATBAIJ or MATSBAIJ) can be used for the components, but it is not possible to use many other methods with MATNEST. See Block Matrices for more on assembling block matrices without depending on a specific matrix format.

The PETSc PCFIELDSPLIT preconditioner is used to implement the “block” solvers in PETSc. There are three ways to provide the information that defines the blocks. If the matrices are stored as interlaced then `PCFieldSplitSetFields()` can be called repeatedly to indicate which fields belong to each block. More generally `PCFieldSplitSetIS()` can be used to indicate exactly which rows/columns of the matrix belong to a particular block. You can provide names for each block with these routines, if you do not provide names they are numbered from 0. With these two approaches the blocks may overlap (though generally they will not). If only one block is defined then the complement of the matrices is used to define the other block. Finally the option `-pc_fieldsplit_detect_saddle_point` causes two diagonal blocks to be found, one associated with all rows/columns that have zeros on the diagonals and the rest.

For simplicity in the rest of the section we restrict our matrices to two-by-two blocks. So the matrix is

\[
\begin{pmatrix}
A_{00} & A_{01} \\
A_{10} & A_{11}
\end{pmatrix}.
\]

On occasion the user may provide another matrix that is used to construct parts of the preconditioner

\[
\begin{pmatrix}
A_{p00} & A_{p01} \\
A_{p10} & A_{p11}
\end{pmatrix}.
\]

For notational simplicity define \(ksp(A, Ap)\) to mean approximately solving a linear system using KSP with operator \(A\) and preconditioner built from matrix \(Ap\).

For matrices defined with any number of blocks there are three “block” algorithms available: block Jacobi,

\[
\begin{pmatrix}
ksp(A_{00}, Ap_{00}) & 0 \\
0 & ksp(A_{11}, Ap_{11})
\end{pmatrix}
\]

block Gauss-Seidel,

\[
\begin{pmatrix}
I & 0 \\
0 & A_{11}^{-1}
\end{pmatrix}
\begin{pmatrix}
I & 0 \\
0 & I
\end{pmatrix}
\begin{pmatrix}
A_{10}^{-1} & 0 \\
0 & I
\end{pmatrix}
\]

which is implemented\(^3\) as

\[
\begin{pmatrix}
I \\
0 & ksp(A_{11}, Ap_{11})
\end{pmatrix}
\begin{pmatrix}
I \\
0 & I
\end{pmatrix}
+ \begin{pmatrix}
I & 0 \\
0 & I
\end{pmatrix}
\begin{pmatrix}
I & 0 \\
0 & I
\end{pmatrix}
\begin{pmatrix}
A_{00}^{-1} & 0 \\
0 & I
\end{pmatrix}
\]

and symmetric block Gauss-Seidel

\[
\begin{pmatrix}
A_{00}^{-1} & 0 \\
0 & I
\end{pmatrix}
\begin{pmatrix}
I & -A_{01} \\
0 & I
\end{pmatrix}
\begin{pmatrix}
A_{00} & 0 \\
0 & A_{11}^{-1}
\end{pmatrix}
\begin{pmatrix}
I & 0 \\
0 & I
\end{pmatrix}
\begin{pmatrix}
A_{00}^{-1} & 0 \\
0 & I
\end{pmatrix}
\]

These can be accessed with `-pc_fieldsplit_type<additive,multiplicative, symmetric_multiplicative>` or the function `PCFieldSplitSetType()`. The option prefixes for the internal KSPs are given by `-fieldsplit_name_`.

By default blocks \(A_{00}, A_{01}\) and so on are extracted out of `Pmat`, the matrix that the KSP uses to build the preconditioner, and not out of `Amat` (i.e., \(A\) itself). As discussed above in Combining Preconditioners,\(^3\) This may seem an odd way to implement since it involves the “extra” multiply by \(-A_{11}\). The reason is this is implemented this way is that this approach works for any number of blocks that may overlap.

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however, it is possible to use \texttt{Amat} instead of \texttt{Pmat} by calling \texttt{PCSetUseAmat(pc)} or using \texttt{-pc_use_amat} on the command line. Alternatively, you can have \texttt{PCFIELDSPLIT} extract the diagonal blocks $A_{00}, A_{11}$ etc. out of \texttt{Amat} by calling \texttt{PCFieldSplitSetDiagUseAmat(pc,\texttt{PETSC\_TRUE})} or supplying command-line argument \texttt{-pc_fieldsplit_diag_use_amat}. Similarly, \texttt{PCFieldSplitSetOffDiagUseAmat(pc,\{\texttt{PETSC\_TRUE}\} or \texttt{-pc_fieldsplit_off_diag_use_amat} will cause the off-diagonal blocks $A_{01}, A_{10}$ etc. to be extracted out of \texttt{Amat}.

For two-by-two blocks only, there is another family of solvers, based on Schur complements. The inverse of the Schur complement factorization is

$$
\begin{pmatrix}
I & A_{10}^{-1}A_{00}^{-1} \\
A_{10}^{-1}A_{00}^{-1} & I
\end{pmatrix}
\begin{pmatrix}
A_{00} & 0 \\
0 & S
\end{pmatrix}
\begin{pmatrix}
I & A_{00}^{-1}A_{01} \\
0 & I
\end{pmatrix}
$$

The preconditioner is accessed with \texttt{-pc_fieldsplit_type schur} and is implemented as

$$
\begin{pmatrix}
ksp(A_{00}, A_{p00}) & 0 \\
0 & I
\end{pmatrix}
\begin{pmatrix}
I & -A_{01} \\
I & -A_{10}ksp(\hat{S}, \hat{Sp})
\end{pmatrix}
\begin{pmatrix}
ksp(A_{00}, A_{p00}) & 0 \\
0 & -A_{10}ksp(\hat{S}, \hat{Sp})
\end{pmatrix}.
$$

Where $\hat{S} = A_{11} - A_{10}ksp(A_{00}, A_{p00})A_{01}$ is the approximate Schur complement.

There are several variants of the Schur complement preconditioner obtained by dropping some of the terms, these can be obtained with \texttt{-pc_fieldsplit_schur_fact_type <diag,lower,upper,full>} or the function \texttt{PCFieldSplitSetSchurFactType()}. Note that the \texttt{diag} form uses the preconditioner

$$
\begin{pmatrix}
ksp(A_{00}, A_{p00}) & 0 \\
0 & -ksp(\hat{S}, \hat{Sp})
\end{pmatrix}.
$$

This is done to ensure the preconditioner is positive definite for a common class of problems, saddle points with a positive definite $A_{00}$: for these the Schur complement is negative definite.

The effectiveness of the Schur complement preconditioner depends on the availability of a good preconditioner $\hat{Sp}$ for the Schur complement matrix. In general, you are responsible for supplying $\hat{Sp}$ via \texttt{PCFieldSplitSetSchurPre(pc,PC\_FIELDSPLIT\_SCHUR\_PRE\_USER,Sp)}. In the absence of a good problem-specific $\hat{Sp}$, you can use some of the built-in options.

Using \texttt{-pc_fieldsplit_schur_precondition user} on the command line activates the matrix supplied programmatically as explained above.

With \texttt{-pc_fieldsplit_schur_precondition all} (default) $\hat{Sp} = A_{11}$ is used to build a preconditioner for $\hat{S}$.

Otherwise, \texttt{-pc_fieldsplit_schur_precondition self} will set $\hat{Sp} = \hat{S}$ and use the Schur complement matrix itself to build the preconditioner.

The problem with the last approach is that $\hat{S}$ is used in unassembled, matrix-free form, and many preconditioners (e.g., ILU) cannot be built out of such matrices. Instead, you can assemble an approximation to $\hat{S}$ by inverting $A_{00}$, but only approximately, so as to ensure the sparsity of $\hat{Sp}$ as much as possible. Specifically, using \texttt{-pc_fieldsplit_schur_precondition selfp} will assemble $\hat{Sp} = A_{11} - A_{10}\inv(A_{00})A_{01}$.

By default $\inv(A_{00})$ is the inverse of the diagonal of $A_{00}$, but using \texttt{-fieldsplit_1_mat_schur_complement_ainv_type lump} will lump $A_{00}$ first. Using
-fieldsplit_1_mat_schur_complement_ainv_type blockdiag will use the inverse of the block diagonal of \( A_{00} \). Option -mat_schur_complement_ainv_type applies to any matrix of MatSchurComplement type and here it is used with the prefix -fieldsplit_1 of the linear system in the second split.

Finally, you can use the PCLSC preconditioner for the Schur complement with -pc_fieldsplit_type schur -fieldsplit_1_pc_type lsc. This uses for the preconditioner to \( \hat{S} \) the operator

\[
ksp(A_{10}A_{01})A_{10}A_{00}A_{01}ksp(A_{10}A_{01}, A_{10}A_{01})
\]

which, of course, introduces two additional inner solves for each application of the Schur complement. The options prefix for this inner KSP is -fieldsplit_1_lsc_. Instead of constructing the matrix \( A_{10}A_{01} \) the user can provide their own matrix. This is done by attaching the matrix/matrices to the \( Sp \) matrix they provide with

```c
PetscObjectCompose((PetscObject)Sp,"LSC_L",(PetscObject)L);
PetscObjectCompose((PetscObject)Sp,"LSC_Lp",(PetscObject)Lp);
```

### 2.3.6 Solving Singular Systems

Sometimes one is required to solve singular linear systems. In this case, the system matrix has a non-trivial null space. For example, the discretization of the Laplacian operator with Neumann boundary conditions has a null space of the constant functions. PETSc has tools to help solve these systems. This approach is only guaranteed to work for left preconditioning (see KSPSetPCSide()); for example it may not work in some situations with KSPFGMRES.

First, one must know what the null space is and store it using an orthonormal basis in an array of PETSc Vecs. The constant functions can be handled separately, since they are such a common case. Create a MatNullSpace object with the command

```c
MatNullSpaceCreate(MPI_Comm,PetscBool hasconstants,PetscInt dim,Vec *basis, MatNullSpace *nsp);
```

Here, \( \text{dim} \) is the number of vectors in \( \text{basis} \) and \( \text{hasconstants} \) indicates if the null space contains the constant functions. If the null space contains the constant functions you do not need to include it in the \( \text{basis} \) vectors you provide, nor in the count \( \text{dim} \).

One then tells the KSP object you are using what the null space is with the call

```c
MatSetNullSpace(Mat Amat,MatNullSpace nsp);
```

The \( \text{Amat} \) should be the first matrix argument used with KSPSetOperators(), SNESSetJacobian(), or TSSetIJacobian(). The PETSc solvers will now handle the null space during the solution process.

If the right hand side of linear system is not in the range of \( \text{Amat} \), that is it is not orthogonal to the null space of \( \text{Amat} \) transpose, then the residual norm of the Krylov iteration will not converge to zero; it will converge to a non-zero value while the solution is converging to the least squares solution of the linear system. One can, if one desires, apply MatNullSpaceRemove() with the null space of \( \text{Amat} \) transpose to the right hand side before calling KSPSolve(). Then the residual norm will converge to zero.

If one chooses a direct solver (or an incomplete factorization) it may still detect a zero pivot. You can run with the additional options or -pc_factor_shift_type NONZERO -pc_factor_shift_amount <dampingfactor> to prevent the zero pivot. A good choice for the dampingfactor is 1.e-10.

If the matrix is non-symmetric and you wish to solve the transposed linear system you must provide the null space of the transposed matrix with MatSetTransposeNullSpace().
2.3.7 Using External Linear Solvers

PETSc interfaces to several external linear solvers (also see acknowledgements). To use these solvers, one may:

1. Run `configure` with the additional options `--download-packagename` e.g. `--download-superlu_dist --download-parmetis` (SuperLU_DIST needs ParMetis) or `--download-mumps --download-scalapack` (MUMPS requires ScaLAPACK).

2. Build the PETSc libraries.

3. Use the runtime option: `-ksp_type preonly` (or equivalently `-ksp_type none`) `-pc_type <pctype> -pc_factor_mat_solver_type <packagename>`. For eg: `-ksp_type preonly -pc_type lu -pc_factor_mat_solver_type superlu_dist`. 
<table>
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</tr>
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The default and available input options for each external software can be found by specifying -help at runtime.

As an alternative to using runtime flags to employ these external packages, procedural calls are provided for some packages. For example, the following procedural calls are equivalent to runtime options -ksp_type preonly (or equivalently -ksp_type none) -pc_type lu -pc_factor_mat_solver_type mumps -mat_mumps_icntl_7 3:

```c
KSPSetType(ksp, KSPPREONLY); (or equivalently KSPSetType(ksp, KSPNONE))
KSPGetPC(ksp, &pc);
PCSetType(pc, PCLU);
PCFactorSetMatSolverType(pc, MATSOLVERMUMPS);
PCFactorSetUpMatSolverType(pc);
PCFactorGetMatrix(pc, &F);
icntl = 7; ival = 3;
MatMumpsSetIcntl(F, icntl, ival);
```

One can also create matrices with the appropriate capabilities by calling MatCreate() followed by MatSetType() specifying the desired matrix type from Options for External Solvers. These matrix types inherit capabilities from their PETSc matrix parents: MATSEQAIJ, MATMPIAIJ, etc. As a result, the preallocation routines MatSeqAIJSetPreallocation(), MatMPIAIJSetPreallocation(), etc. and any other type specific routines of the base class are supported. One can also call MatConvert() inplace to convert the matrix to and from its base class without performing an expensive data copy. MatConvert() cannot be called on matrices that have already been factored.

In Options for External Solvers, the base class aij refers to the fact that inheritance is based on MATSEQAIJ when constructed with a single process communicator, and from MATMPIAIJ otherwise. The same holds for baij and sbaij. For codes that are intended to be run as both a single process or with multiple processes, depending on the mpiexec command, it is recommended that both sets of preallocation routines are called for these communicator morphing types. The call for the incorrect type will simply be ignored without any harm or message.

### 2.3.8 Using PETSc’s MPI parallel linear solvers from a non-MPI program

Using PETSc’s MPI linear solver server it is possible to use multiple MPI processes to solve a linear system when the application code, including the matrix generation, is run on a single MPI rank (with or without OpenMP). The application code must be built with MPI and must call PetscInitialize() at the very beginning of the program and end with PetscFinalize(). The application code may utilize OpenMP. The code may create multiple matrices and KSP objects and call KSPSolve(), similarly the code may utilize the SNES nonlinear solvers, the TS ODE integrators, and the Tao optimization algorithms which use KSP.

The program must then be launched using the standard approaches for launching MPI programs with the additional PETSc option -mpi_linear_solver_server. The linear solves are controlled via the options database in the usual manner (using any options prefix you may have provided via KSPSetOptionsPrefix(), for example -ksp_type cg -ksp_monitor -pc_type bjacobi -ksp_view. The solver options cannot be set via the functional interface, for example KSPSetType() etc.

The option -mpi_linear_solver_server_view will print a summary of all the systems solved by the MPI linear solver server when the program completes. By default the linear solver server will only parallelize the linear solve to the extent that it believes is appropriate to obtain speedup for the parallel solve, for example, if the matrix has 1,000 rows and columns the solution will not be parallelized by default. One can use the option -mpi_linear_solver_server_minimum_count_per_rank 5000 to cause the linear solver server to allow as few as 5,000 unknowns per rank in the parallel solve.
See PCMPI, PCMPIservertBegin(), and PCMPIservertEnd() for more details on the solvers.

Amdahl’s law makes clear that parallelizing only a portion of a numerical code can only provide a limited improvement in the computation time; thus it is crucial to understand what phases of a computation must be parallelized (via MPI, OpenMP, or some other model) to ensure a useful increase in performance. One of the crucial phases is likely the generation of the matrix entries; the use of MatSetPreallocationCOO() and MatSetValuesCOO() in an OpenMP code allows parallelizing the generation of the matrix.

References

2.4 SNES: Nonlinear Solvers

Note: This chapter is being cleaned up by Jed Brown. Contributions are welcome.

The solution of large-scale nonlinear problems pervades many facets of computational science and demands robust and flexible solution strategies. The SNES library of PETSc provides a powerful suite of data-structure-neutral numerical routines for such problems. Built on top of the linear solvers and data structures discussed in preceding chapters, SNES enables the user to easily customize the nonlinear solvers according to the application at hand. Also, the SNES interface is identical for the uniprocess and parallel cases; the only difference in the parallel version is that each process typically forms only its local contribution to various matrices and vectors.

The SNES class includes methods for solving systems of nonlinear equations of the form

\[ F(x) = 0, \]  

where \( F : \mathbb{R}^n \rightarrow \mathbb{R}^n \). Newton-like methods provide the core of the package, including both line search and trust region techniques. A suite of nonlinear Krylov methods and methods based upon problem decomposition are also included. The solvers are discussed further in The Nonlinear Solvers. Following the PETSc design philosophy, the interfaces to the various solvers are all virtually identical. In addition, the SNES software is completely flexible, so that the user can at runtime change any facet of the solution process.

PETSc’s default method for solving the nonlinear equation is Newton’s method. The general form of the \( n \)-dimensional Newton’s method for solving (2.3) is

\[ x_{k+1} = x_k - J(x_k)^{-1}F(x_k), \quad k = 0, 1, \ldots, \]  

where \( x_0 \) is an initial approximation to the solution and \( J(x_k) = F'(x_k) \), the Jacobian, is nonsingular at each iteration. In practice, the Newton iteration (2.4) is implemented by the following two steps:

1. (Approximately) solve \( J(x_k)\Delta x_k = -F(x_k) \).
2. Update \( x_{k+1} \leftarrow x_k + \Delta x_k \).

Other defect-correction algorithms can be implemented by using different choices for \( J(x_k) \).
2.4.1 Basic SNES Usage

In the simplest usage of the nonlinear solvers, the user must merely provide a C, C++, or Fortran routine to evaluate the nonlinear function (2.3). The corresponding Jacobian matrix can be approximated with finite differences. For codes that are typically more efficient and accurate, the user can provide a routine to compute the Jacobian; details regarding these application-provided routines are discussed below. To provide an overview of the use of the nonlinear solvers, browse the concrete example in `ex1.c` or skip ahead to the discussion.

Listing: src/snes/tutorials/ex1.c

```c
static char help[] = "Newton's method for a two-variable system, sequential.\n\n"

/* Include "petscsnes.h" so that we can use SNES solvers. Note that this
   file automatically includes:
   petscsys.h - base PETSc routines      petscvect.h - vectors
   petscmat.h - matrices
   petscisc.h - index sets               petscksp.h - Krylov subspace methods
   petscvviewer.h - viewers              petscpc.h - preconditioners
   petscksp.h - linear solvers
*/

/* This examples solves either
\begin{equation}
   F(x_0, x_1) = \begin{cases}
   x_0^2 + x_0 x_1 - 3
   & x_0 x_1 - 1 + x_1^2 - 6 \\
   \sin(3 x_0) + x_0
   & x_1
   \end{cases}
\end{equation}
   or if the \texttt{-hard} options is given
\begin{equation}
   F(x_0, x_1) = \begin{cases}
   \sin(3 x_0) + x_0
   & x_1
   \end{cases}
\end{equation}
*/
#include <petscsnes.h>

/* User-defined routines */
extern PetscErrorCode FormJacobian1(SNES, Vec, Mat, Mat, void *);
extern PetscErrorCode FormFunction1(SNES, Vec, Vec, void *);
extern PetscErrorCode FormJacobian2(SNES, Vec, Mat, Mat, void *);
extern PetscErrorCode FormFunction2(SNES, Vec, Vec, void *);

int main(int argc, char **argv)
{
    SNES snes; /* nonlinear solver context */
    KSP ksp; /* linear solver context */
    PC pc; /* preconditioner context */
    Vec x, r; /* solution, residual vectors */
    Mat J; /* Jacobian matrix */
    PetscMPIInt size;
    PetscScalar pfive = .5, *xx;
    PetscBool flg;

    PetscFunctionBeginUser;
    PetscCall(PetscInitialize(&argc, &argv, (char *)0, help));
    PetscCallMPI(MPI_Comm_size(PETSC_COMM_WORLD, &size));

    // Code execution...

    PetscFunctionEnd;
}
```

(continues on next page)
PetscCheck(size == 1, PETSC_COMM_WORLD, PETSC_ERR_WRONG_MPI_SIZE, "Example is only for sequential runs");

/* Create nonlinear solver context */
PetscCall(SNESCreate(PETSC_COMM_WORLD, &snes));
PetscCall(SNESSetType(snes, SNESNEWTONLS));
PetscCall(SNESSetOptionsPrefix(snes, "mysolver_"));

/* Create matrix and vector data structures; set corresponding routines */

/* Create vectors for solution and nonlinear function */
PetscCall(VecCreate(PETSC_COMM_WORLD, &x));
PetscCall(VecSetSizes(x, PETSC_DECIDE, 2));
PetscCall(VecSetFromOptions(x));
PetscCall(VecDuplicate(x, &r));

/* Create Jacobian matrix data structure */
PetscCall(MatCreate(PETSC_COMM_WORLD, &J));
PetscCall(MatSetSizes(J, PETSC_DECIDE, PETSC_DECIDE, 2, 2));
PetscCall(MatSetFromOptions(J));
PetscCall(MatSetUp(J));

PetscCall(PetscOptionsHasName(NULL, NULL, "-hard", &flg));
if (!flg) {
  /* Set function evaluation routine and vector. */
  PetscCall(SNESSetFunction(snes, r, FormFunction1, NULL));
  /* Set Jacobian matrix data structure and Jacobian evaluation routine */
  PetscCall(SNESSetJacobian(snes, J, J, FormJacobian1, NULL));
} else {
  PetscCall(SNESSetFunction(snes, r, FormFunction2, NULL));
  PetscCall(SNESSetJacobian(snes, J, J, FormJacobian2, NULL));
}

/* Customize nonlinear solver; set runtime options */

/* Set linear solver defaults for this problem. By extracting the KSP and PC contexts from the SNES context, we can then directly call any KSP and PC routines to set various options. */
PetscCall(SNESGetKSP(snes, &ksp));
PetscCall(KSPGetPC(ksp, &pc));
PetscCall(PCSetType(pc, PCNONE));
PetscCall(KSPSetTolerances(ksp, 1.e-4, PETSC_DEFAULT, PETSC_DEFAULT, 20));

/*
Set SNES/KSP/KSP/PC runtime options, e.g.,
   -snes_view -snes_monitor -ksp_type <ksp> -pc_type <pc>
These options will override those specified above as long as
SNESSetFromOptions() is called _after_ any other customization
routines.
*/
PetscCall(SNESSetFromOptions(snes));

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
Evaluate initial guess; then solve nonlinear system
- - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -*/
if (!flg) {
   PetscCall(VecSet(x, pfive));
} else {
   PetscCall(VecGetArray(x, &xx));
   xx[0] = 2.0;
   xx[1] = 3.0;
   PetscCall(VecRestoreArray(x, &xx));
}

/*
Note: The user should initialize the vector, x, with the initial guess
for the nonlinear solver prior to calling SNESolve(). In particular,
to employ an initial guess of zero, the user should explicitly set
this vector to zero by calling VecSet().
*/
PetscCall(SNESolve(snes, NULL, x));
if (flg) {
   Vec f;
   PetscCall(VecView(x, PETSC_VIEWER_STDOUT_WORLD));
   PetscCall(SNESGetFunction(snes, &f, 0, 0));
   PetscCall(VecView(r, PETSC_VIEWER_STDOUT_WORLD));
}

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
Free work space. All PETSc objects should be destroyed when they
are no longer needed.
- - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -*/
PetscCall(VecDestroy(&x));
PetscCall(VecDestroy(&r));
PetscCall(MatDestroy(&J));
PetscCall(SNESDestroy(&snes));
PetscCall(PetscFinalize());
return 0;

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -*/

FormFunction1 - Evaluates nonlinear function, F(x).

Input Parameters:
. snes - the SNES context
. x - input vector

(continues on next page)
PetscErrorCode FormFunction1(SNES snes, Vec x, Vec f, void *ctx)
{
    const PetscScalar *xx;
    PetscScalar *ff;

    PetscFunctionBeginUser;
    /*
    Get pointers to vector data.
    - For default PETSc vectors, VecGetArray() returns a pointer to
      the data array. Otherwise, the routine is implementation dependent.
    - You MUST call VecRestoreArray() when you no longer need access to
      the array.
    */
    PetscCall(VecGetArrayRead(x, &xx));
    PetscCall(VecGetArray(f, &ff));
    /* Compute function */
    ff[0] = xx[0] * xx[0] + xx[0] * xx[1] - 3.0;
    /* Restore vectors */
    PetscCall(VecRestoreArrayRead(x, &xx));
    PetscCall(VecRestoreArray(f, &ff));
    PetscFunctionReturn(PETSC_SUCCESS);
}

PetscErrorCode FormJacobian1(SNES snes, Vec x, Mat jac, Mat B, void *dummy)
{
    const PetscScalar *xx;
    PetscScalar A[4];
    PetscInt idx[2] = {0, 1};

    PetscFunctionBeginUser;
    /*
    Get pointer to vector data
    */
    PetscCall(VecGetArrayRead(x, &xx));
    /*
    FormJacobian1 - Evaluates Jacobian matrix.
    
    Input Parameters:
    . snes - the SNES context
    . x - input vector
    . dummy - optional user-defined context (not used here)
    
    Output Parameters:
    . jac - Jacobian matrix
    . B - optionally different preconditioning matrix
    . flag - flag indicating matrix structure
    */
    PetscCall(VecGetArrayRead(x, &xx));
    /* ...
    */

(continues on next page)
Compute Jacobian entries and insert into matrix.
- Since this is such a small problem, we set all entries for the matrix at once.

\[
A[0] = 2.0 \times xx[0] + xx[1];
A[1] = xx[0];
A[3] = xx[0] + 2.0 \times xx[1];
\]
PetscCall(MatSetValuesB, 2, idx, 2, idx, A, INSERT_VALUES));

/*
  Restore vector
*/
PetscCall(VecRestoreArrayRead(x, &xx));

/*
  Assemble matrix
*/
PetscCall(MatAssemblyBeginB, MAT_FINAL_ASSEMBLY);
PetscCall(MatAssemblyEndB, MAT_FINAL_ASSEMBLY);
if (jac != B) { 
PetscCall(MatAssemblyBegin(jac, MAT_FINAL_ASSEMBLY));
PetscCall(MatAssemblyEnd(jac, MAT_FINAL_ASSEMBLY));
}
PetscFunctionReturn(PETSC_SUCCESS);
To create a \texttt{SNES} solver, one must first call \texttt{SNESCreate()} as follows:

\begin{verbatim}
SNESCreate(MPI_Comm comm, SNES *snes);
\end{verbatim}

The user must then set routines for evaluating the residual function ($2.3$) and its associated Jacobian matrix, as discussed in the following sections.

To choose a nonlinear solution method, the user can either call

\begin{verbatim}
SNESSetType(SNES snes, SNESType method);
\end{verbatim}

or use the option \texttt{-snes_type <method>}, where details regarding the available methods are presented in

\section*{2.4. \texttt{SNES}: Nonlinear Solvers}
The Nonlinear Solvers. The application code can take complete control of the linear and nonlinear techniques used in the Newton-like method by calling

```c
SNESSetFromOptions(snes);
```

This routine provides an interface to the PETSc options database, so that at runtime the user can select a particular nonlinear solver, set various parameters and customized routines (e.g., specialized line search variants), prescribe the convergence tolerance, and set monitoring routines. With this routine the user can also control all linear solver options in the KSP, and PC modules, as discussed in KSP: Linear System Solvers.

After having set these routines and options, the user solves the problem by calling

```c
SNESolve(SNES snes, Vec b, Vec x);
```

where \( x \) should be initialized to the initial guess before calling and contains the solution on return. In particular, to employ an initial guess of zero, the user should explicitly set this vector to zero by calling `VecZeroEntries(x)`. Finally, after solving the nonlinear system (or several systems), the user should destroy the SNES context with

```c
SNESDestroy(SNES *snes);
```

Nonlinear Function Evaluation

When solving a system of nonlinear equations, the user must provide a residual function (2.3), which is set using

```c
SNESSetFunction(SNES snes, Vec f, PetscErrorCode (*FormFunction)(SNES snes, Vec x, Vec f, ... void *ctx), void *ctx);
```

The argument \( f \) is an optional vector for storing the solution; pass NULL to have the SNES allocate it for you. The argument \( ctx \) is an optional user-defined context, which can store any private, application-specific data required by the function evaluation routine; NULL should be used if such information is not needed. In C and C++, a user-defined context is merely a structure in which various objects can be stashed; in Fortran a user context can be an integer array that contains both parameters and pointers to PETSc objects. SNES Tutorial ex5 and SNES Tutorial ex5f90 give examples of user-defined application contexts in C and Fortran, respectively.

Jacobian Evaluation

The user must also specify a routine to form some approximation of the Jacobian matrix, \( A \), at the current iterate, \( x \), as is typically done with

```c
SNESSetJacobian(SNES snes, Mat Amat, Mat Pmat, PetscErrorCode (*FormJacobian)(SNES snes, Vec x, Mat A, Mat B, void *ctx), void *ctx);
```

The arguments of the routine `FormJacobian()` are the current iterate, \( x \); the (approximate) Jacobian matrix, \( Amat \); the matrix from which the preconditioner is constructed, \( Pmat \) (which is usually the same as \( Amat \)); and an optional user-defined Jacobian context, \( ctx \), for application-specific data. Note that the SNES solvers are all data-structure neutral, so the full range of PETSc matrix formats (including “matrix-free” methods) can be used. Matrices discusses information regarding available matrix formats and options, while Matrix-Free Methods focuses on matrix-free methods in SNES. We briefly touch on a few details of matrix usage that are particularly important for efficient use of the nonlinear solvers.

A common usage paradigm is to assemble the problem Jacobian in the preconditioner storage \( B \), rather than \( A \). In the case where they are identical, as in many simulations, this makes no difference. However, it allows
us to check the analytic Jacobian we construct in FormJacobian() by passing the -snes_mf_operator flag. This causes PETSc to approximate the Jacobian using finite differencing of the function evaluation (discussed in Finite Difference Jacobian Approximations), and the analytic Jacobian becomes merely the preconditioner. Even if the analytic Jacobian is incorrect, it is likely that the finite difference approximation will converge, and thus this is an excellent method to verify the analytic Jacobian. Moreover, if the analytic Jacobian is incomplete (some terms are missing or approximate), -snes_mf_operator may be used to obtain the exact solution, where the Jacobian approximation has been transferred to the preconditioner.

One such approximate Jacobian comes from “Picard linearization” which writes the nonlinear system as

\[ F(x) := A(x)x - b = 0 \]

where \( A(x) \) usually contains the lower-derivative parts of the equation. For example, the nonlinear diffusion problem

\[-\nabla \cdot (\kappa(u)\nabla u) = 0\]

would be linearized as

\[ A(u)v \approx -\nabla \cdot (\kappa(u)\nabla v). \]

Usually this linearization is simpler to implement than Newton and the linear problems are somewhat easier to solve. In addition to using -snes_mf_operator with this approximation to the Jacobian, the Picard iterative procedure can be performed by defining \( J(x) \) to be \( A(x) \). Sometimes this iteration exhibits better global convergence than Newton linearization.

During successive calls to FormJacobian(), the user can either insert new matrix contexts or reuse old ones, depending on the application requirements. For many sparse matrix formats, reusing the old space (and merely changing the matrix elements) is more efficient; however, if the matrix structure completely changes, creating an entirely new matrix context may be preferable. Upon subsequent calls to the FormJacobian() routine, the user may wish to reinitialize the matrix entries to zero by calling MatZeroEntries(). See Other Matrix Operations for details on the reuse of the matrix context.

The directory $PETSC_DIR/src/snes/tutorials provides a variety of examples.

Sometimes a nonlinear solver may produce a step that is not within the domain of a given function, for example one with a negative pressure. When this occurs one can call SNESSetFunctionDomainError() or SNESSetJacobianDomainError() to indicate to SNES the step is not valid. One must also use SNESGetConvergedReason() and check the reason to confirm if the solver succeeded. See Variational Inequalities for how to provide SNES with bounds on the variables to solve (differential) variational inequalities and how to control properties of the line step computed.

### 2.4.2 The Nonlinear Solvers

As summarized in Table PETSc Nonlinear Solvers, SNES includes several Newton-like nonlinear solvers based on line search techniques and trust region methods. Also provided are several nonlinear Krylov methods, as well as nonlinear methods involving decompositions of the problem.

Each solver may have associated with it a set of options, which can be set with routines and options database commands provided for this purpose. A complete list can be found by consulting the manual pages or by running a program with the -help option; we discuss just a few in the sections below.
Table 2.8: PETSc Nonlinear Solvers

<table>
<thead>
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<th>Options Name</th>
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<td>SNESNGS</td>
<td>ngs</td>
<td>—</td>
</tr>
<tr>
<td>Anderson Mixing</td>
<td>SNESANDERSON</td>
<td>anderson</td>
<td>—</td>
</tr>
<tr>
<td>Newton with constraints (1)</td>
<td>SNESVINEWTONR</td>
<td>vinewtonr</td>
<td>SNESLINESEARCHBT</td>
</tr>
<tr>
<td>Newton with constraints (2)</td>
<td>SNESVINEWTON-</td>
<td>vinewton-</td>
<td>SNESLINESEARCHBT</td>
</tr>
<tr>
<td></td>
<td>SSLS</td>
<td>ssls</td>
<td></td>
</tr>
<tr>
<td>Multi-stage Smoothers</td>
<td>SNESMS</td>
<td>ms</td>
<td>—</td>
</tr>
<tr>
<td>Composite</td>
<td>SNESCOMPOSITE</td>
<td>composite</td>
<td>—</td>
</tr>
<tr>
<td>Linear solve only</td>
<td>SNESKSPONLY</td>
<td>ksponly</td>
<td>—</td>
</tr>
<tr>
<td>Python Shell</td>
<td>SNESPYTHON</td>
<td>python</td>
<td>—</td>
</tr>
<tr>
<td>Shell (user-defined)</td>
<td>SNESHELL</td>
<td>shell</td>
<td>—</td>
</tr>
</tbody>
</table>

Line Search Newton

The method `SNESNEWTONLS (-snes_type newtonls)` provides a line search Newton method for solving systems of nonlinear equations. By default, this technique employs cubic backtracking [DennisJrS83]. Alternative line search techniques are listed in Table **PETSc Line Search Methods**.

Table 2.9: PETSc Line Search Methods

<table>
<thead>
<tr>
<th>Line Search</th>
<th>SNESLineSearchType</th>
<th>Options Name</th>
</tr>
</thead>
<tbody>
<tr>
<td>Backtracking</td>
<td>SNESLINESEARCHBT</td>
<td>bt</td>
</tr>
<tr>
<td>(damped) step</td>
<td>SNESLINESEARCHBASIC</td>
<td>basic</td>
</tr>
<tr>
<td>identical to above</td>
<td>SNESLINESEARCHNONE</td>
<td>none</td>
</tr>
<tr>
<td>L2-norm Minimization</td>
<td>SNESLINESEARCHL2</td>
<td>l2</td>
</tr>
<tr>
<td>Critical point</td>
<td>SNESLINESEARCHCP</td>
<td>cp</td>
</tr>
<tr>
<td>Shell</td>
<td>SNESLINESEARCHSHELL</td>
<td>shell</td>
</tr>
</tbody>
</table>

Every `SNES` has a line search context of type `SNESLineSearch` that may be retrieved using

```c
SNESGetLineSearch(SNES snes,SNESLineSearch *ls);
```

There are several default options for the line searches. The order of polynomial approximation may be set with `-snes_linesearch_order` or

```c
SNESLineSearchSetOrder(SNESLineSearch ls, PetscInt order);
```

for instance, 2 for quadratic or 3 for cubic. Sometimes, it may not be necessary to monitor the progress of the nonlinear iteration. In this case, `-snes_linesearch_norms` or
SNESLineSearchSetComputeNorms(SNESLineSearch ls, PetscBool norms);

may be used to turn off function, step, and solution norm computation at the end of the linesearch.

The default line search for the line search Newton method, SNESLINESEARCHBT involves several parameters, which are set to defaults that are reasonable for many applications. The user can override the defaults by using the following options:

- **-snes_linesearch_alpha <alpha>**
- **-snes_linesearch_maxstep <max>**
- **-snes_linesearch_minlambda <tol>**

Besides the backtracking linesearch, there are SNESLINESEARCHL2, which uses a polynomial secant minimization of \(\|F(x)\|_2\), and SNESLINESEARCHCP, which minimizes \(F(x) \cdot Y\) where \(Y\) is the search direction. These are both potentially iterative line searches, which may be used to find a better-fitted steplength in the case where a single secant search is not sufficient. The number of iterations may be set with **-snes_linesearch_max_it**. In addition, the convergence criteria of the iterative line searches may be set using function tolerances **-snes_linesearch_rtol** and **-snes_linesearch_atol**; and steplength tolerances **snes_linesearch_ltol**.

Custom line search types may either be defined using SNESLineSearchShell, or by creating a custom user line search type in the model of the preexisting ones and register it using

SNESLineSearchRegister(const char sname[], PetscErrorCode (*function)(SNESLineSearch));

### Trust Region Methods

The trust region method in SNES for solving systems of nonlinear equations, SNESNEWTONTR (**-snes_type newtontr**), is taken from the MINPACK project [MoreSGH84]. Several parameters can be set to control the variation of the trust region size during the solution process. In particular, the user can control the initial trust region radius, computed by

\[
\Delta = \Delta_0 \|F_0\|_2,
\]

by setting \(\Delta_0\) via the option **-snes_tr_delta0 <delta0>**.

### Nonlinear Krylov Methods

A number of nonlinear Krylov methods are provided, including Nonlinear Richardson, conjugate gradient, GMRES, and Anderson Mixing. These methods are described individually below. They are all instrumental to PETSc’s nonlinear preconditioning.

**Nonlinear Richardson.** The nonlinear Richardson iteration merely takes the form of a line search-damped fixed-point iteration of the form

\[
x_{k+1} = x_k - \lambda F(x_k), \quad k = 0, 1, \ldots,
\]

where the default linesearch is SNESLINESEARCHL2. This simple solver is mostly useful as a nonlinear smoother, or to provide line search stabilization to an inner method.

**Nonlinear Conjugate Gradients.** Nonlinear CG is equivalent to linear CG, but with the steplength determined by line search (SNESLINESEARCHCP by default). Five variants (Fletcher-Reed, Hestenes-Stiefel, Polak-Ribiere-Polyak, Dai-Yuan, and Conjugate Descent) are implemented in PETSc and may be chosen using
Anderson Mixing and Nonlinear GMRES Methods. Nonlinear GMRES and Anderson Mixing methods combine the last $m$ iterates, plus a new fixed-point iteration iterate, into a residual-minimizing new iterate.

Quasi-Newton Methods

Quasi-Newton methods store iterative rank-one updates to the Jacobian instead of computing it directly. Three limited-memory quasi-Newton methods are provided, L-BFGS, which are described in Table PETSc quasi-Newton solvers. These all are encapsulated under -snes_type qn and may be changed with snes_qn_type. The default is L-BFGS, which provides symmetric updates to an approximate Jacobian. This iteration is similar to the line search Newton methods.

<table>
<thead>
<tr>
<th>QN Method</th>
<th>SNESQNType</th>
<th>Options Name</th>
<th>Default Line Search</th>
</tr>
</thead>
<tbody>
<tr>
<td>L-BFGS</td>
<td>SNES_QN_LBFGS</td>
<td>lbfgs</td>
<td>SNESLINESEARCHCP</td>
</tr>
<tr>
<td>“Good” Broyden</td>
<td>SNES_QN_BROYDEN</td>
<td>broyden</td>
<td>SNESLINESEARCHBASIC</td>
</tr>
<tr>
<td>“Bad” Broyden</td>
<td>SNES_QN_BADBROYDEN</td>
<td></td>
<td>SNESLINESEARCHL2</td>
</tr>
</tbody>
</table>

One may also control the form of the initial Jacobian approximation with

SNESQNSetScaleType(SNES snes, SNESQNScaleType stype);

and the restart type with

SNESQNSetRestartType(SNES snes, SNESQNRestartType rtype);

The Full Approximation Scheme

The Full Approximation Scheme is a nonlinear multigrid correction. At each level, there is a recursive cycle control SNES instance, and either one or two nonlinear solvers as smoothers (up and down). Problems set up using the SNES DMDA interface are automatically coarsened. FAS differs slightly from PCMG, in that the hierarchy is constructed recursively. However, much of the interface is a one-to-one map. We describe the “get” operations here, and it can be assumed that each has a corresponding “set” operation. For instance, the number of levels in the hierarchy may be retrieved using

SNESFASGetLevels(SNES snes, PetscInt *levels);

There are four SNESFAS cycle types, SNES_FAS_MULTIPLICATIVE, SNES_FAS_ADDITIVE, SNES_FAS_FULL, and SNES_FAS_KASKADE. The type may be set with

SNESFASSetType(SNES snes, SNESFASType fastype);

and the cycle type, 1 for V, 2 for W, may be set with

SNESFASSetCycles(SNES snes, PetscInt cycles);.
Much like the interface to \texttt{PCMG} described in \textit{Multigrid Preconditioners}, there are interfaces to recover the various levels' cycles and smoothers. The level smoothers may be accessed with

\begin{verbatim}
SNESFASGetSmoothers(SNES snes, PetscInt level, SNES *smooth);
SNESFASGetSmoothersUp(SNES snes, PetscInt level, SNES *smooth);
SNESFASGetSmoothersDown(SNES snes, PetscInt level, SNES *smooth);
\end{verbatim}

and the level cycles with

\begin{verbatim}
SNESFASGetCycleSNES(SNES snes, PetscInt level, SNES *lsnes);
\end{verbatim}

Also akin to \texttt{PCMG}, the restriction and prolongation at a level may be acquired with

\begin{verbatim}
SNESFASGetInterpolation(SNES snes, PetscInt level, Mat *mat);
SNESFASGetRestriction(SNES snes, PetscInt level, Mat *mat);
\end{verbatim}

In addition, FAS requires special restriction for solution-like variables, called injection. This may be set with

\begin{verbatim}
SNESFASGetInjection(SNES snes, PetscInt level, Mat *mat);
\end{verbatim}

The coarse solve context may be acquired with

\begin{verbatim}
SNESFASGetCoarseSolve(SNES snes, SNES *smooth);
\end{verbatim}

\textbf{Nonlinear Additive Schwarz}

Nonlinear Additive Schwarz methods (NASM) take a number of local nonlinear subproblems, solves them independently in parallel, and combines those solutions into a new approximate solution.

\begin{verbatim}
SNESNASMSetSubdomains(SNES snes, PetscInt n, SNES subsnes[], VecScatter isscatter[],...
VecScatter oscatter[], VecScatter gscatter[]);
\end{verbatim}

allows for the user to create these local subdomains. Problems set up using the \texttt{SNES DMDA} interface are automatically decomposed. To begin, the type of subdomain updates to the whole solution are limited to two types borrowed from \texttt{PCASM: PC_ASM_BASIC}, in which the overlapping updates added. \texttt{PC_ASM_RESTRICT} updates in a nonoverlapping fashion. This may be set with

\begin{verbatim}
SNESNASMSetType(SNES snes, PCASMType type);
\end{verbatim}

\texttt{SNESASPIN} is a helper \texttt{SNES} type that sets up a nonlinearly preconditioned Newton’s method using NASM as the preconditioner.

\subsection*{2.4.3 General Options}

This section discusses options and routines that apply to all \texttt{SNES} solvers and problem classes. In particular, we focus on convergence tests, monitoring routines, and tools for checking derivative computations.
Convergence Tests

Convergence of the nonlinear solvers can be detected in a variety of ways; the user can even specify a customized test, as discussed below. Most of the nonlinear solvers use `SNESConvergenceTestDefault()`, however, `SNESNEWTONTR` uses a method-specific additional convergence test as well. The convergence tests involves several parameters, which are set by default to values that should be reasonable for a wide range of problems. The user can customize the parameters to the problem at hand by using some of the following routines and options.

One method of convergence testing is to declare convergence when the norm of the change in the solution between successive iterations is less than some tolerance, `stol`. Convergence can also be determined based on the norm of the function. Such a test can use either the absolute size of the norm, `atol`, or its relative decrease, `rtol`, from an initial guess. The following routine sets these parameters, which are used in many of the default `SNES` convergence tests:

```c
SNESSetTolerances(SNES snes, PetscReal atol, PetscReal rtol, PetscReal stol, PetscInt its, PetscInt fcts);
```

This routine also sets the maximum numbers of allowable nonlinear iterations, `its`, and function evaluations, `fcts`. The corresponding options database commands for setting these parameters are:

- `-snes_atol <atol>`
- `-snes_rtol <rtol>`
- `-snes_stol <stol>`
- `-snes_max_it <its>`
- `-snes_max_funcs <fcts>`

A related routine is `SNESGetTolerances()`.

Convergence tests for trust regions methods often use an additional parameter that indicates the minimum allowable trust region radius. The user can set this parameter with the option `-snes_tr_tol <trtol>` or with the routine

```c
SNESSetTrustRegionTolerance(SNES snes, PetscReal trtol);
```

Users can set their own customized convergence tests in `SNES` by using the command

```c
SNESSetConvergenceTest(SNES snes, PetscErrorCode (*test)(SNES snes, PetscInt it, PetscReal xnorm, PetscReal gnorm, PetscReal f, SNESConvergedReason reason, void *cctx), void *cctx, PetscErrorCode (*destroy)(void *cctx));
```

The final argument of the convergence test routine, `cctx`, denotes an optional user-defined context for private data. When solving systems of nonlinear equations, the arguments `xnorm`, `gnorm`, and `f` are the current iterate norm, current step norm, and function norm, respectively. `SNESConvergedReason` should be set positive for convergence and negative for divergence. See `include/petscsnes.h` for a list of values for `SNESConvergedReason`. 


Convergence Monitoring

By default the SNES solvers run silently without displaying information about the iterations. The user can initiate monitoring with the command

\[
\text{SNESMonitorSet}(\text{SNES } \text{snes}, \text{PetscErrorCode} \ (\text{*mon})(\text{SNES}, \text{PetscInt } \text{its}, \text{PetscReal } \text{norm}, \text{void}*, \text{PetscErrorCode} \ (*\text{monitordestroy})(\text{void}**));)
\]

The routine, mon, indicates a user-defined monitoring routine, where its and mctx respectively denote the iteration number and an optional user-defined context for private data for the monitor routine. The argument norm is the function norm.

The routine set by SNESMonitorSet() is called once after every successful step computation within the nonlinear solver. Hence, the user can employ this routine for any application-specific computations that should be done after the solution update. The option -snes_monitor activates the default SNES monitor routine, SNESMonitorDefault(), while -snes_monitor_lg_residualnorm draws a simple line graph of the residual norm's convergence.

One can cancel hardwired monitoring routines for SNES at runtime with -snes_monitor_cancel.

As the Newton method converges so that the residual norm is small, say $10^{-10}$, many of the final digits printed with the -snes_monitor option are meaningless. Worse, they are different on different machines; due to different round-off rules used by, say, the IBM RS6000 and the Sun SPARC. This makes testing between different machines difficult. The option -snes_monitor_short causes PETSc to print fewer of the digits of the residual norm as it gets smaller; thus on most of the machines it will always print the same numbers making cross-process testing easier.

The routines

\[
\text{SNESGetSolution}(\text{SNES } \text{snes}, \text{Vec } \text{*x});
\]
\[
\text{SNESGetFunction}(\text{SNES } \text{snes}, \text{Vec } \text{*r}, \text{void } \text{*ctx}, \text{int} (**\text{func})(\text{SNES, Vec, Vec, void}*));
\]

return the solution vector and function vector from a SNES context. These routines are useful, for instance, if the convergence test requires some property of the solution or function other than those passed with routine arguments.

Checking Accuracy of Derivatives

Since hand-coding routines for Jacobian matrix evaluation can be error prone, SNES provides easy-to-use support for checking these matrices against finite difference versions. In the simplest form of comparison, users can employ the option -snes_test_jacobian to compare the matrices at several points. Although not exhaustive, this test will generally catch obvious problems. One can compare the elements of the two matrices by using the option -snes_test_jacobian_view, which causes the two matrices to be printed to the screen.

Another means for verifying the correctness of a code for Jacobian computation is running the problem with either the finite difference or matrix-free variant. -snes_fd or -snes_mf; see Finite Difference Jacobian Approximations or Matrix-Free Methods. If a problem converges well with these matrix approximations but not with a user-provided routine, the problem probably lies with the hand-coded matrix. See the note in Jacobian Evaluation about assembling your Jacobian in the “preconditioner” slot Pmat.

The correctness of user provided MATSHELL Jacobians in general can be checked with MatShellTestMultTranspose() and MatShellTestMult().

The correctness of user provided MATSHELL Jacobians via TSSetRHSJacobian() can be checked with TSRSJacobianTestTranspose() and TSRSJacobianTest() that check the correction of the matrix-transpose vector product and the matrix-product. From the command line, these can be checked with
2.4.4 Inexact Newton-like Methods

Since exact solution of the linear Newton systems within (2.4) at each iteration can be costly, modifications are often introduced that significantly reduce these expenses and yet retain the rapid convergence of Newton’s method. Inexact or truncated Newton techniques approximately solve the linear systems using an iterative scheme. In comparison with using direct methods for solving the Newton systems, iterative methods have the virtue of requiring little space for matrix storage and potentially saving significant computational work. Within the class of inexact Newton methods, of particular interest are Newton-Krylov methods, where the subsidiary iterative technique for solving the Newton system is chosen from the class of Krylov subspace projection methods. Note that at runtime the user can set any of the linear solver options discussed in KSP: Linear System Solvers, such as -ksp_type <ksp_method> and -pc_type <pc_method>, to set the Krylov subspace and preconditioner methods.

Two levels of iterations occur for the inexact techniques, where during each global or outer Newton iteration a sequence of subsidiary inner iterations of a linear solver is performed. Appropriate control of the accuracy to which the subsidiary iterative method solves the Newton system at each global iteration is critical, since these inner iterations determine the asymptotic convergence rate for inexact Newton techniques. While the Newton systems must be solved well enough to retain fast local convergence of the Newton’s iterates, use of excessive inner iterations, particularly when \( \|x_k - x^*\| \) is large, is neither necessary nor economical. Thus, the number of required inner iterations typically increases as the Newton process progresses, so that the truncated iterates approach the true Newton iterates.

A sequence of nonnegative numbers \( \{\eta_k\} \) can be used to indicate the variable convergence criterion. In this case, when solving a system of nonlinear equations, the update step of the Newton process remains unchanged, and direct solution of the linear system is replaced by iteration on the system until the residuals

\[
r_k^{(i)} = F'(x^*_k) \Delta x_k + F(x_k)
\]

satisfy

\[
\frac{\|r_k^{(i)}\|}{\|F(x_k)\|} \leq \eta_k \leq \eta < 1.
\]

Here \( x_0 \) is an initial approximation of the solution, and \( \| \cdot \| \) denotes an arbitrary norm in \( \mathbb{R}^n \).

By default a constant relative convergence tolerance is used for solving the subsidiary linear systems within the Newton-like methods of SNES. When solving a system of nonlinear equations, one can instead employ the techniques of Eisenstat and Walker [EW96] to compute \( \eta_k \) at each step of the nonlinear solver by using the option -snes_ksp_ew. In addition, by adding one’s own KSP convergence test (see Convergence Tests), one can easily create one’s own, problem-dependent, inner convergence tests.
2.4.5 Matrix-Free Methods

The SNES class fully supports matrix-free methods. The matrices specified in the Jacobian evaluation routine need not be conventional matrices; instead, they can point to the data required to implement a particular matrix-free method. The matrix-free variant is allowed only when the linear systems are solved by an iterative method in combination with no preconditioning (PCNONE or -pc_type none), a user-provided preconditioner matrix, or a user-provided preconditioner shell (PCSHELL, discussed in Preconditioners); that is, obviously matrix-free methods cannot be used with a direct solver, approximate factorization, or other preconditioner which requires access to explicit matrix entries.

The user can create a matrix-free context for use within SNES with the routine

```c
MatCreateSNESMF(SNES snes, Mat *mat);
```

This routine creates the data structures needed for the matrix-vector products that arise within Krylov space iterative methods [BS90]. The default SNES matrix-free approximations can also be invoked with the command -snes_mf. Or, one can retain the user-provided Jacobian preconditioner, but replace the user-provided Jacobian matrix with the default matrix-free variant with the option -snes_mf_operator.

The user can set one parameter to control the Jacobian-vector product approximation with the command

```c
MatMFFDSetFunctionError(Mat mat, PetscReal rerror);
```

The parameter `rerror` should be set to the square root of the relative error in the function evaluations, $e_{rel}$; the default is the square root of machine epsilon (about $10^{-8}$ in double precision), which assumes that the functions are evaluated to full floating-point precision accuracy. This parameter can also be set from the options database with `-mat_mffd_err <err>`.

In addition, PETSc provides ways to register new routines to compute the differencing parameter ($h$); see the manual page for `MatMFFDSetType()` and `MatMFFDRegister()`. We currently provide two default routines accessible via `-mat_mffd_type <ds or wp>`. For the default approach there is one “tuning” parameter, set with

```c
MatMFFDDSSetUmin(Mat mat, PetscReal umin);
```

This parameter, $u_{\text{min}}$ (or $u_{\text{min}}$), is a bit involved; its default is $10^{-6}$ . Its command line form is `-mat_mffd_umin <umin>`.

The Jacobian-vector product is approximated via the formula

$$F'(u)a \approx \frac{F(u + h * a) - F(u)}{h}$$

where $h$ is computed via

$$h = e_{rel} \cdot \begin{cases} u^T a / \| a \|^2_2 & \text{if } |u^T a| > u_{\text{min}} \| a \|_1 \\ u_{\text{min}} \text{sign}(u^T a) \| a \|_1 / \| a \|^2_2 & \text{otherwise.} \end{cases}$$

This approach is taken from Brown and Saad [BS90]. The second approach, taken from Walker and Pernice, [PW98], computes $h$ via

$$h = \frac{\sqrt{1 + \|u\|_2^2 e_{rel}}}{\| a \|}$$
This has no tunable parameters, but note that inside the nonlinear solve for the entire linear iterative process $u$ does not change hence $\sqrt{1+||u||}$ need be computed only once. This information may be set with the options

\begin{verbatim}
MatMFFDWPSetComputeNormU(Mat mat, PetscBool);
\end{verbatim}

or -mat_mffd_compute_normu <true or false>. This information is used to eliminate the redundant computation of these parameters, therefore reducing the number of collective operations and improving the efficiency of the application code. This takes place automatically for the PETSc GMRES solver with left preconditioning.

It is also possible to monitor the differing parameters $h$ that are computed via the routines

\begin{verbatim}
MatMFFDSetHHistory(Mat mat, PetscScalar *, int);
MatMFFDResetHHistory(Mat mat, PetscScalar *, int);
MatMFFDGetH(Mat mat, PetscScalar *);
\end{verbatim}

We include an explicit example of using matrix-free methods in `ex3.c`. Note that by using the option -snes_mf one can easily convert any SNES code to use a matrix-free Newton-Krylov method without a preconditioner. As shown in this example, SNESSetFromOptions() must be called after SNESSetJacobian() to enable runtime switching between the user-specified Jacobian and the default SNES matrix-free form.

Listing: src/snes/tutorials/ex3.c

```
static char help[] = "Newton methods to solve $u'' + u^2 = f$ in parallel.\n";
This example employs a user-defined monitoring routine and optionally a user-defined\nroutine to check candidate iterates produced by line search routines.\nThe command line options include:\n-\npre_check_iterates : activate checking of iterates\n-post_check_iterates : activate checking of iterates\n-check_tol <tol>: set tolerance for iterate checking\n-user_precond : activate a (trivial) user-defined preconditioner\n"

#include "petscdm.h" so that we can use data management objects (DMs)\n#include "petscdmda.h" so that we can use distributed arrays (DMDAs).\n#include "petscsnes.h" so that we can use SNES solvers. Note that this\nfile automatically includes:\n  petscsys.h - base PETSc routines\npetscvec.h - vectors\npetscmat.h - matrices\npetscis.h - index sets\npetscksp.h - Krylov subspace methods\npetscviewer.h - viewers\npetscpc.h - preconditioners\npetscksp.h - linear solvers

/*

User-defined routines.
*/
```

(continues on next page)
PetscErrorCode FormJacobian(SNES, Vec, Mat, Mat, void *);
PetscErrorCode FormFunction(SNES, Vec, Vec, void *);
PetscErrorCode FormInitialGuess(Vec);
PetscErrorCode Monitor(SNES, PetscInt, PetscReal, void *);
PetscErrorCode PreCheck(SNESLineSearch, Vec, Vec, PetscBool *, void *);
PetscErrorCode PostCheck(SNESLineSearch, Vec, Vec, Vec, PetscBool *, PetscBool *, void *);
PetscErrorCode PostSetSubKSP(SNESLineSearch, Vec, Vec, Vec, PetscBool *, void *);
PetscErrorCode MatrixFreePreconditioner(PC, Vec, Vec);

/* User-defined application context */
typedef struct {
    DM da; /* distributed array */
    Vec F; /* right-hand-side of PDE */
    PetscMPIInt rank; /* rank of processor */
    PetscMPIInt size; /* size of communicator */
    PetscReal h; /* mesh spacing */
    PetscBool sjerr; /* if or not to test jacobian domain error */
} ApplicationCtx;

/* User-defined context for monitoring */
typedef struct {
    PetscViewer viewer;
} MonitorCtx;

/* User-defined context for checking candidate iterates that are determined by line search methods */
typedef struct {
    Vec last_step; /* previous iterate */
    PetscReal tolerance; /* tolerance for changes between successive iterates */
    ApplicationCtx *user;
} StepCheckCtx;

typedef struct {
    PetscInt its0; /* num of previous outer KSP iterations */
} SetSubKSPCtx;

int main(int argc, char **argv) {
    SNES snes; /* SNES context */
    SNESLineSearch linesearch; /* SNESLineSearch context */
    Mat J; /* Jacobian matrix */
    ApplicationCtx ctx; /* user-defined context */
    Vec x, r, U, F; /* vectors */
    MonitorCtx monP; /* monitoring context */
    StepCheckCtx checkP; /* step-checking context */
    SetSubKSPCtx checkP1;
    PetscBool pre_check, post_check, post_setsubksp; /* flag indicating whether we're checking candidate iterates */
PetscScalar xp, *FF, *UU, none = -1.0;
PetscInt its, N = 5, i, maxit, maxf, xs, xm;
PetscReal abstol, rtol, stol, norm;
PetscBool flg, viewinitial = PETSC_FALSE;

PetsfuncBeginUser;
PetscCall(PetscInitialize(&argc, &argv, (char *)0, help));
PetscCallMPI(MPI_Comm_rank(PETSC_COMM_WORLD, &ctx.rank));
PetscCallMPI(MPI_Comm_size(PETSC_COMM_WORLD, &ctx.size));
PetscCall(PetscOptionsGetInt(NULL, NULL, "-n", &N, NULL));
ctx.h = 1.0 / (N - 1);
ctx.sjerr = PETSC_FALSE;
PetscCall(PetscOptionsGetBool(NULL, NULL, "-test_jacobian_domain_error", &ctx.sjerr, NULL));
PetscCall(PetscOptionsGetBool(NULL, NULL, "-view_initial", &viewinitial, NULL));

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
Create nonlinear solver context
- - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - */
PetscCall(SNESCreate(PETSC_COMM_WORLD, &snes));

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
Create vector data structures; set function evaluation routine
- - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - */

/* Create distributed array (DMDA) to manage parallel grid and vectors */
PetscCall(DMDACreate1d(PETSC_COMM_WORLD, DM_BOUNDARY_NONE, N, 1, 1, NULL, &ctx.da));
PetscCall(DMSetFromOptions(ctx.da));
PetscCall(DMSetUp(ctx.da));

/* Extract global and local vectors from DMDA; then duplicate for remaining vectors that are the same types */
PetscCall(DMCreateGlobalVector(ctx.da, &x));
PetscCall(VecDuplicate(x, &r));
PetscCall(VecDuplicate(x, &F));
ctx.F = F;
PetscCall(VecDuplicate(x, &U));

/* Set function evaluation routine and vector. Whenever the nonlinear solver needs to compute the nonlinear function, it will call this routine. - Note that the final routine argument is the user-defined context that provides application-specific data for the function evaluation routine. */
PetscCall(SNNESSetFunction(snes, r, FormFunction, &ctx));

/* - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
Create matrix data structure; set Jacobian evaluation routine
- - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - */

(continues on next page)
PetscCall(MatCreate(PETSC_COMM_WORLD, &J));
PetscCall(MatSetSizes(J, PETSC_DECIDE, PETSC_DECIDE, N, N));
PetscCall(MatSetFromOptions(J));
PetscCall(MatSeqAIJSetPreallocation(J, 3, NULL));
PetscCall(MatMPIAIJSetPreallocation(J, 3, NULL, 3, NULL));

/*
 Set Jacobian matrix data structure and default Jacobian evaluation
 routine. Whenever the nonlinear solver needs to compute the
 Jacobian matrix, it will call this routine.
 - Note that the final routine argument is the user-defined
   context that provides application-specific data for the
   Jacobian evaluation routine.
 */
PetscCall(SNESSetJacobian(snes, J, J, FormJacobian, &ctx));

/* Optionally allow user-provided preconditioner */
PetscCall(PetscOptionsHasName(NULL, NULL, "-user_precond", &flg));
if (flg) {
  KSP ksp;
  PC pc;
  PetscCall(SNESGetKSP(snes, &ksp));
  PetscCall(KSPGetPC(ksp, &pc));
  PetscCall(PCSetType(pc, PCSHELL));
  PetscCall(PCShellSetApply(pc, MatrixFreePreconditioner));
}

/* Customize nonlinear solver; set runtime options */

/* Set an optional user-defined monitoring routine */
PetscCall(PetscViewerDrawOpen(PETSC_COMM_WORLD, 0, 0, 0, 0, 400, 400, &monP.
  viewer));
PetscCall(SNESMonitorSet(snes, Monitor, 0, &monP, 0));

/* Set names for some vectors to facilitate monitoring (optional) */
PetscCall(PetscObjectSetName((PetscObject)x, "Approximate Solution"));
PetscCall(PetscObjectSetName((PetscObject)U, "Exact Solution"));

/* Set SNES/KSP/KSP/PC runtime options, e.g.,
   -snes_view -snes_monitor -ksp_type <ksp> -pc_type <pc>
 */
PetscCall(SNESSetFromOptions(snes));

/* Set an optional user-defined routine to check the validity of candidate
 iterates that are determined by line search methods */

(continues on next page)
PetscCall(SNESGetLineSearch(snes, &linesearch));
PetscCall(PetscOptionsHasName(NULL, NULL, "-post_check_iterates", &post_check));

if (post_check) {
PetscCall(PetscPrintf(PETSC_COMM_WORLD, "Activating post step checking routine\n"));
PetscCall(SNESLineSearchSetPostCheck(linesearch, PostCheck, &checkP));
PetscCall(VecDuplicate(x, &(checkP.last_step)));

checkP.tolerance = 1.0;
checkP.user = &ctx;

PetscCall(PetscOptionsGetReal(NULL, NULL, "-check_tol", &checkP.tolerance, NULL));
}

PetscCall(PetscOptionsHasName(NULL, NULL, "-post_setsubksp", &post_setsubksp));
if (post_setsubksp) {
PetscCall(PetscPrintf(PETSC_COMM_WORLD, "Activating post setsubksp\n"));
PetscCall(SNESLineSearchSetPostCheck(linesearch, PostSetSubKSP, &checkP1));
}

PetscCall(PetscOptionsHasName(NULL, NULL, "-pre_check_iterates", &pre_check));
if (pre_check) {
PetscCall(PetscPrintf(PETSC_COMM_WORLD, "Activating pre step checking routine\n"));
PetscCall(SNESLineSearchSetPreCheck(linesearch, PreCheck, &checkP));
}

/* Print parameters used for convergence testing (optional) ... just to demonstrate this routine; this information is also printed with the option -snes_view */
PetscCall(SNESGetTolerances(snes, &abstol, &rtol, &stol, &maxit, &maxf));
PetscCall(PetscPrintf(PETSC_COMM_WORLD, "atol=%g, rtol=%g, stol=%g, maxit=%d, maxf=%d\n", &abstol, &rtol, &stol, &maxit, &maxf));

/* Initialize application: Store right-hand-side of PDE and exact solution */
PetscCall(DMDAGetCorners(ctx.da, &xs, NULL, NULL, &xm, NULL, NULL));
PetscCall(DMDAVecGetArray(ctx.da, F, &FF));
PetscCall(DMDAVecGetArray(ctx.da, U, &UU));

(continues on next page)
Compute local vector entries

```c
xp = ctx.h * xs;
for (i = xs; i < xs + xm; i++) {
    FF[i] = 6.0 * xp + PetscPowScalar(xp + 1.e-12, 6.0); /* +1.e-12 is to prevent 0^6, */
    UU[i] = xp * xp * xp;
    xp += ctx.h;
}
```

Restore vectors

```c
PetscCall(DMDAVecRestoreArray(ctx.da, F, &FF));
PetscCall(DMDAVecRestoreArray(ctx.da, U, &UU));
if (viewinitial) {
    PetscCall(VecView(U, 0));
    PetscCall(VecView(F, 0));
}
```

Evaluate initial guess; then solve nonlinear system

```c
PetscCall(FormInitialGuess(x));
PetscCall(SNESSolve(snes, NULL, x));
PetscCall(SNESGetIterationNumber(snes, &its));
PetcCall(PetscPrintf(PETSC_COMM_WORLD, "Number of SNES iterations = %d \n", its));
```

Check solution and clean up

```c
PetscCall(VecAXPY(x, none, U));
PetscCall(VecNorm(x, NORM_2, &norm));
PetscCall(PetscPrintf(PETSC_COMM_WORLD, "Norm of error %g Iterations %d \n", norm, its));
if (ctx.sjerr) {
    SNESType snestype;
    PetscCall(SNESGetType(snes, &snestype));
    PetscCall(PetscPrintf(PETSC_COMM_WORLD, "SNES Type %s\n", snestype));
}
```

Free work space. All PETSc objects should be destroyed when they
are no longer needed.
/*
PetscCall(PetscViewerDestroy(&monP.viewer));
if (post_check) PetscCall(VecDestroy(&checkP.last_step));
PetscCall(VecDestroy(&x));
PetscCall(VecDestroy(&r));
PetscCall(VecDestroy(&U));
PetscCall(VecDestroy(&F));
PetscCall(MatDestroy(&J));
PetscCall(SNESDestroy(&snes));
PetscCall(DMDestroy(&ctx.da));
PetscCall(PetscFinalize());
return 0;
}
/* --------------------------------------------------------------- */
/*
FormInitialGuess - Computes initial guess.
Input/Output Parameter:
  x - the solution vector
*/
PetscErrorCode FormInitialGuess(Vec x)
{
  PetscScalar pfive = .50;
  PetscFunctionBeginUser;
  PetscCall(VecSet(x, pfive));
  PetscFunctionReturn(PETSC_SUCCESS);
}
/* --------------------------------------------------------------- */
/*
FormFunction - Evaluates nonlinear function, F(x).
Input Parameters:
  snes - the SNES context
  x - input vector
  ctx - optional user-defined context, as set by SNESetFunction()
Output Parameter:
  f - function vector
Note:
The user-defined context can contain any application-specific
data needed for the function evaluation.
*/
PetscErrorCode FormFunction(SNES snes, Vec x, Vec f, void *ctx)
{
  ApplicationCtx  *user = (ApplicationCtx *)ctx;
  DM               da = user->da;
  PetscScalar      *ff, d;
  const PetscScalar *xx, *FF;
  PetscInt          i, M, xs, xm;
  Vec               xlocal;

PetscFunctionBeginUser;
PetscCall(DMGetLocalVector(da, &xlocal));
/*
  Scatter ghost points to local vector, using the 2-step process
  DMGlobalToLocalBegin(), DMGlobalToLocalEnd().
  By placing code between these two statements, computations can
  be done while messages are in transition.
*/
PetscCall(DMGlobalToLocalBegin(da, x, INSERT_VALUES, xlocal));
PetscCall(DMGlobalToLocalEnd(da, x, INSERT_VALUES, xlocal));
/*
  Get pointers to vector data.
  - The vector xlocal includes ghost point; the vectors x and f do
    NOT include ghost points.
  - Using DMDAVecGetArray() allows accessing the values using global ordering
*/
PetscCall(DMDAVecGetArrayRead(da, xlocal, (void *)&xx));
PetscCall(DMDAVecGetArray(da, f, &ff));
PetscCall(DMDAVecGetArrayRead(da, user->F, (void *)&FF));
/*
  Get local grid boundaries (for 1-dimensional DMDA):
  xs, xm - starting grid index, width of local grid (no ghost points)
*/
PetscCall(DMDAGetCorners(da, &xs, NULL, NULL, &xm, NULL, NULL));
PetscCall(DMDAGetInfo(da, NULL, &M, NULL, NULL, NULL, NULL, NULL, NULL, NULL, NULL));
/*
  Set function values for boundary points; define local interior grid point range:
  xsi - starting interior grid index
  xei - ending interior grid index
*/
if (xs == 0) { /* left boundary */
  ff[0] = xx[0];
  xs++;
  xm--;
}
if (xs + xm == M) { /* right boundary */
  ff[xs + xm - 1] = xx[xs + xm - 1] - 1.0;
  xm--;
}
/*
  Compute function over locally owned part of the grid (interior points only)
*/
d = 1.0 / (user->h * user->h);
for (i = xs; i < xs + xm; i++) ff[i] = d * (xx[i - 1] - 2.0 * xx[i] + xx[i + 1]) +...
x[i] * xx[i] - FF[i];
/*
  Restore vectors
*/
PetscCall(DMDAVecRestoreArrayRead(da, xlocal, (void *)&xx));
PetscCall(DMDAVecRestoreArray(da, f, &ff));
(continues on next page)
PetscCall(DMDAVecRestoreArrayRead(da, user->F, (void *)&F));
PetscCall(DMRestoreLocalVector(da, &xlocal));
PetscFunctionReturn(PETSC_SUCCESS);
}

/* FormJacobian - Evaluates Jacobian matrix. */

PetscFunctionCode FormJacobian(SNES snes, Vec x, Mat jac, Mat B, void *ctx)
{
    ApplicationCtx *user = (ApplicationCtx *)ctx;
    PetscScalar *xx, d, A[3];
    PetscInt i, j[3], M, xs, xm;
    DM da = user->da;

    PetscFunctionBeginUser;
    if (user->sjerr) {
        PetscCall(SNESSetJacobianDomainError(snes));
        PetscFunctionReturn(PETSC_SUCCESS);
    }

    if (xs == 0) { /* left boundary */
        i = 0;
        A[0] = 1.0;

        PetscCall(MatSetValues(jac, 1, &i, 1, &i, A, INSERT_VALUES));
        xs++;
        xm--;
    }
}
if (xs + xm == M) { /* right boundary */
    i = M - 1;
    A[0] = 1.0;
    PetscCall(MatSetValues(jac, 1, &i, 1, &i, A, INSERT_VALUES));
    xm -=;
}

/*
 * Interior grid points
 * - Note that in this case we set all elements for a particular row at once.
 */
    d = 1.0 / (user->h * user->h);
for (i = xs; i < xs + xm; i++) {
    j[0] = i - 1;
    j[1] = i;
    j[2] = i + 1;
    A[1] = -2.0 * d + 2.0 * xx[i];
    PetscCall(MatSetValues(jac, 1, &i, 3, j, A, INSERT_VALUES));
}

/*
Assemble matrix, using the 2-step process:
  MatAssemblyBegin(), MatAssemblyEnd().
By placing code between these two statements, computations can be done while messages are in transition.

Also, restore vector.
*/
PetscCall(MatAssemblyBegin(jac, MAT_FINAL_ASSEMBLY));
PetscCall(DMDAVecRestoreArrayRead(da, x, &xx));
PetscCall(MatAssemblyEnd(jac, MAT_FINAL_ASSEMBLY));
PetscFunctionReturn(PETSC_SUCCESS);
}

/* --------------------------------------------------------------- */

Monitor - Optional user-defined monitoring routine that views the current iterate with an x-window plot. Set by SNESMonitorSet().

Input Parameters:
  snes - the SNES context
  its - iteration number
  norm - 2-norm function value (may be estimated)
  ctx - optional user-defined context for private data for the monitor routine, as set by SNESMonitorSet()

Note:
  See the manpage for PetscViewerDrawOpen() for useful runtime options, such as -nox to deactivate all x-window output.
*/
PetscErrorCode Monitor(SNES snes, PetscInt its, PetscReal fnorm, void *ctx) {

(continues on next page)
MonitorCtx *monP = (MonitorCtx *)ctx;
Vec x;

PetscFunctionBeginUser;
PetscCall(PetscPrintf(PETSC_COMM_WORLD, "iter = %" PetscInt_FMT ", SNES Function
-norm %g\n", its, (double)fnorm));
PetscCall(SNESGetSolution(snes, &x));
PetcCall(VecView(x, monP->viewer));
PetcFunctionReturn(PETSC_SUCCESS);
}

/* ----------------------------- */

PreCheck - Optional user-defined routine that checks the validity of
candidate steps of a line search method. Set by SNESLineSearchSetPreCheck().

Input Parameters:
snes - the SNES context
xcurrent - current solution
y - search direction and length

Output Parameters:
y - proposed step (search direction and length) (possibly changed)
changed_y - tells if the step has changed or not

PetscErrorCode PreCheck(SNESLineSearch linesearch, Vec xcurrent, Vec y, PetscBool *changed_y, void *ctx)
{
  PetscFunctionBeginUser;
  *changed_y = PETSC_FALSE;
  PetscFunctionReturn(PETSC_SUCCESS);
}

/* ----------------------------- */

PostCheck - Optional user-defined routine that checks the validity of
candidate steps of a line search method. Set by SNESLineSearchSetPostCheck().

Input Parameters:
snes - the SNES context
cx - optional user-defined context for private data for the
  monitor routine, as set by SNESLineSearchSetPostCheck()
xcurrent - current solution
y - search direction and length
x - the new candidate iterate

Output Parameters:
y - proposed step (search direction and length) (possibly changed)
x - current iterate (possibly modified)

PetscErrorCode PostCheck(SNESLineSearch linesearch, Vec xcurrent, Vec y, Vec x, 
PetcBool *changed_y, PetscBool *changed_x, void *ctx)
{
  PetscInt i, iter, xs, xm;
  StepCheckCtx *check;
ApplicationCtx *user;
PetscScalar *xa, *xa_last, tmp;
PetscReal rdiff;
DM da;
SNES snes;

PetscFunctionBeginUser;
*changed_x = PETSC_FALSE;
*changed_y = PETSC_FALSE;
PetscCall(SNESLineSearchGetSNES(linesearch, &snes));
check = (StepCheckCtx *)ctx;
user = check->user;
PetscCall(SNESGetIterationNumber(snes, &iter));

/* iteration 1 indicates we are working on the second iteration */
if (iter > 0) {
da = user->da;
PetscCall(PetscPrintf(PETSC_COMM_WORLD, "Checking candidate step at iteration %", iter, " with tolerance %g
", check->tolerance));

/* Access local array data */
PetscCall(DMDAVecGetArray(da, check->last_step, &xa_last));
PetscCall(DMDAVecGetArray(da, x, &xa));
PetscCall(DMDAGetCorners(da, &xs, NULL, NULL, &xm, NULL, NULL));

/* If we fail the user-defined check for validity of the candidate iterate,
then modify the iterate as we like. (Note that the particular modification
below is intended simply to demonstrate how to manipulate this data, not
as a meaningful or appropriate choice.) */
for (i = xs; i < xs + xm; i++) {
    if (!PetscAbsScalar(xa[i])) rdiff = 2 * check->tolerance;
    else rdiff = PetscAbsScalar((xa[i] - xa_last[i]) / xa[i]);
    if (rdiff > check->tolerance) {
        tmp = xa[i];
        xa[i] = .5 * (xa[i] + xa_last[i]);
        *changed_x = PETSC_TRUE;
PetscCall(PetscPrintf(PETSC_COMM_WORLD, " Altering entry %g, x_last=%g, diff=%g, x_new=%g\n", i, (double)PetscAbsScalar(tmp), (double)PetscAbsScalar(xa_last[i]), (double)rdiff, (double)PetscAbsScalar(xa[i])));
    }
PetscCall(DMDAVecRestoreArray(da, check->last_step, &xa_last));
PetscCall(DMDAVecRestoreArray(da, x, &xa));
}
PetscCall(VecCopy(x, check->last_step));
PetscFunctionReturn(PETSC_SUCCESS);
}

/* *------------------------------------------------------------------*/

PostSetSubKSP - Optional user-defined routine that reset SubKSP options when,
hierarchical bjacobi PC is used

(continues on next page)
Input Parameters:
linesearch - the LineSearch context
xcurrent - current solution
y - search direction and length
x - the new candidate iterate

Output Parameters:
y - proposed step (search direction and length) (possibly changed)
x - current iterate (possibly modified)

PetscErrorCode PostSetSubKSP(SNESLineSearch linesearch, Vec xcurrent, Vec y, Vec x, _PetscBool *changed_y, PetscBool *changed_x, void *ctx)
{
  SetSubKSPCtx *check;
PetscInt iter, its, sub_its, maxit;
  KSP ksp, sub_ksp, *sub_ksps;
  PC pc;
PetscReal ksp_ratio;
  SNES snes;
  PetscFunctionBeginUser;
  PetscCall(SNESLineSearchGetSNES(linesearch, &snes));
  check = (SetSubKSPCtx *)ctx;
  PetscCall(SNESGetIterationNumber(snes, &iter));
  PetscCall(SNESGetKSP(snes, &ksp));
  PetscCall(KSPGetPC(ksp, &pc));
  PetscCall(PCBJacobiGetSubKSP(pc, NULL, NULL, &sub_ksps));
  sub_ksp = sub_ksps[0];
  PetscCall(KSPGetIterationNumber(ksp, &its)); /* outer KSP iteration number */
  PetscCall(KSPGetIterationNumber(sub_ksp, &sub_its)); /* inner KSP iteration number */

  if (iter) {
    PetscCall(PetscPrintf(PETSC_COMM_WORLD, "...PostCheck snes iteration %d", iter, check->its0, its, sub_its));
    ksp_ratio = ((PetscReal)(its)) / check->its0;
    maxit = (PetscInt)(ksp_ratio * sub_its + 0.5);
    if (maxit < 2) maxit = 2;
    PetscCall(KSPSetTolerances(sub_ksp, PETSC_DEFAULT, PETSC_DEFAULT, PETSC_DEFAULT, maxit));
    PetscCall(PetscPrintf(PETSC_COMM_WORLD, "...ksp_ratio %g, new maxit %d", ksp_ratio, maxit));
  }
  check->its0 = its; /* save current outer KSP iteration number */
PetscFunctionReturn(PETSC_SUCCESS);
}
/* */
**MatrixFreePreconditioner** - This routine demonstrates the use of a user-provided preconditioner. This code implements just the null preconditioner, which of course is not recommended for general use.

**Input Parameters:**
+ pc - preconditioner
- x - input vector

**Output Parameter:**
- y - preconditioned vector

PetscErrorCode MatrixFreePreconditioner(PC pc, Vec x, Vec y)
{
    PetscFunctionBeginUser;
    PetscCall(VecCopy(x, y));
    PetscFunctionReturn(PETSC_SUCCESS);
}

Table *Jacobian Options* summarizes the various matrix situations that SNES supports. In particular, different linear system matrices and preconditioning matrices are allowed, as well as both matrix-free and application-provided preconditioners. If `ex3.c` is run with the options *-snes_mf* and *-user_precond* then it uses a matrix-free application of the matrix-vector multiple and a user provided matrix-free Jacobian.

<table>
<thead>
<tr>
<th>Matrix Use</th>
<th>Conventional Matrix Formats</th>
<th>Matrix-free versions</th>
</tr>
</thead>
<tbody>
<tr>
<td>Jacobian Matrix</td>
<td>Create matrix with MatCreate(). Assemble matrix</td>
<td>Create matrix with MatCreateShell(). Use MatShellSetOperation() to set various matrix actions, or use MatCreateMFFD() or MatCreateSNESMF().</td>
</tr>
<tr>
<td></td>
<td>with user-defined routine ^</td>
<td></td>
</tr>
<tr>
<td>Preconditioning</td>
<td>Create matrix with MatCreate(). Assemble matrix</td>
<td>Use SNESGetRSP() and KSPGetPC() to access the PC, then use PCSetType(pc, PCSHELL) followed by PCShellSetApply().</td>
</tr>
<tr>
<td>Matrix</td>
<td>with user-defined routine †</td>
<td></td>
</tr>
</tbody>
</table>

^ Use either the generic MatCreate() or a format-specific variant such as MatCreateAIJ().
† Set user-defined matrix formation routine with SNESSetJacobian() or with a DM variant such as DMDASNESSetJacobianLocal().

SNES also provides some less well-integrated code to apply matrix-free finite differencing using an automatically computed measurement of the noise of the functions. This can be selected with *-snes_mf_version 2*; it does not use MatCreateMFFD() but has similar options that start with *-snes_mf_* instead of *-mat_mffd_*.. Note that this alternative prefix only works for version 2 differencing.
2.4.6 Finite Difference Jacobian Approximations

PETSc provides some tools to help approximate the Jacobian matrices efficiently via finite differences. These tools are intended for use in certain situations where one is unable to compute Jacobian matrices analytically, and matrix-free methods do not work well without a preconditioner, due to very poor conditioning. The approximation requires several steps:

- First, one colors the columns of the (not yet built) Jacobian matrix, so that columns of the same color do not share any common rows.
- Next, one creates a `MatFDColoring` data structure that will be used later in actually computing the Jacobian.
- Finally, one tells the nonlinear solvers of `SNES` to use the `SNESComputeJacobianDefaultColor()` routine to compute the Jacobians.

A code fragment that demonstrates this process is given below.

```c
ISColoring iscoloring;
MatFDColoring fdcoloring;
MatColoring coloring;

/*
 This initializes the nonzero structure of the Jacobian. This is artificial
 because clearly if we had a routine to compute the Jacobian we wouldn't
 need to use finite differences.
*/
FormJacobian(snes,x,&J,&J,&user);

/*
 Color the matrix, i.e. determine groups of columns that share no common
 rows. These columns in the Jacobian can all be computed simultaneously.
*/
MatColoringCreate(J,&coloring);
MatColoringSetType(coloring,MATCOLORINGSL);
MatColoringSetFromOptions(coloring);
MatColoringApply(coloring,&iscoloring);
MatColoringDestroy(&coloring);

/*
 Create the data structure that SNESComputeJacobianDefaultColor() uses
 to compute the actual Jacobians via finite differences.
*/
MatFDColoringCreate(J,iscoloring,&fdcoloring);
ISColoringDestroy(&iscoloring);
MatFDColoringSetFunction(fdcoloring,(PetscErrorCode (*)(void))FormFunction,&user);
MatFDColoringSetFromOptions(fdcoloring);

/*
 Tell SNES to use the routine SNESComputeJacobianDefaultColor()
 to compute Jacobians.
*/
SNESetJacobian(snes,J,J,SNESComputeJacobianDefaultColor,fdcoloring);
```

Of course, we are cheating a bit. If we do not have an analytic formula for computing the Jacobian, then how do we know what its nonzero structure is so that it may be colored? Determining the structure is problem dependent, but fortunately, for most structured grid problems (the class of problems for which PETSc was originally designed) if one knows the stencil used for the nonlinear function one can usually fairly easily obtain an estimate of the location of nonzeros in the matrix. This is harder in the unstructured case, but one typically knows where the nonzero entries are from the mesh topology and distribution of degrees of
freedom. If using DMplex (DMplex: Unstructured Grids) for unstructured meshes, the nonzero locations will be identified in DMCreateMatrix() and the procedure above can be used. Most external packages for unstructured meshes have similar functionality.

One need not necessarily use a MatColoring object to determine a coloring. For example, if a grid can be colored directly (without using the associated matrix), then that coloring can be provided to MatFDColoringCreate(). Note that the user must always preset the nonzero structure in the matrix regardless of which coloring routine is used.

PETSc provides the following coloring algorithms, which can be selected using MatColoringSetType() or via the command line argument -mat_coloring_type.

<table>
<thead>
<tr>
<th>Algorithm</th>
<th>MatColoringType</th>
<th>-mat_coloring_type</th>
<th>Parallel</th>
</tr>
</thead>
<tbody>
<tr>
<td>smallest-last [MoreSGH84]</td>
<td>MATCOLORINGSL</td>
<td>sl</td>
<td>No</td>
</tr>
<tr>
<td>largest-first [MoreSGH84]</td>
<td>MATCOLORINGLF</td>
<td>lf</td>
<td>No</td>
</tr>
<tr>
<td>incidence-degree [MoreSGH84]</td>
<td>MATCOLORINGID</td>
<td>id</td>
<td>No</td>
</tr>
<tr>
<td>Jones-Plassmann [JP93]</td>
<td>MATCOLORINGJP</td>
<td>jp</td>
<td>Yes</td>
</tr>
<tr>
<td>Greedy</td>
<td>MATCOLORINGGREEDY</td>
<td>greedy</td>
<td>Yes</td>
</tr>
<tr>
<td>Natural (1 color per column)</td>
<td>MATCOLORINGNATURAL</td>
<td>natural</td>
<td>Yes</td>
</tr>
<tr>
<td>Power ($A^k$ followed by 1-coloring)</td>
<td>MATCOLORINGPOWER</td>
<td>power</td>
<td>Yes</td>
</tr>
</tbody>
</table>

As for the matrix-free computation of Jacobians (Matrix-Free Methods), two parameters affect the accuracy of the finite difference Jacobian approximation. These are set with the command

```
MatFDColoringSetParameters(MatFDColoring fdcoloring, PetscReal rerror, PetscReal umin);
```

The parameter $rerror$ is the square root of the relative error in the function evaluations, $e_{rel}$; the default is the square root of machine epsilon (about $10^{-8}$ in double precision), which assumes that the functions are evaluated approximately to floating-point precision accuracy. The second parameter, $umin$, is a bit more involved; its default is $10^{e-6}$. Column $i$ of the Jacobian matrix (denoted by $F_{i,i}$) is approximated by the formula

$$F'_{i,i} \approx \frac{F(u + h * dx_i) - F(u)}{h}$$

where $h$ is computed via:

$$h = e_{rel} \cdot \begin{cases} u_i, & \text{if } |u_i| > u_{\text{min}} \\ u_{\text{min}} \cdot \text{sign}(u_i), & \text{otherwise.} \end{cases}$$

for MATMFFD_DS or:

$$h = e_{rel} \sqrt{\|u\|}$$

for MATMFFD_WP (default). These parameters may be set from the options database with

```
-mat_fd_coloring_err <err>
-mat_fd_coloring_umin <umin>
-mat_fd_type <htype>
```

Note that MatColoring type MATCOLORINGSL, MATCOLORINGLF, and MATCOLORINGID are sequential algorithms. MATCOLORINGJP and MATCOLORINGGREEDY are parallel algorithms, although in practice they may create more colors than the sequential algorithms. If one computes the coloring iscoloring reasonably with a parallel algorithm or by knowledge of the discretization, the routine MatFDColoringCreate() is scalable. An example of this for 2D distributed arrays is given below that uses the utility routine DMCreatColoring().
Note that the routine `MatFDColoringCreate()` currently is only supported for the AIJ and BAIJ matrix formats.

### 2.4.7 Variational Inequalities

`SNES` can also solve (differential) variational inequalities with box (bound) constraints. These are nonlinear algebraic systems with additional inequality constraints on some or all of the variables: \( L_i \leq u_i \leq H_i \). For example, the pressure variable cannot be negative. Some, or all, of the lower bounds may be negative infinity (indicated to PETSc with `SNES_VI_NINF`) and some, or all, of the upper bounds may be infinity (indicated by `SNES_VI_INF`). The commands

```c
SNESVIISetVariableBounds(SNES, Vec L, Vec H);
SNESVIISetComputeVariableBounds(SNES snes, PetscErrorCode (*compute)(SNES, Vec, Vec))
```

are used to indicate that one is solving a variational inequality. Problems with box constraints can be solved with the reduced space, `SNESNEWTONRQLS`, and semi-smooth `SNESNEWTONSLS` solvers.

The option `-snse_vi_monitor` turns on extra monitoring of the active set associated with the bounds and `-snse_vi_type` allows selecting from several VI solvers, the default is preferred.

`SNESLineSearchSetPreCheck()` and `SNESLineSearchSetPostCheck()` can also be used to control properties of the steps selected by `SNES`.

### 2.4.8 Nonlinear Preconditioning

The mathematical framework of nonlinear preconditioning is explained in detail in [BKST15]. Nonlinear preconditioning in PETSc involves the use of an inner `SNES` instance to define the step for an outer `SNES` instance. The inner instance may be extracted using

```c
SNESGetNPC(SNES snes, SNES *npc);
```

and passed run-time options using the `-npc_` prefix. Nonlinear preconditioning comes in two flavors: left and right. The side may be changed using `-snse_npc_side` or `SNESSetNPCSide()`. Left nonlinear preconditioning redefines the nonlinear function as the action of the nonlinear preconditioner \( M \);

\[
F_M(x) = M(x, b) - x.
\]

Right nonlinear preconditioning redefines the nonlinear function as the function on the action of the nonlinear preconditioner;

\[
F(M(x, b)) = b,
\]

which can be interpreted as putting the preconditioner into "striking distance" of the solution by outer acceleration.

In addition, basic patterns of solver composition are available with the `SNESType SNESCOMPOSITE`. This allows for two or more `SNES` instances to be combined additively or multiplicatively. By command line, a set of `SNES` types may be given by comma separated list argument to `-snse_composite_sneses`. There are additive (`SNES_COMPOSITE_ADDITIVE`), additive with optimal damping (`SNES_COMPOSITE_ADDITIVEOPTIMAL`), and multiplicative (`SNES_COMPOSITE_MULTIPLICATIVE`) variants which may be set with
SNESCompositeSetType(SNES,SNESCompositeType);

New subsolvers may be added to the composite solver with
SNESCompositeAddSNES(SNES,SNESType);

and accessed with
SNESCompositeGetSNES(SNES,PetscInt,SNES *);

2.5 TS: Scalable ODE and DAE Solvers

The TS library provides a framework for the scalable solution of ODEs and DAEs arising from the discretization of time-dependent PDEs.

**Simple Example:** Consider the PDE
\[ u_t = u_{xx} \]
discretized with centered finite differences in space yielding the semi-discrete equation
\[ (u_i)_t = \frac{u_{i+1} - 2u_i + u_{i-1}}{h^2}, \]
\[ u_t = \bar{A}u; \]
or with piecewise linear finite elements approximation in space \( u(x,t) \approx \sum_i \xi_i(t) \phi_i(x) \) yielding the semi-discrete equation
\[ B\xi'(t) = A\xi(t) \]
Now applying the backward Euler method results in
\[ (B - dt^n A)u^{n+1} = Bu^n, \]
in which
\[ u^n_i = \xi_i(t_n) = u(x_i,t_n), \]
\[ \xi'(t_{n+1}) = \frac{u^{n+1}_i - u^n_i}{dt^n}, \]
\( A \) is the stiffness matrix, and \( B \) is the identity for finite differences or the mass matrix for the finite element method.

The PETSc interface for solving time dependent problems assumes the problem is written in the form
\[ F(t,u,\dot{u}) = G(t,u), \quad u(t_0) = u_0. \]

In general, this is a differential algebraic equation (DAE). For ODE with nontrivial mass matrices such as arise in FEM, the implicit/DAE interface significantly reduces overhead to prepare the system for algebraic solvers (SNES/KSP) by having the user assemble the correctly shifted matrix. Therefore this interface is also useful for ODE systems.

To solve an ODE or DAE one uses:
- Function \( F(t,u,\dot{u}) \)

\[ \text{if the matrix } F_{\dot{u}}(t) = \frac{\partial F}{\partial \dot{u}} \text{ is nonsingular then it is an ODE and can be transformed to the standard explicit form, although this transformation may not lead to efficient algorithms.} \]

2.5. TS: Scalable ODE and DAE Solvers 135
The vector \( R \) is an optional location to store the residual. The arguments to the function \( f() \) are the timestep context, current time, input state \( u \), input time derivative \( \dot{u} \), and the (optional) user-provided context \( \text{funP} \). If \( F(t, u, \dot{u}) = \dot{u} \) then one need not call this function.

- **Function \( G(t, u) \), if it is nonzero, is provided with the function**

  \[
  \text{TSSetRHSFunction(TS ts, Vec R, PetscErrorCode (*f)(TS, PetscReal, Vec, Vec, \text{void*}), void*, \text{~funP};)}
  \]

- **Jacobian \( \sigma F_u(t^n, u^n, \dot{u}^n) + F_u(t^n, u^n, \dot{u}^n) \)**

  If using a fully implicit or semi-implicit (IMEX) method one also can provide an appropriate (approximate) Jacobian matrix of \( F() \).

  \[
  \text{TSSetIJacobian(TS ts, Mat A, Mat B, PetscErrorCode (*fjac)(TS, PetscReal, Vec, Vec, \text{PetscReal, Mat, Mat, \text{void*}}), \text{void *jacP};)}
  \]

The arguments for the function \( fjac() \) are the timestep context, current time, input state \( u \), input derivative \( \dot{u} \), input shift \( \sigma \), matrix \( A \), preconditioning matrix \( B \), and the (optional) user-provided context \( \text{jacP} \).

The Jacobian needed for the nonlinear system is, by the chain rule,

\[
\frac{dF}{du^n} = \frac{\partial F}{\partial \dot{u}} \frac{\partial \dot{u}}{\partial u^n} u^n + \frac{\partial F}{\partial u} u^n.
\]

For any ODE integration method the approximation of \( \dot{u} \) is linear in \( u^n \) hence \( \frac{\partial \dot{u}}{\partial u} u^n = \sigma \), where the shift \( \sigma \) depends on the ODE integrator and time step but not on the function being integrated. Thus

\[
\frac{dF}{du^n} = \sigma F_u(t^n, u^n, \dot{u}^n) + F_u(t^n, u^n, \dot{u}^n).
\]

This explains why the user provide Jacobian is in the given form for all integration methods. An equivalent way to derive the formula is to note that

\[
F(t^n, u^n, \dot{u}^n) = F(t^n, u^n, w + \sigma * u^n)
\]

where \( w \) is some linear combination of previous time solutions of \( u \) so that

\[
\frac{dF}{du^n} = \sigma F_u(t^n, u^n, \dot{u}^n) + F_u(t^n, u^n, \dot{u}^n)
\]

again by the chain rule.

For example, consider backward Euler’s method applied to the ODE \( F(t, u, \dot{u}) = \dot{u} - f(t, u) \) with \( \dot{u} = (u^n - u^{n-1})/\delta t \) and \( \frac{\partial \dot{u}}{\partial u} u^n = 1/\delta t \) resulting in

\[
\frac{dF}{du^n} = (1/\delta t)F_u + F_u(t^n, u^n, \dot{u}^n).
\]

But \( F_u = 1 \), in this special case, resulting in the expected Jacobian \( 1/\delta t - f_u(t, u^n) \).

- **Jacobian \( G_u \)**

  If using a fully implicit method and the function \( G() \) is provided, one also can provide an appropriate (approximate) Jacobian matrix of \( G() \).
The arguments for the function `fjac()` are the timestep context, current time, input state \(u\), matrix \(A\), preconditioning matrix \(B\), and the (optional) user-provided context `jacP`.

Providing appropriate \(F()\) and \(G()\) for your problem allows for the easy runtime switching between explicit, semi-implicit (IMEX), and fully implicit methods.

### 2.5.1 Basic TS Options

The user first creates a TS object with the command

```c
int TSCreate(MPI_Comm comm, TS *ts);
```

```c
int TSSetProblemType(TS ts, TSProblemType problemtype);
```

The `TSProblemType` is one of `TS_LINEAR` or `TS_NONLINEAR`.

To set up TS for solving an ODE, one must set the “initial conditions” for the ODE with

```c
TSSetSolution(TS ts, Vec initialsolution);
```

One can set the solution method with the routine

```c
TSSetType(TS ts, TSType type);
```

Some of the currently supported types are `TSEULER`, `TSRK` (Runge-Kutta), `TSBEULER`, `TSCN` (Crank-Nicolson), `TSTHETA`, `TSGLLE` (generalized linear), `TSPSEUDO`, and `TSSUNDIALS` (only if the Sundials package is installed), or the command line option `-ts_type euler,rk,beuler,cn,theta,gl,pseudo,sundials,eimex,arkimex,rosw`.

A list of available methods is given in `integrator_table`.

Set the initial time with the command

```c
TSSetTime(TS ts, PetscReal time);
```

One can change the timestep with the command

```c
TSSetTimeStep(TS ts, PetscReal dt);
```

One can determine the current timestep with the routine

```c
TSGetTimeStep(TS ts, PetscReal* dt);
```

Here, “current” refers to the timestep being used to attempt to promote the solution from \(u^n\) to \(u^{n+1}\).

One sets the total number of timesteps to run or the total time to run (whatever is first) with the commands

```c
TSSetMaxSteps(TS ts, PetscInt maxsteps);
TSSetMaxTime(TS ts, PetscReal maxtime);
```

and determines the behavior near the final time with
TSSetExactFinalTime(TS ts, TSExactFinalTimeOption eftopt);

where `eftopt` is one of `TS_EXACTFINALTIME_STEPOVER`, `TS_EXACTFINALTIME_INTERPOLATE`, or `TS_EXACTFINALTIME_MATCHSTEP`. One performs the requested number of time steps with

TSSolve(TS ts, Vec U);

The solve call implicitly sets up the timestep context; this can be done explicitly with

TSSetUp(TS ts);

One destroys the context with

TSDestroy(TS *ts);

and views it with

TView(TS ts, PetscViewer viewer);

In place of `TSSolve()`, a single step can be taken using

TSStep(TS ts);

### 2.5.2 DAE Formulations

You can find a discussion of DAEs in [AP98] or Scholarpedia. In PETSc, TS deals with the semi-discrete form of the equations, so that space has already been discretized. If the DAE depends explicitly on the coordinate `x`, then this will just appear as any other data for the equation, not as an explicit argument. Thus we have

\[ F(t, u, \dot{u}) = 0 \]

In this form, only fully implicit solvers are appropriate. However, specialized solvers for restricted forms of DAE are supported by PETSc. Below we consider an ODE which is augmented with algebraic constraints on the variables.

**Hessenberg Index-1 DAE**

This is a Semi-Explicit Index-1 DAE which has the form

\[ \dot{u} = f(t, u, z) \]
\[ 0 = h(t, u, z) \]

where `z` is a new constraint variable, and the Jacobian \( \frac{dh}{dz} \) is non-singular everywhere. We have suppressed the `x` dependence since it plays no role here. Using the non-singularity of the Jacobian and the Implicit Function Theorem, we can solve for `z` in terms of `u`. This means we could, in principle, plug `z(u)` into the first equation to obtain a simple ODE, even if this is not the numerical process we use. Below we show that this type of DAE can be used with IMEX schemes.
Hessenberg Index-2 DAE

This DAE has the form
\[ \dot{u} = f(t, u, z) \]
\[ 0 = h(t, u) \]

Notice that the constraint equation \( h \) is not a function of the constraint variable \( z \). This means that we cannot naively invert as we did in the index-1 case. Our strategy will be to convert this into an index-1 DAE using a time derivative, which loosely corresponds to the idea of an index being the number of derivatives necessary to get back to an ODE. If we differentiate the constraint equation with respect to time, we can use the ODE to simplify it,
\[ 0 = \dot{h}(t, u) = \frac{dh}{du} \dot{u} + \frac{\partial h}{\partial t} = \frac{dh}{du} f(t, u, z) + \frac{\partial h}{\partial t} \]

If the Jacobian \( \frac{dh}{du} \frac{df}{dz} \) is non-singular, then we have precisely a semi-explicit index-1 DAE, and we can once again use the PETSc IMEX tools to solve it. A common example of an index-2 DAE is the incompressible Navier-Stokes equations, since the continuity equation \( \nabla \cdot u = 0 \) does not involve the pressure. Using PETSc IMEX with the above conversion then corresponds to the Segregated Runge-Kutta method applied to this equation [ColomesB16].

2.5.3 Using Implicit-Explicit (IMEX) Methods

For “stiff” problems or those with multiple time scales \( F() \) will be treated implicitly using a method suitable for stiff problems and \( G() \) will be treated explicitly when using an IMEX method like TSARKIMEX. \( F() \) is typically linear or weakly nonlinear while \( G() \) may have very strong nonlinearities such as arise in non-oscillatory methods for hyperbolic PDE. The user provides three pieces of information, the APIs for which have been described above.

- “Slow” part \( G(t, u) \) using \text{TSSetRHSFunction}().
- “Stiff” part \( F(t, u, \dot{u}) \) using \text{TSSetIFunction}().
- Jacobian \( F_u + \sigma F_\dot{u} \) using \text{TSSetIJacobian}().

The user needs to set \text{TSSetEquationType()} to \text{TS_EQ_IMPLICIT} or higher if the problem is implicit; e.g., \( F(t, u, \dot{u}) = M \dot{u} - f(t, u) \), where \( M \) is not the identity matrix:

- the problem is an implicit ODE (defined implicitly through \text{TSSetIFunction}()) or
- a DAE is being solved.

An IMEX problem representation can be made implicit by setting \text{TSARKIMEXSetFullyImplicit}().

In PETSc, DAEs and ODEs are formulated as \( F(t, u, \dot{u}) = G(t, u) \), where \( F() \) is meant to be integrated implicitly and \( G() \) explicitly. An IMEX formulation such as \( M \dot{u} = f(t, u) + g(t, u) \) requires the user to provide \( M^{-1} g(t, u) \) or solve \( g(t, u) - M x = 0 \) in place of \( G(t, u) \). General cases such as \( F(t, u, \dot{u}) = G(t, u) \) are not amenable to IMEX Runge-Kutta, but can be solved by using fully implicit methods. Some use-case examples for \text{TSARKIMEX} are listed in Table 2.12 and a list of methods with a summary of their properties is given in IMEX Runge-Kutta schemes.
Table 2.12: Use case examples for **TSARKIMEX**

<table>
<thead>
<tr>
<th>Equation</th>
<th>Type</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>( \dot{u} = g(t, u) )</td>
<td>nonstiff ODE</td>
<td>( F(t, u, \dot{u}) = \dot{u} )</td>
</tr>
<tr>
<td></td>
<td></td>
<td>( G(t, u) = g(t, u) )</td>
</tr>
<tr>
<td>( M\dot{u} = g(t, u) )</td>
<td>nonstiff ODE with mass matrix</td>
<td>( F(t, u, \dot{u}) = \dot{u} )  ( G(t, u) = M^{-1}g(t, u) )</td>
</tr>
<tr>
<td>( \dot{u} = f(t, u) )</td>
<td>stiff ODE</td>
<td>( F(t, u, \dot{u}) = \dot{u} - f(t, u) )</td>
</tr>
<tr>
<td></td>
<td></td>
<td>( G(t, u) = 0 )</td>
</tr>
<tr>
<td>( M\dot{u} = f(t, u) )</td>
<td>stiff ODE with mass matrix</td>
<td>( F(t, u, \dot{u}) = M\dot{u} - f(t, u) )</td>
</tr>
<tr>
<td></td>
<td></td>
<td>( G(t, u) = 0 )</td>
</tr>
<tr>
<td>( \dot{u} = f(t, u) + g(t, u) )</td>
<td>stiff-nonstiff ODE</td>
<td>( F(t, u, \dot{u}) = \dot{u} - f(t, u) )</td>
</tr>
<tr>
<td></td>
<td></td>
<td>( G(t, u) = g(t, u) )</td>
</tr>
<tr>
<td>( M\dot{u} = f(t, u) + g(t, u) )</td>
<td>stiff-nonstiff ODE with mass matrix</td>
<td>( F(t, u, \dot{u}) = M\dot{u} - f(t, u) )</td>
</tr>
<tr>
<td></td>
<td></td>
<td>( G(t, u) = M^{-1}g(t, u) )</td>
</tr>
<tr>
<td>( \dot{u} = f(t, u, z) + g(t, u, z) )</td>
<td>semi-explicit index-1 DAE</td>
<td>( F(t, u, \dot{u}) = \begin{pmatrix} \dot{u} - f(t, u, z) \ h(t, u, z) \end{pmatrix} )</td>
</tr>
<tr>
<td>0 = h(t, y, z)</td>
<td></td>
<td>( G(t, u) = g(t, u) )</td>
</tr>
<tr>
<td>( f(t, u, \dot{u}) = 0 )</td>
<td>fully implicit ODE/DAE</td>
<td>( F(t, u, \dot{u}) = f(t, u, \dot{u}) ); the user needs to set TS-SetEquationType() to TS_EQ_IMPLICIT or higher</td>
</tr>
</tbody>
</table>

Table 2.13 lists the currently available IMEX Runge-Kutta schemes. For each method, it gives the **-ts_arkimex_type** name, the reference, the total number of stages/implicit stages, the order/stage-order, the implicit stability properties (IM), stiff accuracy (SA), the existence of an embedded scheme, and dense output (DO).
<table>
<thead>
<tr>
<th>Name</th>
<th>Reference</th>
<th>Stages (IM)</th>
<th>Order (Stage)</th>
<th>IM</th>
<th>SA</th>
<th>Embed</th>
<th>DO</th>
<th>Remarks</th>
</tr>
</thead>
<tbody>
<tr>
<td>a2</td>
<td>based on CN</td>
<td>2 (1)</td>
<td>2 (2)</td>
<td>A-Stable</td>
<td>yes</td>
<td>yes (1)</td>
<td>yes (2)</td>
<td></td>
</tr>
<tr>
<td>l2</td>
<td>SSP2(2,2,2) [PR05]</td>
<td>2 (2)</td>
<td>2 (1)</td>
<td>L-Stable</td>
<td>yes</td>
<td>yes (1)</td>
<td>yes (2)</td>
<td>SSP SDIRK</td>
</tr>
<tr>
<td>ars122</td>
<td>ARS122 [ARS97]</td>
<td>2 (1)</td>
<td>3 (1)</td>
<td>A-Stable</td>
<td>yes</td>
<td>yes (1)</td>
<td>yes (2)</td>
<td></td>
</tr>
<tr>
<td>2c</td>
<td>[GKC13]</td>
<td>3 (2)</td>
<td>2 (2)</td>
<td>L-Stable</td>
<td>yes</td>
<td>yes (1)</td>
<td>yes (2)</td>
<td>SDIRK</td>
</tr>
<tr>
<td>2d</td>
<td>[GKC13]</td>
<td>3 (2)</td>
<td>2 (2)</td>
<td>L-Stable</td>
<td>yes</td>
<td>yes (1)</td>
<td>yes (2)</td>
<td>SDIRK</td>
</tr>
<tr>
<td>2e</td>
<td>[GKC13]</td>
<td>3 (2)</td>
<td>2 (2)</td>
<td>L-Stable</td>
<td>yes</td>
<td>yes (1)</td>
<td>yes (2)</td>
<td>SDIRK</td>
</tr>
<tr>
<td>prssp2</td>
<td>PRS(3,3,2) [PR05]</td>
<td>3 (3)</td>
<td>3 (1)</td>
<td>L-Stable</td>
<td>yes</td>
<td>no</td>
<td>no</td>
<td>SSP</td>
</tr>
<tr>
<td>3</td>
<td>[KC03]</td>
<td>4 (3)</td>
<td>3 (2)</td>
<td>L-Stable</td>
<td>yes</td>
<td>yes (2)</td>
<td>yes (2)</td>
<td>SDIRK</td>
</tr>
<tr>
<td>bpr3</td>
<td>[BPR11]</td>
<td>5 (4)</td>
<td>3 (2)</td>
<td>L-Stable</td>
<td>yes</td>
<td>no</td>
<td>no</td>
<td>SDIRK</td>
</tr>
<tr>
<td>ars443</td>
<td>ARS97 [ARS97]</td>
<td>5 (4)</td>
<td>3 (1)</td>
<td>L-Stable</td>
<td>yes</td>
<td>no</td>
<td>no</td>
<td>SDIRK</td>
</tr>
<tr>
<td>4</td>
<td>[KC03]</td>
<td>6 (5)</td>
<td>4 (2)</td>
<td>L-Stable</td>
<td>yes</td>
<td>yes (3)</td>
<td>yes</td>
<td>SDIRK</td>
</tr>
<tr>
<td>5</td>
<td>[KC03]</td>
<td>8 (7)</td>
<td>5 (2)</td>
<td>L-Stable</td>
<td>yes</td>
<td>yes (4)</td>
<td>yes (3)</td>
<td>SDIRK</td>
</tr>
</tbody>
</table>

ROSW are linearized implicit Runge-Kutta methods known as Rosenbrock W-methods. They can accommodate inexact Jacobian matrices in their formulation. A series of methods are available in PETSc are listed in Table 2.14 below. For each method, it gives the reference, the total number of stages and implicit stages, the scheme order and stage order, the implicit stability properties (IM), stiff accuracy (SA), the existence of an embedded scheme, dense output (DO), the capacity to use inexact Jacobian matrices (-W), and high order integration of differential algebraic equations (PDAE).
Table 2.14: Rosenbrock W-schemes

<table>
<thead>
<tr>
<th>TS</th>
<th>Reference</th>
<th>Stages (IM)</th>
<th>Order (Stage)</th>
<th>IM</th>
<th>SA</th>
<th>Embed</th>
<th>DO</th>
<th>-W</th>
<th>PDAE</th>
<th>Remarks</th>
</tr>
</thead>
<tbody>
<tr>
<td>theta1</td>
<td>classical</td>
<td>1(1)</td>
<td>1(1)</td>
<td>L-Stable</td>
<td>•</td>
<td>•</td>
<td>•</td>
<td>•</td>
<td>•</td>
<td></td>
</tr>
<tr>
<td>theta2</td>
<td>classical</td>
<td>1(1)</td>
<td>2(2)</td>
<td>A-Stable</td>
<td>•</td>
<td>•</td>
<td>•</td>
<td>•</td>
<td>•</td>
<td></td>
</tr>
<tr>
<td>2m</td>
<td>Zoltan</td>
<td>2(2)</td>
<td>2(1)</td>
<td>L-Stable</td>
<td>No</td>
<td>Yes(1)</td>
<td>Yes(2)</td>
<td>Yes</td>
<td>No</td>
<td>SSP</td>
</tr>
<tr>
<td>2p</td>
<td>Zoltan</td>
<td>2(2)</td>
<td>2(1)</td>
<td>L-Stable</td>
<td>No</td>
<td>Yes(1)</td>
<td>Yes(2)</td>
<td>Yes</td>
<td>No</td>
<td>SSP</td>
</tr>
<tr>
<td>ra3pw</td>
<td>[RA05]</td>
<td>3(3)</td>
<td>3(1)</td>
<td>A-Stable</td>
<td>No</td>
<td>Yes</td>
<td>Yes(2)</td>
<td>No</td>
<td>Yes(3)</td>
<td></td>
</tr>
<tr>
<td>ra34pw2</td>
<td>[RA05]</td>
<td>4(4)</td>
<td>3(1)</td>
<td>L-Stable</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes(3)</td>
<td>Yes</td>
<td>Yes(3)</td>
<td></td>
</tr>
<tr>
<td>rodas3</td>
<td>[SVB+97]</td>
<td>4(4)</td>
<td>3(1)</td>
<td>L-Stable</td>
<td>Yes</td>
<td>Yes</td>
<td>No</td>
<td>No</td>
<td>Yes</td>
<td></td>
</tr>
<tr>
<td>sandu3</td>
<td>[SVB+97]</td>
<td>3(3)</td>
<td>3(1)</td>
<td>L-Stable</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes(2)</td>
<td>No</td>
<td>No</td>
<td></td>
</tr>
<tr>
<td>assp3p3</td>
<td>un-pub.</td>
<td>3(2)</td>
<td>3(1)</td>
<td>A-Stable</td>
<td>No</td>
<td>Yes</td>
<td>Yes(2)</td>
<td>Yes</td>
<td>No</td>
<td>SSP</td>
</tr>
<tr>
<td>lassp3p4p2</td>
<td>un-pub.</td>
<td>4(3)</td>
<td>3(1)</td>
<td>L-Stable</td>
<td>No</td>
<td>Yes</td>
<td>Yes(3)</td>
<td>Yes</td>
<td>No</td>
<td>SSP</td>
</tr>
<tr>
<td>lassp3p4p2</td>
<td>un-pub.</td>
<td>4(3)</td>
<td>3(1)</td>
<td>L-Stable</td>
<td>No</td>
<td>Yes</td>
<td>Yes(3)</td>
<td>Yes</td>
<td>No</td>
<td>SSP</td>
</tr>
<tr>
<td>ark3</td>
<td>un-pub.</td>
<td>4(3)</td>
<td>3(1)</td>
<td>L-Stable</td>
<td>No</td>
<td>Yes</td>
<td>Yes(3)</td>
<td>Yes</td>
<td>No</td>
<td>IMEX-RK</td>
</tr>
</tbody>
</table>

2.5.4 GLEE methods

In this section, we describe explicit and implicit time stepping methods with global error estimation that are introduced in [Con16]. The solution vector for a GLEE method is either $[y, \tilde{y}]$ or $[y, \varepsilon]$, where $y$ is the solution, $\tilde{y}$ is the “auxiliary solution,” and $\varepsilon$ is the error. The working vector that TSGLEE uses is $Y = [y, \tilde{y}]$, or $[y, \varepsilon]$. A GLEE method is defined by

- $(p, r, s)$: (order, steps, and stages),
- $\gamma$: factor representing the global error ratio,
- $A, U, B, V$: method coefficients,
- $S$: starting method to compute the working vector from the solution (say at the beginning of time integration) so that $Y = Sy$,
- $F$: finalizing method to compute the solution from the working vector $y = FY$,
- $F_{\text{embed}}$: coefficients for computing the auxiliary solution $\tilde{y}$ from the working vector $(\tilde{y} = F_{\text{embed}}Y)$,
- $F_{\text{error}}$: coefficients to compute the estimated error vector from the working vector $(\varepsilon = F_{\text{error}}Y)$,
- $S_{\text{error}}$: coefficients to initialize the auxiliary solution $(\tilde{y} \text{ or } \varepsilon)$ from a specified error vector $(\varepsilon)$. It is currently implemented only for $r = 2$. We have $y_{\text{aux}} = S_{\text{error}}[0] \ast \varepsilon + S_{\text{error}}[1] \ast y$, where $y_{\text{aux}}$ is the 2nd component of the working vector $Y$. 

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The methods can be described in two mathematically equivalent forms: propagate two components ("$y\tilde{y}$ form") and propagating the solution and its estimated error ("$y\varepsilon$ form"). The two forms are not explicitly specified in TSGLEE; rather, the specific values of $B, U, S, F, F_{\text{embed}}$, and $F_{\text{error}}$ characterize whether the method is in $y\tilde{y}$ or $y\varepsilon$ form.

The API used by this TS method includes:

- **TSGetSolutionComponents**: Get all the solution components of the working vector
  \[
  \text{ierr} = \text{TSGetSolutionComponents}(\text{TS}, \text{int}^*, \text{Vec}^*)
  \]
  Call with **NULL** as the last argument to get the total number of components in the working vector $Y$ (this is $r$ (not $r - 1$)), then call to get the $i$-th solution component.

- **TSGetAuxSolution**: Returns the auxiliary solution $\tilde{y}$ (computed as $F_{\text{embed}}Y$)
  \[
  \text{ierr} = \text{TSGetAuxSolution}(\text{TS}, \text{Vec}^*)
  \]

- **TSGetTimeError**: Returns the estimated error vector $\varepsilon$ (computed as $F_{\text{error}}Y$ if $n = 0$ or restores the error estimate at the end of the previous step if $n = -1$)
  \[
  \text{ierr} = \text{TSGetTimeError}(\text{TS}, \text{PetscInt } n, \text{Vec}^*)
  \]

- **TSSetTimeError**: Initializes the auxiliary solution ($\tilde{y}$ or $\varepsilon$) for a specified initial error.
  \[
  \text{ierr} = \text{TSSetTimeError}(\text{TS}, \text{Vec})
  \]

The local error is estimated as $\varepsilon(n+1) - \varepsilon(n)$. This is to be used in the error control. The error in $y\tilde{y}$ GLEE is $\varepsilon(n) = \frac{1}{1-\gamma} * (\tilde{y}(n) - y(n))$.

Note that $y$ and $\tilde{y}$ are reported to TSAdapt basic (TSADAPTBASIC), and thus it computes the local error as $\varepsilon_{\text{loc}} = (\tilde{y} - y)$. However, the actual local error is $\varepsilon_{\text{loc}} = \varepsilon_{n+1} - \varepsilon_{n} = \frac{1}{1-\gamma} * [(\tilde{y} - y)_{n+1} - (\tilde{y} - y)_{n}]$.

Table 2.15 lists currently available GL schemes with global error estimation [Con16].

<table>
<thead>
<tr>
<th>TS</th>
<th>Reference</th>
<th>IM/EX</th>
<th>$(p, r, s)$</th>
<th>$\gamma$</th>
<th>Form</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td>TSGLEEi1</td>
<td>BE1</td>
<td>IM</td>
<td>(1, 3, 2)</td>
<td>0.5</td>
<td>$y\varepsilon$</td>
<td>Based on backward Euler</td>
</tr>
<tr>
<td>TSGLEE23</td>
<td>23</td>
<td>EX</td>
<td>(2, 3, 2)</td>
<td>0</td>
<td>$y\varepsilon$</td>
<td></td>
</tr>
<tr>
<td>TSGLEE24</td>
<td>24</td>
<td>EX</td>
<td>(2, 4, 2)</td>
<td>0</td>
<td>$y\tilde{y}$</td>
<td></td>
</tr>
<tr>
<td>TSGLEE251</td>
<td>25i</td>
<td>EX</td>
<td>(2, 5, 2)</td>
<td>0</td>
<td>$y\tilde{y}$</td>
<td></td>
</tr>
<tr>
<td>TSGLEE35</td>
<td>35</td>
<td>EX</td>
<td>(3, 5, 2)</td>
<td>0</td>
<td>$y\tilde{y}$</td>
<td></td>
</tr>
<tr>
<td>TSGLEEEXRK2A</td>
<td>exrk2a</td>
<td>EX</td>
<td>(2, 6, 2)</td>
<td>0.25</td>
<td>$y\varepsilon$</td>
<td></td>
</tr>
<tr>
<td>TSGLEERK32G1</td>
<td>rk32g1</td>
<td>EX</td>
<td>(3, 8, 2)</td>
<td>0</td>
<td>$y\varepsilon$</td>
<td></td>
</tr>
<tr>
<td>TSGLEERK285EX</td>
<td>rk285ex</td>
<td>EX</td>
<td>(2, 9, 2)</td>
<td>0.25</td>
<td>$y\varepsilon$</td>
<td></td>
</tr>
</tbody>
</table>

### 2.5.5 Using fully implicit methods

To use a fully implicit method like TSTHETA, TSBDF or TSDIRK, either provide the Jacobian of $F()$ (and $G()$ if $G()$ is provided) or use a DM that provides a coloring so the Jacobian can be computed efficiently via finite differences.
2.5.6 Using the Explicit Runge-Kutta timestepper with variable timesteps

The explicit Euler and Runge-Kutta methods require the ODE be in the form
\[ \dot{u} = G(u,t). \]

The user can either call `TSSetRHSFunction()` and/or they can call `TSSetIFunction()` (so long as the function provided to `TSSetIFunction()` is equivalent to \( \dot{u} + \tilde{F}(t, u) \)) but the Jacobians need not be provided.\(^5\)

The Explicit Runge-Kutta timestepper with variable timesteps is an implementation of the standard Runge-Kutta with an embedded method. The error in each timestep is calculated using the solutions from the Runge-Kutta method and its embedded method (the 2-norm of the difference is used). The default method is the 3rd-order Bogacki-Shampine method with a 2nd-order embedded method (TSRK3BS). Other available methods are the 5th-order Fehlberg RK scheme with a 4th-order embedded method (TSRK5F), the 5th-order Dormand-Prince RK scheme with a 4th-order embedded method (TSRK5DP), the 5th-order Bogacki-Shampine RK scheme with a 4th-order embedded method (TSRK5BS), and the 6th-, 7th, and 8th-order robust Verner RK schemes with a 5th-, 6th, and 7th-order embedded method, respectively (TSRK6VR, TSRK7VR, TSRK8VR). Variable timesteps cannot be used with RK schemes that do not have an embedded method (TSRK1FE - 1st-order, 1-stage forward Euler, TSRK2A - 2nd-order, 2-stage RK scheme, TSRK3 - 3rd-order, 3-stage RK scheme, TSRK4 - 4th-order, 4-stage RK scheme).

2.5.7 Special Cases

- **\( \dot{u} = Au \).** First compute the matrix \( A \) then call

\begin{verbatim}
TSSetProblemType(ts, TS_LINEAR);
TSSetRHSFunction(ts, NULL, TSComputeRHSFunctionLinear, NULL);
TSSetRHSJacobian(ts, A, A, TSComputeRHSJacobianConstant, NULL);
\end{verbatim}

or

\begin{verbatim}
TSSetProblemType(ts, TS_LINEAR);
TSSetIFunction(ts, NULL, TSComputeIFunctionLinear, NULL);
TSSetIJacobian(ts, A, A, TSComputeIJacobianConstant, NULL);
\end{verbatim}

- **\( \dot{u} = A(t)u \).** Use

\begin{verbatim}
TSSetProblemType(ts, TS_LINEAR);
TSSetRHSFunction(ts, NULL, TSComputeRHSFunctionLinear, NULL);
TSSetRHSJacobian(ts, A, A, YourComputeRHSJacobian, &appctx);
\end{verbatim}

where `YourComputeRHSJacobian()` is a function you provide that computes \( A \) as a function of time. Or use

\begin{verbatim}
TSSetProblemType(ts, TS_LINEAR);
TSSetIFunction(ts, NULL, TSComputeIFunctionLinear, NULL);
TSSetIJacobian(ts, A, A, YourComputeIJacobian, &appctx);
\end{verbatim}

\(^5\) PETSc will automatically translate the function provided to the appropriate form.
2.5.8 Monitoring and visualizing solutions

- `-ts_monitor` - prints the time and timestep at each iteration.
- `-ts_adapt_monitor` - prints information about the timestep adaptation calculation at each iteration.
- `-ts_monitor_lg_timestep` - plots the size of each timestep, `TSMonitorLGTimeStep()`.
- `-ts_monitor_lg_solution` - for ODEs with only a few components (not arising from the discretization of a PDE) plots the solution as a function of time, `TSMonitorLGSolution()`.
- `-ts_monitor_lg_error` - for ODEs with only a few components plots the error as a function of time, only if `TSSetSolutionFunction()` is provided, `TSMonitorLGEError()`.
- `-ts_monitor_draw_solution` - plots the solution at each iteration, `TSMonitorDrawSolution()`.
- `-ts_monitor_draw_error` - plots the error at each iteration only if `TSSetSolutionFunction()` is provided, `TSMonitorDrawError()`.
- `-ts_monitor_solution_binary[:filename]` - saves the solution at each iteration to a binary file, `TSMonitorSolution()`.
- `-ts_monitor_solution_vtk <filename-%03D.vts>` - saves the solution at each iteration to a file in vtk format, `TSMonitorSolutionVTK()`.

2.5.9 Error control via variable time-stepping

Most of the time stepping methods available in PETSc have an error estimation and error control mechanism. This mechanism is implemented by changing the step size in order to maintain user specified absolute and relative tolerances. The PETSc object responsible with error control is `TSAdapt`. The available `TSAdapt` types are listed in the following table.

<table>
<thead>
<tr>
<th>ID</th>
<th>Name</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td>TSADAPT NONE</td>
<td>none</td>
<td>no adaptivity</td>
</tr>
<tr>
<td>TSADAPT BASIC</td>
<td>basic</td>
<td>the default adaptor</td>
</tr>
<tr>
<td>TSADAPT GLEE</td>
<td>glee</td>
<td>extension of the basic adaptor to treat $Tol_A$ and $Tol_R$ as separate criteria. It can also control global errors if the integrator (e.g., <code>TSGLEE</code>) provides this information</td>
</tr>
</tbody>
</table>

When using `TSADAPT_BASIC` (the default), the user typically provides a desired absolute $Tol_A$ or a relative $Tol_R$ error tolerance by invoking `TSSetTolerances()` or at the command line with options `-ts_atol` and `-ts_rtol`. The error estimate is based on the local truncation error, so for every step the algorithm verifies that the estimated local truncation error satisfies the tolerances provided by the user and computes a new step size to be taken. For multistage methods, the local truncation is obtained by comparing the solution $y$ to a lower order $\tilde{y} = p - 1$ approximation, where $p$ is the order of the method and $\tilde{p}$ the order of $\tilde{y}$.

The adaptive controller at step $n$ computes a tolerance level

$$Tol_n(i) = Tol_A(i) + \max(y_n(i), \tilde{y}_n(i))Tol_R(i),$$

2.5. TS: Scalable ODE and DAE Solvers
and forms the acceptable error level

\[ \text{wlte}_n = \frac{1}{m} \sum_{i=1}^{m} \sqrt{\frac{\|y_n(i) - \hat{y}_n(i)\|}{Tol(i)}}, \]

where the errors are computed componentwise, \( m \) is the dimension of \( y \) and \(-\text{ts_adapt_wnormtype}\) is 2 (default). If \(-\text{ts_adapt_wnormtype}\) is \( \text{infinity} \) (max norm), then

\[ \text{wlte}_n = \max_{1 \leq i \leq m} \frac{\|y_n(i) - \hat{y}_n(i)\|}{Tol(i)}. \]

The error tolerances are satisfied when \( \text{wlte} \leq 1.0 \).

The next step size is based on this error estimate, and determined by

\[ \Delta t_{\text{new}}(t) = \Delta t_{\text{old}} \min(\alpha_{\text{max}}, \max(\alpha_{\text{min}}, \beta(1/\text{wlte})^{1/\alpha})) \tag{2.5} \]

where \( \alpha_{\text{min}} = -\text{ts_adapt_clip}[0] \) and \( \alpha_{\text{max}} = -\text{ts_adapt_clip}[1] \) keep the change in \( \Delta t \) to within a certain factor, and \( \beta < 1 \) is chosen through \(-\text{ts_adapt_safety}\) so that there is some margin to which the tolerances are satisfied and so that the probability of rejection is decreased.

This adaptive controller works in the following way. After completing step \( k \), if \( \text{wlte}_{k+1} \leq 1.0 \), then the step is accepted and the next step is modified according to eq:hnnew; otherwise, the step is rejected and retaken with the step length computed in (2.5).

\text{TSAADAPTGLEE} is an extension of the basic adaptor to treat \( \text{Tol}_A \) and \( \text{Tol}_R \) as separate criteria. It can also control global errors if the integrator (e.g., \text{TSGLEE}) provides this information.

### 2.5.10 Handling of discontinuities

For problems that involve discontinuous right hand sides, one can set an “event” function \( g(t, u) \) for PETSc to detect and locate the times of discontinuities (zeros of \( g(t, u) \)). Events can be defined through the event monitoring routine

```c
TSSetEventHandler(TS ts, PetscInt nevents, PetscInt *direction, PetscBool *terminate, 
    PetscErrorCode (*eventhandler)(TS, PetscReal, Vec, PetscScalar *, void *eventP), 
    PetscErrorCode (*postevent)(TS, PetscInt, PetscInt[], PetscReal, Vec, PetscBool, void *eventP));
```

Here, \text{nevents} denotes the number of events, \text{direction} sets the type of zero crossing to be detected for an event (+1 for positive zero-crossing, -1 for negative zero-crossing, and 0 for both), \text{terminate} conveys whether the time-stepping should continue or halt when an event is located, \text{eventmonitor} is a user-defined routine that specifies the event description, \text{postevent} is an optional user-defined routine to take specific actions following an event.

The arguments to \text{eventhandler}() are the timestep context, current time, input state \( u \), array of event function value, and the (optional) user-provided context \text{eventP}.

The arguments to \text{postevent}() routine are the timestep context, number of events occurred, indices of events occured, current time, input state \( u \), a boolean flag indicating forward solve (1) or adjoint solve (0), and the (optional) user-provided context \text{eventP}.
2.5.11 Explicit integrators with finite element mass matrices

Discretized finite element problems often have the form \( M \dot{u} = G(t, u) \) where \( M \) is the mass matrix. Such problems can be solved using \texttt{DMTSSetIFunction()} with implicit integrators. When \( M \) is nonsingular (i.e., the problem is an ODE, not a DAE), explicit integrators can be applied to \( \dot{u} = M^{-1}G(t, u) \) or \( \dot{u} = \hat{M}^{-1}G(t, u) \), where \( \hat{M} \) is the lumped mass matrix. While the true mass matrix generally has a dense inverse and thus must be solved iteratively, the lumped mass matrix is diagonal (e.g., computed via collocated quadrature or row sums of \( M \)). To have PETSc create and apply a (lumped) mass matrix automatically, first use \texttt{DMTSSetRHSFunction()} to specify \( :math:`G` \) and set a ``PetscFE`` using ``DMAddField()`` and ``DMCreateDS()``, then call either \texttt{DMTSCreateRHSMassMatrix()} or \texttt{DMTSCreateRHSMassMatrixLumped()} to automatically create the mass matrix and a KSP that will be used to apply \( M^{-1} \). This KSP can be customized using the "mass_" prefix.

2.5.12 Performing sensitivity analysis with the TS ODE Solvers

The \texttt{TS} library provides a framework based on discrete adjoint models for sensitivity analysis for ODEs and DAEs. The ODE/DAE solution process (henceforth called the forward run) can be obtained by using either explicit or implicit solvers in \texttt{TS}, depending on the problem properties. Currently supported method types are \texttt{TSRK} (Runge-Kutta) explicit methods and \texttt{TSTHETA} implicit methods, which include \texttt{TSBEULER} and \texttt{TSCN}.

Using the discrete adjoint methods

Consider the ODE/DAE

\[
F(t, y, \dot{y}, p) = 0, \quad y(t_0) = y_0(p) \quad t_0 \leq t \leq t_F
\]

and the cost function(s)

\[
\Psi_i(y_0, p) = \Phi_i(y_F, p) + \int_{t_0}^{t_F} r_i(y(t), p, t) dt \quad i = 1, \ldots, n_{\text{cost}}.
\]

The \texttt{TSAdjoint} routines of PETSc provide

\[
\frac{\partial \Psi_i}{\partial y_0} = \lambda_i
\]

and

\[
\frac{\partial \Psi_i}{\partial p} = \mu_i + \lambda_i \left( \frac{\partial y_0}{\partial p} \right).
\]

To perform the discrete adjoint sensitivity analysis one first sets up the \texttt{TS} object for a regular forward run but with one extra function call

\texttt{TSSetSaveTrajectory(TS ts)},

then calls \texttt{TSSolve()} in the usual manner.

One must create two arrays of \( n_{\text{cost}} \) vectors \( \lambda \) and \( \mu \) (if there are no parameters \( p \) then one can use \texttt{NULL} for the \( \mu \) array). The \( \lambda \) vectors are the same dimension and parallel layout as the solution vector for the ODE, the \( \mu \) vectors are of dimension \( p \); when \( p \) is small usually all its elements are on the first MPI process, while the vectors have no entries on the other processes. \( \lambda_i \) and \( \mu_i \) should be initialized with the values \( d\Phi_i/dy|_{t=t_F} \) and \( d\Phi_i/dp|_{t=t_F} \) respectively. Then one calls
TSSetCostGradients(TS ts, PetscInt numcost, Vec *lambda, Vec *mu);

where `numcost` denotes $n_{\text{cost}}$. If $F()$ is a function of $p$ one needs to also provide the Jacobian $-F_p$ with

TSSetRHSJacobianP(TS ts, Mat Amat, PetscErrorCode (*fp)(TS, PetscReal, Vec, Mat, void*), void *ctx)

or

TSSetIJacobianP(TS ts, Mat Amat, PetscErrorCode (*fp)(TS, PetscReal, Vec, Vec, PetscReal, Mat, void*), void *ctx)

or both, depending on which form is used to define the ODE.

The arguments for the function `fp()` are the timestep context, current time, $y$, and the (optional) user-provided context.

If there is an integral term in the cost function, i.e. $r$ is nonzero, it can be transformed into another ODE that is augmented to the original ODE. To evaluate the integral, one needs to create a child TS objective by calling

TSCreateQuadratureTS(TS ts, PetscBool fwd, TS *quadts);

and provide the ODE RHS function (which evaluates the integrand $r$) with

TSSetRHSFunction(TS quadts, Vec R, PetscErrorCode (*rf)(TS, PetscReal, Vec, void*), void *ctx)

Similar to the settings for the original ODE, Jacobians of the integrand can be provided with

TSSetRHSJacobian(TS quadts, Vec DRDU, PetscErrorCode (*drdyf)(TS, PetscReal, Vec, Vec, PetscReal, Vec*, void*), void *ctx)
TSSetRHSJacobianP(TS quadts, Vec DRDU, PetscErrorCode (*drdpf)(TS, PetscReal, Vec, Vec, PetscReal, Vec*, void*), void *ctx)

where $drdyf = dr/dy$, $drdpf = dr/dp$. Since the integral term is additive to the cost function, its gradient information will be included in $\lambda$ and $\mu$.

Lastly, one starts the backward run by calling

TSAdjointSolve(TS ts).

One can obtain the value of the integral term by calling

TSGetCostIntegral(TS ts, Vec *q).

or accessing directly the solution vector used by `quadts`.

The second argument of `TSCreateQuadratureTS()` allows one to choose if the integral term is evaluated in the forward run (inside `TSSolve()`) or in the backward run (inside `TSAdjointSolve()`) when `TSSetCostGradients()` and `TSSetCostIntegrand()` are called before `TSSolve()`. Note that this also allows for evaluating the integral without having to use the adjoint solvers.

To provide a better understanding of the use of the adjoint solvers, we introduce a simple example, corresponding to TS Power Grid Tutorial ex3sa. The problem is to study dynamic security of power system when there are credible contingencies such as short-circuits or loss of generators, transmission lines, or loads. The dynamic security constraints are incorporated as equality constraints in the form of discretized differential
equations and inequality constraints for bounds on the trajectory. The governing ODE system is

\[ \dot{\phi} = \omega_B (\omega - \omega_S) \]

\[ 2H/\omega_S \omega' = p_m - p_{\text{max}} \sin(\phi) - D(\omega - \omega_S), \quad t_0 \leq t \leq t_F, \]

where \( \phi \) is the phase angle and \( \omega \) is the frequency.

The initial conditions at time \( t_0 \) are

\[ \phi(t_0) = \arcsin \left( \frac{p_m}{p_{\text{max}}} \right), \]

\[ w(t_0) = 1. \]

\( p_{\text{max}} \) is a positive number when the system operates normally. At an event such as fault incidence/removal, \( p_{\text{max}} \) will change to 0 temporarily and back to the original value after the fault is fixed. The objective is to maximize \( p_m \) subject to the above ODE constraints and \( \phi < \phi_S \) during all times. To accommodate the inequality constraint, we want to compute the sensitivity of the cost function

\[ \Psi(p_m, \phi) = -p_m + c \int_{t_0}^{t_F} (\max(0, \phi - \phi_S))^2 \, dt \]

with respect to the parameter \( p_m \). numcost is 1 since it is a scalar function.

For ODE solution, PETSc requires user-provided functions to evaluate the system \( F(t, y, \dot{y}, p) \) (set by TS-SetIFunction()) and its corresponding Jacobian \( F_y + \sigma F_{\dot{y}} \) (set by TSSetIJacobian()). Note that the solution state \( y \) is \([\phi \ \omega]^T\) here. For sensitivity analysis, we need to provide a routine to compute \( f_p = [0 \ 1]^T \) using TSSetRHSJacobianP(), and three routines corresponding to the integrand \( r = c(\max(0, \phi - \phi_S))^2 \), \( r_p = [0 \ 0]^T \) and \( r_y = [2c(\max(0, \phi - \phi_S)) \ 0]^T \) using TSSetCostIntegrand().

In the adjoint run, \( \lambda \) and \( \mu \) are initialized as \([0 \ 0]^T \) and \([-1 \ -1]^T \) at the final time \( t_F \). After TSAdjointSolve(), the sensitivity of the cost function w.r.t. initial conditions is given by the sensitivity variable \( \lambda \) (at time \( t_0 \)) directly. And the sensitivity of the cost function w.r.t. the parameter \( p_m \) can be computed (by users) as

\[ \frac{d\Psi}{dp_m} = \mu(t_0) + \lambda(t_0) \frac{d[\phi(t_0) \ \omega(t_0)]'}{dp_m}. \]

For explicit methods where one does not need to provide the Jacobian \( F_y \) for the forward solve one still does need it for the backward solve and thus must call

| TSSetRHSJacobian(TS ts,Mat Amat, Mat Pmat,PetscErrorCode (*f)(TS,PetscReal,Vec,Mat, _Mat, void*), void *fP); |

Examples include:

- discrete adjoint sensitivity using explicit and implicit time stepping methods for an ODE problem TS Tutorial ex20adj,
- an optimization problem using the discrete adjoint models of the ERK (for nonstiff ODEs) and the Theta methods (for stiff DAEs) TS Tutorial ex20opt_ic and TS Tutorial ex20opt_p,
- an ODE-constrained optimization using the discrete adjoint models of the Theta methods for cost function with an integral term TS Power Grid Tutorial ex3opt,
- discrete adjoint sensitivity using the Crank-Nicolson methods for DAEs with discontinuities TS Power Grid Stability Tutorial ex9busadj,
- a DAE-constrained optimization problem using the discrete adjoint models of the Crank-Nicolson methods for cost function with an integral term TS Power Grid Tutorial ex9busopt,
Checkpointing

The discrete adjoint model requires the states (and stage values in the context of multistage timestepping methods) to evaluate the Jacobian matrices during the adjoint (backward) run. By default, PETSc stores the whole trajectory to disk as binary files, each of which contains the information for a single time step including state, time, and stage values (optional). One can also make PETSc store the trajectory to memory with the option `-ts_trajectory_type memory`. However, there might not be sufficient memory capacity especially for large-scale problems and long-time integration.

A so-called checkpointing scheme is needed to solve this problem. The scheme stores checkpoints at selective time steps and recomputes the missing information. The revolve library is used by PETSc TSTrajectory to generate an optimal checkpointing schedule that minimizes the recomputations given a limited number of available checkpoints. One can specify the number of available checkpoints with the option `-ts_trajectory_max_cps_ram [maximum number of checkpoints in RAM]`. Note that one checkpoint corresponds to one time step.

The revolve library also provides an optimal multistage checkpointing scheme that uses both RAM and disk for storage. This scheme is automatically chosen if one uses both the option `-ts_trajectory_max_cps_ram [maximum number of checkpoints in RAM]` and the option `-ts_trajectory_max_cps_disk [maximum number of checkpoints on disk]`.

Some other useful options are listed below.

- `-ts_trajectory_view` prints the total number of recomputations,
- `-ts_monitor` and `-ts_adjoint_monitor` allow users to monitor the progress of the adjoint workflow,
- `-ts_trajectory_type visualization` may be used to save the whole trajectory for visualization. It stores the solution and the time, but no stage values. The binary files generated can be read into MATLAB via the script `$PETSC_DIR/share/petsc/matlab/PetscReadBinaryTrajectory.m`.

2.5.13 Using Sundials from PETSc

Sundials is a parallel ODE solver developed by Hindmarsh et al. at LLNL. The TS library provides an interface to use the CVODE component of Sundials directly from PETSc. (To configure PETSc to use Sundials, see the installation guide, installation/index.htm.)

To use the Sundials integrators, call

```c
TSSetType(TS ts, TSType TSSUNDIALS);
```

or use the command line option `-ts_type sundials`.

Sundials’ CVODE solver comes with two main integrator families, Adams and BDF (backward differentiation formula). One can select these with

```c
TSSundialsSetType(TS ts, TSSundialsLmmType [SUNDIALS_ADAMS, SUNDIALS_BDF]);
```

or the command line option `-ts_sundials_type <adams,bdf>`. BDF is the default.

Sundials does not use the SNES library within PETSc for its nonlinear solvers, so one cannot change the nonlinear solver options via SNES. Rather, Sundials uses the preconditioners within the PC package of PETSc, which can be accessed via

```c
TSSundialsGetPC(TS ts, PC *pc);
```
The user can then directly set preconditioner options; alternatively, the usual runtime options can be employed via 
\texttt{--pc xxx}.

Finally, one can set the Sundials tolerances via

\begin{verbatim}
TSSundialsSetTolerance(TS ts, double abs, double rel);
\end{verbatim}

where \texttt{abs} denotes the absolute tolerance and \texttt{rel} the relative tolerance.

Other PETSc-Sundials options include

\begin{verbatim}
TSSundialsSetGramSchmidtType(TS ts, TSSundialsGramSchmidtType type);
\end{verbatim}

where \texttt{type} is either \texttt{SUNDIALS_MODIFIED_GS} or \texttt{SUNDIALS_UNMODIFIED_GS}. This may be set via the options data base with 
\texttt{-ts_sundials_gramschmidt_type <modified,unmodified>}

The routine

\begin{verbatim}
TSSundialsSetMaxl(TS ts, PetscInt restart);
\end{verbatim}

sets the number of vectors in the Krylov subspace used by GMRES. This may be set in the options database with 
\texttt{-ts_sundials_maxl maxl}.

\subsection*{2.5.14 Using TChem from PETSc}

TChem\textsuperscript{6} is a package originally developed at Sandia National Laboratory that can read in CHEMKIN\textsuperscript{7} data files and compute the right hand side function and its Jacobian for a reaction ODE system. To utilize PETSc’s ODE solvers for these systems, first install PETSc with the additional \texttt{configure} option 
\texttt{--download-tchem}. We currently provide two examples of its use; one for single cell reaction and one for an “artificial” one dimensional problem with periodic boundary conditions and diffusion of all species. The self-explanatory examples are the \texttt{The TS tutorial extchem} and \texttt{The TS tutorial extchemfield}.

\section*{2.6 Solving Steady-State Problems with Pseudo-Timestepping}

\textbf{Simple Example:} \texttt{TS} provides a general code for performing pseudo timestepping with a variable timestep at each physical node point. For example, instead of directly attacking the steady-state problem

\[ G(u) = 0, \]

we can use pseudo-transient continuation by solving

\[ u_t = G(u). \]

Using time differencing

\[ u_t \equiv \frac{u^{n+1} - u^n}{dt^n} \]

with the backward Euler method, we obtain nonlinear equations at a series of pseudo-timesteps

\[ \frac{1}{dt^n} B(u^{n+1} - u^n) = G(u^{n+1}). \]

\begin{footnotesize}
\begin{enumerate}
\item bitbucket.org/jedbrown/tchem
\item en.wikipedia.org/wiki/CHEMKIN
\end{enumerate}
\end{footnotesize}
For this problem the user must provide $G(u)$, the time steps $dt^n$ and the left-hand-side matrix $B$ (or optionally, if the timestep is position independent and $B$ is the identity matrix, a scalar timestep), as well as optionally the Jacobian of $G(u)$.

More generally, this can be applied to implicit ODE and DAE for which the transient form is

$$F(u, \dot{u}) = 0.$$ 

For solving steady-state problems with pseudo-timestepping one proceeds as follows.

- Provide the function $G(u)$ with the routine

  ```c
  TSSetRHSFunction(TS ts, Vec r, PetscErrorCode (*f)(TS, PetscReal, Vec, Vec, void*), void*, *fP);
  ```

  The arguments to the function $f()$ are the timestep context, the current time, the input for the function, the output for the function and the (optional) user-provided context variable $fP$.

- Provide the (approximate) Jacobian matrix of $G(u)$ and a function to compute it at each Newton iteration. This is done with the command

  ```c
  TSSetRHSJacobian(TS ts, Mat Amat, Mat Pmat, PetscErrorCode (*f)(TS, PetscReal, Vec, Mat, Mat, void*), void* fP);
  ```

  The arguments for the function $f()$ are the timestep context, the current time, the location where the Jacobian is to be computed, the (approximate) Jacobian matrix, an alternative approximate Jacobian matrix used to construct the preconditioner, and the optional user-provided context, passed in as $fP$. The user must provide the Jacobian as a matrix; thus, if using a matrix-free approach, one must create a MATSHELL matrix.

In addition, the user must provide a routine that computes the pseudo-timestep. This is slightly different depending on if one is using a constant timestep over the entire grid, or it varies with location.

- For location-independent pseudo-timestepping, one uses the routine

  ```c
  TSPseudoSetTimeStep(TS ts, PetscInt(*dt)(TS, PetscReal*, void*), void* dtctx);
  ```

  The function $dt$ is a user-provided function that computes the next pseudo-timestep. As a default one can use `TSPseudoTimeStepDefault(TS,PetscReal*,void*)` for $dt$. This routine updates the pseudo-timestep with one of two strategies: the default

  $$dt^n = dt_{\text{increment}} * dt^{n-1} * \frac{||F(u^{n-1})||}{||F(u^n)||}$$

  or, the alternative,

  $$dt^n = dt_{\text{increment}} * dt^0 * \frac{||F(u^0)||}{||F(u^n)||}$$

  which can be set with the call

  ```c
  TSPseudoIncrementDtFromInitialDt(TS ts);
  ```

  or the option `-ts_pseudo_increment_dt_from_initial_dt`. The value $dt_{\text{increment}}$ is by default 1.1, but can be reset with the call

  ```c
  TSPseudoSetTimeStepIncrement(TS ts, PetscReal inc);
  ```

  or the option `-ts_pseudo_increment <inc>`.

- For location-dependent pseudo-timestepping, the interface function has not yet been created.
2.7 TAO: Optimization Solvers

The Toolkit for Advanced Optimization (TAO) focuses on algorithms for the solution of large-scale optimization problems on high-performance architectures. Methods are available for

- Nonlinear Least-Squares
- Quadratic Solvers
- Unconstrained Minimization
- Bound-Constrained Optimization
- Generally Constrained Solvers
- Complementarity
- PDE-constrained Optimization

2.7.1 Getting Started: A Simple TAO Example

To help the user start using TAO immediately, we introduce here a simple uniprocessor example. Please read TAO Algorithms for a more in-depth discussion on using the TAO solvers. The code presented below minimizes the extended Rosenbrock function \( f: \mathbb{R}^n \to \mathbb{R} \) defined by

\[
  f(x) = \sum_{i=0}^{n-1} \left( \alpha(x_{2i+1} - x_{2i}^2)^2 + (1 - x_{2i})^2 \right),
\]

where \( n = 2m \) is the number of variables. Note that while we use the C language to introduce the TAO software, the package is fully usable from C++ and Fortran. PETSc for Fortran Users discusses additional issues concerning Fortran usage.

The code in the example contains many of the components needed to write most TAO programs and thus is illustrative of the features present in complex optimization problems. Note that for display purposes we have omitted some nonessential lines of code as well as the (essential) code required for the routine FormFunctionGradient, which evaluates the function and gradient, and the code for FormHessian, which evaluates the Hessian matrix for Rosenbrock’s function. The complete code is available in $TAO_DIR/src/unconstrained/tutorials/rosenbrock1.c. The following sections annotate the lines of code in the example.

Listing: src/tao/unconstrained/tutorials/rosenbrock1.c

```c
#include <petsctao.h>

typedef struct {
  PetscInt n; /* dimension */
  PetscReal alpha; /* condition parameter */
  PetscBool chained;
} AppCtx;

PetscErrorCode FormFunctionGradient(Tao, Vec, PetscReal *, Vec, void *);
PetscErrorCode FormHessian(Tao, Vec, Mat, Mat, void *);

int main(int argc, char **argv) {
  PetscReal zero = 0.0;
```

(continues on next page)
Vec x; /* solution vector */
Mat H;
Tao tao; /* Tao solver context */
PetscBool flg, test_lmvm = PETSC_FALSE;
PetscMPIInt size; /* number of processes running */
AppCtx user; /* user-defined application context */
KSP ksp;
PC pc;
Mat M;
Vec in, out, out2;
PetscReal mult_solve_dist;

/* Initialize TAO and PETSc */
PetscFunctionBeginUser;
PetscCall(PetscInitialize(&argc, &argv, (char *)0, help));
PetscCall(MPI_Comm_size(PETSC_COMM_WORLD, &size));
PetscCheck(size == 1, PETSC_COMM_WORLD, PETSC_ERR_WRONG_MPI_SIZE, "Incorrect number of processors");

/* Initialize problem parameters */
user.n = 2;
user.alpha = 99.0;
user.chained = PETSC_FALSE;
/* Check for command line arguments to override defaults */
PetscCall(PetscOptionsGetInt(NULL, NULL, "-n", &user.n, &flg));
PetscCall(PetscOptionsGetReal(NULL, NULL, "-alpha", &user.alpha, &flg));
PetscCall(PetscOptionsGetBool(NULL, NULL, "-chained", &user.chained, &flg));
PetscCall(PetscOptionsGetBool(NULL, NULL, "-test_lmvm", &test_lmvm, &flg));

/* Allocate vectors for the solution and gradient */
PetscCall(VecCreateSeq(PETSC_COMM_SELF, user.n, &x));
PetscCall(MatCreateSeqBAIJ(PETSC_COMM_SELF, 2, user.n, user.n, 1, NULL, &H));

/* The TAO code begins here */

/* Create TAO solver with desired solution method */
PetscCall(TaoCreate(PETSC_COMM_SELF, &tao));
PetscCall(TaoSetType(tao, TAOLMVM));

/* Set solution vec and an initial guess */
PetscCall(VecSet(x, zero));
PetscCall(TaoSetSolution(tao, x));

/* Set routines for function, gradient, hessian evaluation */
PetscCall(TaoSetObjectiveAndGradient(tao, NULL, FormFunctionGradient, &user));
PetscCall(TaoSetHessian(tao, H, H, FormHessian, &user));

/* Test the LMVM matrix */
if (test_lmvm) PetscCall(PetscOptionsSetValue(NULL, "-tao_type", "bqnktr"));

/* Check for TAO command line options */
PetscCall(TaoSetFromOptions(tao));

/* SOLVE THE APPLICATION */
PetscCall(TaoSolve(tao));

(continues on next page)
/* Test the LMVM matrix */
if (test_lmvm) {
    PetscCall(TaoGetKSP(tao, &ksp));
    PetscCall(KSPGetPC(ksp, &pc));
    PetscCall(PCLMVMGetMatLMVM(pc, &M));
    PetscCall(VecDuplicate(x, &in));
    PetscCall(VecDuplicate(x, &out));
    PetscCall(VecDuplicate(x, &out2));
    PetscCall(VecSet(in, 1.0));
    PetscCall(MatMult(M, in, out));
    PetscCall(MatSolve(M, out, out2));
    PetscCall(VecAXPY(out2, -1.0, in));
    PetscCall(VecNorm(out2, NORM_2, &mult_solve_dist));
    if (mult_solve_dist < 1.e-11) {
        PetscCall(PetscPrintf(PetscObjectComm((PetscObject)tao),
            "error between LMVM and MatMult and MatSolve: < 1.e-11\n"));
    } else if (mult_solve_dist < 1.e-6) {
        PetscCall(PetscPrintf(PetscObjectComm((PetscObject)tao),
            "error between LMVM and MatMult and MatSolve: < 1.e-6\n"));
    } else {
        PetscCall(PetscPrintf(PetscObjectComm((PetscObject)tao),
            "error between LMVM and MatMult and MatSolve: %e\n", (double)mult_solve_dist));
    }
    PetscCall(VecDestroy(&in));
    PetscCall(VecDestroy(&out));
    PetscCall(VecDestroy(&out2));
}
PetscCall(TaoDestroy(&tao));
PetscCall(VecDestroy(&x));
PetscCall(MatDestroy(&H));
PetscCall(PetscFinalize());
return ierr;

2.7.2 TAO Workflow

Many TAO applications will follow an ordered set of procedures for solving an optimization problem: The user creates a Tao context and selects a default algorithm. Call-back routines as well as vector (Vec) and matrix (Mat) data structures are then set. These call-back routines will be used for evaluating the objective function, gradient, and perhaps the Hessian matrix. The user then invokes TAO to solve the optimization problem and finally destroys the Tao context. A list of the necessary functions for performing these steps using TAO is shown below.

TaoCreate(MPI_Comm comm, Tao *tao);
TaoSetType(Tao tao, TaoType type);
TaoSetSolution(Tao tao, Vec x);
TaoSetObjectiveAndGradient(Tao tao, Vec g, PetscErrorCode (*FormFGradient)(Tao, Vec, void *user), PetscReal *, void *user);
TaoSetHessian(Tao tao, Mat H, Mat Hpre, PetscErrorCode (*FormHessian)(Tao, Vec, Mat, void *user), void *user);
TaoSolve(Tao tao);
TaoDestroy(Tao tao);
Note that the solver algorithm selected through the function \texttt{TaoSetType()} can be overridden at runtime by using an options database. Through this database, the user not only can select a minimization method (e.g., limited-memory variable metric, conjugate gradient, Newton with line search or trust region) but also can prescribe the convergence tolerance, set various monitoring routines, set iterative methods and preconditions for solving the linear systems, and so forth. See \textit{TAO Algorithms} for more information on the solver methods available in TAO.

\subsection*{Header File}

TAO applications written in C/C++ should have the statement
\verbatim{include <petsctao.h>}
in each file that uses a routine in the TAO libraries.

\subsection*{Creation and Destruction}

A TAO solver can be created by calling the
\verbatim{TaoCreate(MPI_Comm, Tao*);}
routine. Much like creating PETSc vector and matrix objects, the first argument is an MPI \texttt{communicator}. An MPI\footnote{For more on MPI and PETSc, see \textit{Running PETSc Programs}.} communicator indicates a collection of processors that will be used to evaluate the objective function, compute constraints, and provide derivative information. When only one processor is being used, the communicator \texttt{PETSC_COMM_SELF} can be used with no understanding of MPI. Even parallel users need to be familiar with only the basic concepts of message passing and distributed-memory computing. Most applications running TAO in parallel environments can employ the communicator \texttt{PETSC_COMM_WORLD} to indicate all processes known to PETSc in a given run.

The routine
\verbatim{TaoSetType(Tao, TaoType);} can be used to set the algorithm TAO uses to solve the application. The various types of TAO solvers and the flags that identify them will be discussed in the following sections. The solution method should be carefully chosen depending on the problem being solved. Some solvers, for instance, are meant for problems with no constraints, whereas other solvers acknowledge constraints in the problem and handle them accordingly. The user must also be aware of the derivative information that is available. Some solvers require second-order information, while other solvers require only gradient or function information. The command line option \texttt{-tao_type} followed by a TAO method will override any method specified by the second argument. The command line option \texttt{-tao_type bqnls}, for instance, will specify the limited-memory quasi-Newton line search method for bound-constrained problems. Note that the \texttt{TaoType} variable is a string that requires quotation marks in an application program, but quotation marks are not required at the command line.

Each TAO solver that has been created should also be destroyed by using the
\verbatim{TaoDestroy(Tao tao);} command. This routine frees the internal data structures used by the solver.
Command-line Options

Additional options for the TAO solver can be set from the command line by using the `TaoSetFromOptions(Tao)` routine. This command also provides information about runtime options when the user includes the `-help` option on the command line.

In addition to common command line options shared by all TAO solvers, each TAO method also implements its own specialized options. Please refer to the documentation for individual methods for more details.

Defining Variables

In all the optimization solvers, the application must provide a `Vec` object of appropriate dimension to represent the variables. This vector will be cloned by the solvers to create additional work space within the solver. If this vector is distributed over multiple processors, it should have a parallel distribution that allows for efficient scaling, inner products, and function evaluations. This vector can be passed to the application object by using the `TaoSetSolution(Tao, Vec);` routine. When using this routine, the application should initialize the vector with an approximate solution of the optimization problem before calling the TAO solver. This vector will be used by the TAO solver to store the solution. Elsewhere in the application, this solution vector can be retrieved from the application object by using the `TaoGetSolution(Tao, Vec*);` routine. This routine takes the address of a `Vec` in the second argument and sets it to the solution vector used in the application.

User Defined Call-back Routines

Users of TAO are required to provide routines that perform function evaluations. Depending on the solver chosen, they may also have to write routines that evaluate the gradient vector and Hessian matrix.

Application Context

Writing a TAO application may require use of an application context. An application context is a structure or object defined by an application developer, passed into a routine also written by the application developer, and used within the routine to perform its stated task.

For example, a routine that evaluates an objective function may need parameters, work vectors, and other information. This information, which may be specific to an application and necessary to evaluate the objective, can be collected in a single structure and used as one of the arguments in the routine. The address of this structure will be cast as type `(void*)` and passed to the routine in the final argument. Many examples of these structures are included in the TAO distribution.

This technique offers several advantages. In particular, it allows for a uniform interface between TAO and the applications. The fundamental information needed by TAO appears in the arguments of the routine, while data specific to an application and its implementation is confined to an opaque pointer. The routines can access information created outside the local scope without the use of global variables. The TAO solvers
and application objects will never access this structure, so the application developer has complete freedom to define it. If no such structure or needed by the application then a NULL pointer can be used.

**Objective Function and Gradient Routines**

TAO solvers that minimize an objective function require the application to evaluate the objective function. Some solvers may also require the application to evaluate derivatives of the objective function. Routines that perform these computations must be identified to the application object and must follow a strict calling sequence.

Routines should follow the form

```c
PetscErrorCode EvaluateObjective(Tao, Vec, PetscReal*, void*);
```

in order to evaluate an objective function \( f : \mathbb{R}^n \to \mathbb{R} \). The first argument is the TAO Solver object, the second argument is the \( n \)-dimensional vector that identifies where the objective should be evaluated, and the fourth argument is an application context. This routine should use the third argument to return the objective value evaluated at the point specified by the vector in the second argument.

This routine, and the application context, should be passed to the application object by using the

```c
TaoSetObjective(Tao, PetscErrorCode(*)(Tao,Vec,PetscReal*,void*), void*);
```

routine. The first argument in this routine is the TAO solver object, the second argument is a function pointer to the routine that evaluates the objective, and the third argument is the pointer to an appropriate application context. Although the final argument may point to anything, it must be cast as a \((\text{void}*)\) type. This pointer will be passed back to the developer in the fourth argument of the routine that evaluates the objective. In this routine, the pointer can be cast back to the appropriate type. Examples of these structures and their usage are provided in the distribution.

Many TAO solvers also require gradient information from the application. The gradient of the objective function is specified in a similar manner. Routines that evaluate the gradient should have the calling sequence

```c
PetscErrorCode EvaluateGradient(Tao, Vec, Vec, void*);
```

where the first argument is the TAO solver object, the second argument is the variable vector, the third argument is the gradient vector, and the fourth argument is the user-defined application context. Only the third argument in this routine is different from the arguments in the routine for evaluating the objective function. The numbers in the gradient vector have no meaning when passed into this routine, but they should represent the gradient of the objective at the specified point at the end of the routine. This routine, and the user-defined pointer, can be passed to the application object by using the

```c
TaoSetGradient(Tao, Vec, PetscErrorCode(*)(Tao, Vec, Vec, void*), void*);
```

routine. In this routine, the first argument is the Tao object, the second argument is the optional vector to hold the computed gradient, the third argument is the function pointer, and the fourth object is the application context, cast to \((\text{void}*)\).

Instead of evaluating the objective and its gradient in separate routines, TAO also allows the user to evaluate the function and the gradient in the same routine. In fact, some solvers are more efficient when both function and gradient information can be computed in the same routine. These routines should follow the form

```c
PetscErrorCode EvaluateFunctionAndGradient(Tao, Vec, PetscReal*, Vec, void*);
```
where the first argument is the TAO solver and the second argument points to the input vector for use in evaluating the function and gradient. The third argument should return the function value, while the fourth argument should return the gradient vector. The fifth argument is a pointer to a user-defined context. This context and the name of the routine should be set with the call

```
TaoSetObjectiveAndGradient(Tao, Vec PetscErrorCode (*)(Tao,Vec,PetscReal*,Vec,void*), ω →void*);
```

where the arguments are the TAO application, the optional vector to be used to hold the computed gradient, a function pointer, and a pointer to a user-defined context.

The TAO example problems demonstrate the use of these application contexts as well as specific instances of function, gradient, and Hessian evaluation routines. All these routines should return the integer 0 after successful completion and a nonzero integer if the function is undefined at that point or an error occurred.

**Hessian Evaluation**

Some optimization routines also require a Hessian matrix from the user. The routine that evaluates the Hessian should have the form

```
PetscErrorCode EvaluateHessian(Tao, Vec, Mat, Mat, ω void*);
```

where the first argument of this routine is a TAO solver object. The second argument is the point at which the Hessian should be evaluated. The third argument is the Hessian matrix, and the sixth argument is a user-defined context. Since the Hessian matrix is usually used in solving a system of linear equations, a preconditioner for the matrix is often needed. The fourth argument is the matrix that will be used for preconditioning the linear system; in most cases, this matrix will be the same as the Hessian matrix. The fifth argument is the flag used to set the Hessian matrix and linear solver in the routine `KSPSetOperators()`.

One can set the Hessian evaluation routine by calling the

```
TaoSetHessian(Tao, Mat, Mat, PetscErrorCode (*)(Tao,Vec,Mat,Mat,void*), void*);
```

routine. The first argument is the TAO Solver object. The second and third arguments are, respectively, the Mat object where the Hessian will be stored and the Mat object that will be used for the preconditioning (they may be the same). The fourth argument is the function that evaluates the Hessian, and the fifth argument is a pointer to a user-defined context, cast to `(void*)`.

**Finite Differences**

Finite-difference approximations can be used to compute the gradient and the Hessian of an objective function. These approximations will slow the solve considerably and are recommended primarily for checking the accuracy of hand-coded gradients and Hessians. These routines are

```
TaoDefaultComputeGradient(Tao, Vec, Vec, void*);
```

and

```
TaoDefaultComputeHessian(Tao, Vec, Mat*, Mat*, void*);
```

respectively. They can be set by using `TaoSetGradient()` and `TaoSetHessian()` or through the options database with the options `-tao_fdgrad` and `-tao_fd`, respectively.
The efficiency of the finite-difference Hessian can be improved if the coloring of the matrix is known. If the application programmer creates a PETSc `MatFDColoring` object, it can be applied to the finite-difference approximation by setting the Hessian evaluation routine to

```
TaoDefaultComputeHessianColor(Tao, Vec, Mat*, Mat*, void*);
```

and using the `MatFDColoring` object as the last (`void *`) argument to `TaoSetHessian()`.

One also can use finite-difference approximations to directly check the correctness of the gradient and/or Hessian evaluation routines. This process can be initiated from the command line by using the special TAO solver `tao_fd_test` together with the option `-tao_test_gradient` or `-tao_test_hessian`.

### Matrix-Free Methods

TAO fully supports matrix-free methods. The matrices specified in the Hessian evaluation routine need not be conventional matrices; instead, they can point to the data required to implement a particular matrix-free method. The matrix-free variant is allowed only when the linear systems are solved by an iterative method in combination with no preconditioning (`PCNONE` or `-pc_type none`), a user-provided preconditioner matrix, or a user-provided preconditioner shell (`PCSHELL`). In other words, matrix-free methods cannot be used if a direct solver is to be employed. Details about using matrix-free methods are provided in the *User-Guide.*

![Fig. 2.5: Tao use of PETSc and callbacks](image)
Constraints

Some optimization problems also impose constraints on the variables or intermediate application states. The user defines these constraints through the appropriate TAO interface functions and call-back routines where necessary.

Variable Bounds

The simplest type of constraint on an optimization problem puts lower or upper bounds on the variables. Vectors that represent lower and upper bounds for each variable can be set with the

\[ \text{TaoSetVariableBounds(Tao, Vec, Vec);} \]

command. The first vector and second vector should contain the lower and upper bounds, respectively. When no upper or lower bound exists for a variable, the bound may be set to \text{PETSC_INFINITY} or \text{PETSC_NINFINITY}. After the two bound vectors have been set, they may be accessed with the command \text{TaoGetVariableBounds()}. Since not all solvers recognize the presence of bound constraints on variables, the user must be careful to select a solver that acknowledges these bounds.

General Constraints

Some TAO algorithms also support general constraints as a linear or nonlinear function of the optimization variables. These constraints can be imposed either as equalities or inequalities. TAO currently does not make any distinctions between linear and nonlinear constraints, and implements them through the same software interfaces.

In the equality constrained case, TAO assumes that the constraints are formulated as \( c_e(x) = 0 \) and requires the user to implement a call-back routine for evaluating \( c_e(x) \) at a given vector of optimization variables,

\[ \text{PetscErrorCode EvaluateEqualityConstraints(Tao, Vec, Vec, void*);} \]

As in the previous call-back routines, the first argument is the TAO solver object. The second and third arguments are the vector of optimization variables (input) and vector of equality constraints (output), respectively. The final argument is a pointer to the user-defined application context, cast into \text{(void*)}. Generally constrained TAO algorithms also require a second user call-back function to compute the constraint Jacobian matrix \( \nabla_x c_e(x) \),

\[ \text{PetscErrorCode EvaluateEqualityJacobian(Tao, Vec, Mat, Mat, void*);} \]

where the first and last arguments are the TAO solver object and the application context pointer as before. The second argument is the vector of optimization variables at which the computation takes place. The third and fourth arguments are the constraint Jacobian and its pseudo-inverse (optional), respectively. The pseudoinverse is optional, and if not available, the user can simply set it to the constraint Jacobian itself. These call-back functions are then given to the TAO solver using the interface functions

\[ \text{TaoSetEqualityConstraintsRoutine(Tao, Vec, PetscErrorCode (*)(Tao,Vec,Vec,void*), \_void*);} \]

and

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Inequality constraints are assumed to be formulated as \( c_i(x) \leq 0 \) and follow the same workflow as equality constraints using the \texttt{TaoSetInequalityConstraintsRoutine()} and \texttt{TaoSetJacobianInequalityRoutine()} interfaces.

Some TAO algorithms may adopt an alternative double-sided \( c_l \leq c_i(x) \leq c_u \) formulation and require the lower and upper bounds \( c_l \) and \( c_u \) to be set using the \texttt{TaoSetInequalityBounds(Tao,Vec,Vec)} interface. Please refer to the documentation for each TAO algorithm for further details.

### Solving

Once the application and solver have been set up, the solver can be

\[
\text{TaoSolve(Tao);} 
\]

routine. We discuss several universal options below.

### Convergence

Although TAO and its solvers set default parameters that are useful for many problems, the user may need to modify these parameters in order to change the behavior and convergence of various algorithms.

One convergence criterion for most algorithms concerns the number of digits of accuracy needed in the solution. In particular, the convergence test employed by TAO attempts to stop when the error in the constraints is less than \( \epsilon_{crtol} \) and either

\[
\begin{align*}
||g(X)|| & \leq \epsilon_{gatol}, \\
||g(X)||/|f(X)| & \leq \epsilon_{grtol}, \text{ or} \\
||g(X)||/|g(X_0)| & \leq \epsilon_{gttol},
\end{align*}
\]

where \( X \) is the current approximation to the true solution \( X^* \) and \( X_0 \) is the initial guess. \( X^* \) is unknown, so TAO estimates \( f(X) - f(X^*) \) with either the square of the norm of the gradient or the duality gap. A relative tolerance of \( \epsilon_{frtol} = 0.01 \) indicates that two significant digits are desired in the objective function. Each solver sets its own convergence tolerances, but they can be changed by using the routine \texttt{TaoSetTolerances()}.

Another set of convergence tolerances terminates the solver when the norm of the gradient function (or Lagrangian function for bound-constrained problems) is sufficiently close to zero.

Other stopping criteria include a minimum trust-region radius or a maximum number of iterations. These parameters can be set with the routines \texttt{TaoSetTrustRegionTolerance()} and \texttt{TaoSetMaximumIterations()} Similarly, a maximum number of function evaluations can be set with the command \texttt{TaoSetMaximumFunctionEvaluations()} (\texttt{-tao_max_it}) and \texttt{-tao_max_funcs}.

### Viewing Status

To see parameters and performance statistics for the solver, the routine

\[
\text{TaoView(Tao tao)} 
\]

can be used. This routine will display to standard output the number of function evaluations need by the solver and other information specific to the solver. This same output can be produced by using the command line option \texttt{-tao_view}.
The progress of the optimization solver can be monitored with the runtime option -tao_monitor. Although monitoring routines can be customized, the default monitoring routine will print out several relevant statistics to the screen.

The user also has access to information about the current solution. The current iteration number, objective function value, gradient norm, infeasibility norm, and step length can be retrieved with the following command.

```c
TaoGetSolutionStatus(Tao tao, PetscInt* iterate, PetscReal* f,
                     PetscReal* gnorm, PetscReal* cnorm, PetscReal* xdiff,
                     TaoConvergedReason* reason)
```

The last argument returns a code that indicates the reason that the solver terminated. Positive numbers indicate that a solution has been found, while negative numbers indicate a failure. A list of reasons can be found in the manual page for TaoGetConvergedReason().

**Obtaining a Solution**

After exiting the TaoSolve() function, the solution and the gradient can be recovered with the following routines.

```c
TaoGetSolution(Tao, Vec*);
TaoGetGradient(Tao, Vec*, NULL, NULL);
```

Note that the Vec returned by TaoGetSolution() will be the same vector passed to TaoSetSolution(). This information can be obtained during user-defined routines such as a function evaluation and customized monitoring routine or after the solver has terminated.

**Special Problem structures**

Certain special classes of problems solved with TAO utilize specialized code interfaces that are described below per problem type.

**PDE-constrained Optimization**

TAO solves PDE-constrained optimization problems of the form

\[
\min_{u,v} \quad f(u,v) \\
\text{subject to} \quad g(u,v) = 0,
\]

where the state variable \( u \) is the solution to the discretized partial differential equation defined by \( g \) and parametrized by the design variable \( v \), and \( f \) is an objective function. The Lagrange multipliers on the constraint are denoted by \( y \). This method is set by using the linearly constrained augmented Lagrangian TAO solver tao_lcl.

We make two main assumptions when solving these problems: the objective function and PDE constraints have been discretized so that we can treat the optimization problem as finite dimensional and \( \nabla_u g(u,v) \) is invertible for all \( u \) and \( v \).

Unlike other TAO solvers where the solution vector contains only the optimization variables, PDE-constrained problems solved with tao_lcl combine the design and state variables together in a monolithic solution vector \( x^T = [u^T, v^T] \). Consequently, the user must provide index sets to separate the two,
where the first IS is a PETSc IndexSet containing the indices of the state variables and the second IS the design variables.

PDE constraints have the general form $g(x) = 0$, where $c: \mathbb{R}^n \rightarrow \mathbb{R}^m$. These constraints should be specified in a routine, written by the user, that evaluates $g(x)$. The routine that evaluates the constraint equations should have the form

$$\text{PetscErrorCode \ EvaluateConstraints(Tao, Vec, Vec, void*)};$$

The first argument of this routine is a TAO solver object. The second argument is the variable vector at which the constraint function should be evaluated. The third argument is the vector of function values $g(x)$, and the fourth argument is a pointer to a user-defined context. This routine and the user-defined context should be set in the TAO solver with the

$$\text{TaoSetConstraintsRoutine(Tao, Vec, PetscErrorCode (*)(Tao, Vec, Vec, void*), void*)};$$

command. In this function, the first argument is the TAO solver object, the second argument a vector in which to store the constraints, the third argument is a function point to the routine for evaluating the constraints, and the fourth argument is a pointer to a user-defined context.

The Jacobian of $g(x)$ is the matrix in $\mathbb{R}^{m \times n}$ such that each column contains the partial derivatives of $g(x)$ with respect to one variable. The evaluation of the Jacobian of $g$ should be performed by calling the

$$\text{PetscErrorCode \ JacobianState(Tao, Vec, Mat, Mat, Mat, void*)};$$
$$\text{PetscErrorCode \ JacobianDesign(Tao, Vec, Mat*, void*)};$$

routines. In these functions, The first argument is the TAO solver object. The second argument is the variable vector at which to evaluate the Jacobian matrix, the third argument is the Jacobian matrix, and the last argument is a pointer to a user-defined context. The fourth and fifth arguments of the Jacobian evaluation with respect to the state variables are for providing PETSc matrix objects for the preconditioner and for applying the inverse of the state Jacobian, respectively. This inverse matrix may be `PETSC_NULL`, in which case TAO will use a PETSc Krylov subspace solver to solve the state system. These evaluation routines should be registered with TAO by using the

$$\text{TaoSetJacobianStateRoutine(Tao, Mat, Mat, Mat, PetscErrorCode (*)(Tao, Vec, Mat, Mat, void*), void*)};$$
$$\text{TaoSetJacobianDesignRoutine(Tao, Mat, PetscErrorCode (*)(Tao, Vec, Mat*, void*), void*)};$$

routines. The first argument is the TAO solver object, and the second argument is the matrix in which the Jacobian information can be stored. For the state Jacobian, the third argument is the matrix that will be used for preconditioning, and the fourth argument is an optional matrix for the inverse of the state Jacobian. One can use `PETSC_NULL` for this inverse argument and let PETSc apply the inverse using a KSP method, but faster results may be obtained by manipulating the structure of the Jacobian and providing an inverse. The fifth argument is the function pointer, and the sixth argument is an optional user-defined context. Since no solve is performed with the design Jacobian, there is no need to provide preconditioner or inverse matrices.
Nonlinear Least Squares

For nonlinear least squares applications, we are solving the optimization problem

$$
\min_x \frac{1}{2} ||r(x)||_2^2.
$$

For these problems, the objective function value should be computed as a vector of residuals, $r(x)$, computed with a function of the form

```c
PetscErrorCode EvaluateResidual(Tao, Vec, Vec, void*);
```

and set with the

```c
TaoSetResidualRoutine(Tao, PetscErrorCode (*)(Tao,Vec,Vec,void*), void*);
```

routine. If required by the algorithm, the Jacobian of the residual, $J = \partial r(x)/\partial x$, should be computed with a function of the form

```c
PetscErrorCode EvaluateJacobian(Tao, Vec, Mat, void*);
```

and set with the

```c
TaoSetJacobianResidualRoutine(Tao, PetscErrorCode (*)(Tao,Vec,Mat,void*), void *);
```

Complementarity

Complementarity applications have equality constraints in the form of nonlinear equations $C(X) = 0$, where $C : \mathbb{R}^n \rightarrow \mathbb{R}^m$. These constraints should be specified in a routine written by the user with the form

```c
PetscErrorCode EqualityConstraints(Tao, Vec, Vec, void*);
```

that evaluates $C(X)$. The first argument of this routine is a TAO Solver object. The second argument is the variable vector $X$ at which the constraint function should be evaluated. The third argument is the output vector of function values $C(X)$, and the fourth argument is a pointer to a user-defined context.

This routine and the user-defined context must be registered with TAO by using the

```c
TaoSetConstraintRoutine(Tao, Vec, PetscErrorCode (*)(Tao,Vec,Vec,void*), void *);
```

command. In this command, the first argument is TAO Solver object, the second argument is vector in which to store the function values, the third argument is the user-defined routine that evaluates $C(X)$, and the fourth argument is a pointer to a user-defined context that will be passed back to the user.

The Jacobian of the function is the matrix in $\mathbb{R}^{m \times n}$ such that each column contains the partial derivatives of $f$ with respect to one variable. The evaluation of the Jacobian of $C$ should be performed in a routine of the form

```c
PetscErrorCode EvaluateJacobian(Tao, Vec, Mat, void*);
```

In this function, the first argument is the TAO Solver object and the second argument is the variable vector at which to evaluate the Jacobian matrix. The third argument is the Jacobian matrix, and the sixth argument is a pointer to a user-defined context. Since the Jacobian matrix may be used in solving a system of linear equations, a preconditioner for the matrix may be needed. The fourth argument is the matrix that will
be used for preconditioning the linear system; in most cases, this matrix will be the same as the Hessian matrix. The fifth argument is the flag used to set the Jacobian matrix and linear solver in the routine `KSPSetOperators()`.

This routine should be specified to TAO by using the

```c
TaoSetJacobianRoutine(Tao, Mat, Mat, PetscErrorCode (*)(Tao,Vec,Mat,Mat,void*), _void*);
```

command. The first argument is the TAO Solver object; the second and third arguments are the Mat objects in which the Jacobian will be stored and the Mat object that will be used for the preconditioning (they may be the same), respectively. The fourth argument is the function pointer; and the fifth argument is an optional user-defined context. The Jacobian matrix should be created in a way such that the product of it and the variable vector can be stored in the constraint vector.

### 2.7.3 TAO Algorithms

TAO includes a variety of optimization algorithms for several classes of problems (unconstrained, bound-constrained, and PDE-constrained minimization, nonlinear least-squares, and complementarity). The TAO algorithms for solving these problems are detailed in this section, a particular algorithm can chosen by using the `TaoSetType()` function or using the command line arguments `-tao_type <name>`. For those interested in extending these algorithms or using new ones, please see *Adding a Solver* for more information.

#### Unconstrained Minimization

Unconstrained minimization is used to minimize a function of many variables without any constraints on the variables, such as bounds. The methods available in TAO for solving these problems can be classified according to the amount of derivative information required:

1. Function evaluation only – Nelder-Mead method (**tao_nm**)
2. Function and gradient evaluations – limited-memory, variable-metric method (**tao_lmmv**), and nonlinear conjugate gradient method (**tao_cg**)
3. Function, gradient, and Hessian evaluations – Newton-Krylov methods: Newton line search (**tao_nls**), Newton trust-region (**tao_ntr**), and Newton trust-region line-search (**tao_ntl**)

The best method to use depends on the particular problem being solved and the accuracy required in the solution. If a Hessian evaluation routine is available, then the Newton line search and Newton trust-region methods will likely perform best. When a Hessian evaluation routine is not available, then the limited-memory, variable-metric method is likely to perform best. The Nelder-Mead method should be used only as a last resort when no gradient information is available.

Each solver has a set of options associated with it that can be set with command line arguments. These algorithms and the associated options are briefly discussed in this section.
Newton-Krylov Methods

TAO features three Newton-Krylov algorithms, separated by their globalization methods for unconstrained optimization: line search (NLS), trust region (NTR), and trust region with a line search (NTL). They are available via the TAO solvers TAO_NLS, TAO_NTR and TAO_NTL, respectively, or the -tao_type=nls/ntr/ntl flag.

Newton Line Search Method (NLS)

The Newton line search method solves the symmetric system of equations

\[ H_k d_k = -g_k \]

to obtain a step \( d_k \), where \( H_k \) is the Hessian of the objective function at \( x_k \) and \( g_k \) is the gradient of the objective function at \( x_k \). For problems where the Hessian matrix is indefinite, the perturbed system of equations

\[(H_k + \rho_k I)d_k = -g_k\]

is solved to obtain the direction, where \( \rho_k \) is a positive constant. If the direction computed is not a descent direction, the (scaled) steepest descent direction is used instead. Having obtained the direction, a Moré-Thuente line search is applied to obtain a step length, \( \tau_k \), that approximately solves the one-dimensional optimization problem

\[
\min_{\tau} f(x_k + \tau d_k).
\]

The Newton line search method can be selected by using the TAO solver tao_nls. The options available for this solver are listed in Table 2.17. For the best efficiency, function and gradient evaluations should be performed simultaneously when using this algorithm.

<table>
<thead>
<tr>
<th>Name -tao_nls_</th>
<th>Value</th>
<th>Default</th>
<th>Description</th>
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<td></td>
</tr>
<tr>
<td>pc_type</td>
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<td>PCType for linear system</td>
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</tr>
<tr>
<td>imin</td>
<td>real</td>
<td>(10^{-4})</td>
<td>Minimum initial perturbation value</td>
</tr>
<tr>
<td>imax</td>
<td>real</td>
<td>100</td>
<td>Maximum initial perturbation value</td>
</tr>
<tr>
<td>imfac</td>
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<td>0.1</td>
<td>Gradient norm factor when initializing perturbation</td>
</tr>
<tr>
<td>pmax</td>
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<td>100</td>
<td>Maximum perturbation when increasing value</td>
</tr>
<tr>
<td>pgfac</td>
<td>real</td>
<td>10</td>
<td>Perturbation growth when increasing value</td>
</tr>
<tr>
<td>pmgfac</td>
<td>real</td>
<td>0.1</td>
<td>Gradient norm factor when increasing perturbation</td>
</tr>
<tr>
<td>pm1n</td>
<td>real</td>
<td>(10^{-12})</td>
<td>Minimum non-zero perturbation when decreasing value</td>
</tr>
<tr>
<td>psfac</td>
<td>real</td>
<td>0.4</td>
<td>Perturbation shrink factor when decreasing value</td>
</tr>
<tr>
<td>pm2fac</td>
<td>real</td>
<td>0.1</td>
<td>Gradient norm factor when decreasing perturbation</td>
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<td>(\nu_1) in step update</td>
</tr>
<tr>
<td>nu2</td>
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<td>(\nu_2) in step update</td>
</tr>
<tr>
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<td>(\nu_3) in step update</td>
</tr>
<tr>
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<td>real</td>
<td>1.25</td>
<td>(\nu_4) in step update</td>
</tr>
<tr>
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<td>(\omega_1) in step update</td>
</tr>
<tr>
<td>omegal2</td>
<td>real</td>
<td>0.50</td>
<td>(\omega_2) in step update</td>
</tr>
<tr>
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<td>1.00</td>
<td>(\omega_3) in step update</td>
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continues on next page
Table 2.17 – continued from previous page

<table>
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<th>Description</th>
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</tr>
<tr>
<td>alpha3</td>
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<td>1.00</td>
<td>$\alpha_3$ in reduction update</td>
</tr>
<tr>
<td>alpha4</td>
<td>real</td>
<td>2.00</td>
<td>$\alpha_4$ in reduction update</td>
</tr>
<tr>
<td>alpha5</td>
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<td>4.00</td>
<td>$\alpha_5$ in reduction update</td>
</tr>
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</tr>
<tr>
<td>mu2</td>
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<td>$\mu_2$ in interpolation update</td>
</tr>
<tr>
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<td>$\gamma_1$ in interpolation update</td>
</tr>
<tr>
<td>gamma2</td>
<td>real</td>
<td>0.50</td>
<td>$\gamma_2$ in interpolation update</td>
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<td>2.00</td>
<td>$\gamma_3$ in interpolation update</td>
</tr>
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</tr>
<tr>
<td>theta</td>
<td>real</td>
<td>0.05</td>
<td>$\theta$ in interpolation update</td>
</tr>
</tbody>
</table>

The system of equations is approximately solved by applying the conjugate gradient method, Nash conjugate gradient method, Steihaug-Toint conjugate gradient method, generalized Lanczos method, or an alternative Krylov subspace method supplied by PETSc. The method used to solve the systems of equations is specified with the command line argument `-tao_nls_ksp_type <cg,nash,stcg,gltr,gmres,...>`; `stcg` is the default. See the PETSc manual for further information on changing the behavior of the linear system solvers.

A good preconditioner reduces the number of iterations required to solve the linear system of equations. For the conjugate gradient methods and generalized Lanczos method, this preconditioner must be symmetric and positive definite. The available options are to use no preconditioner, the absolute value of the diagonal of the Hessian matrix, a limited-memory BFGS approximation to the Hessian matrix, or one of the other preconditioners provided by the PETSc package. These preconditioners are specified by the command line arguments `-tao_nls_pc_type <none,jacobi,icc,ilu,lmvm>` respectively. The default is the `lmvm` preconditioner, which uses a BFGS approximation of the inverse Hessian. See the PETSc manual for further information on changing the behavior of the preconditioners.

The perturbation $\rho_k$ is added when the direction returned by the Krylov subspace method is not a descent direction, the Krylov method diverged due to an indefinite preconditioner or matrix, or a direction of negative curvature was found. In the last two cases, if the step returned is a descent direction, it is used during the line search. Otherwise, a steepest descent direction is used during the line search. The perturbation is decreased as long as the Krylov subspace method reports success and increased if further problems are encountered. There are three cases: initializing, increasing, and decreasing the perturbation. These cases are described below.

1. If $\rho_k$ is zero and a problem was detected with either the direction or the Krylov subspace method, the perturbation is initialized to

   $$\rho_{k+1} = \text{median} \{ \text{imin}, \text{imfac} \ast \|g(x_k)\|, \text{imax} \},$$

   where $g(x_k)$ is the gradient of the objective function and $\text{imin}$ is set with the command line argument `-tao_nls_imin <real>` with a default value of $10^{-4}$, $\text{imfac}$ by `-tao_nls_imfac` with a default value of 0.1, and $\text{imax}$ by `-tao_nls_imax` with a default value of 100. When using the `gltr` method...
to solve the system of equations, an estimate of the minimum eigenvalue $\lambda_1$ of the Hessian matrix is available. This value is used to initialize the perturbation to $\rho_{k+1} = \max \{\rho_{k+1}, -\lambda_1\}$ in this case.

2. If $\rho_k$ is nonzero and a problem was detected with either the direction or Krylov subspace method, the perturbation is increased to

$$\rho_{k+1} = \min \{p_{\text{max}}, \max \{p_{\text{gfac}} \ast \rho_k, p_{\text{mgf}a}\ast \|g(x_k)\|\}\},$$

where $g(x_k)$ is the gradient of the objective function and $p_{\text{gfac}}$ is set with the command line argument -tao_nls_pgfac with a default value of 10, $p_{\text{mgf}a}$ by -tao_nls_pmgf with a default value of 0.1, and $p_{\text{max}}$ by -tao_nls_pmax with a default value of 100.

3. If $\rho_k$ is nonzero and no problems were detected with either the direction or Krylov subspace method, the perturbation is decreased to

$$\rho_{k+1} = \min \{p_{\text{psf}a} \ast \rho_k, p_{\text{msf}a} \ast \|g(x_k)\|\},$$

where $g(x_k)$ is the gradient of the objective function, $p_{\text{psf}a}$ is set with the command line argument -tao_nls_psfac with a default value of 0.4, and $p_{\text{msf}a}$ is set by -tao_nls_pmsf with a default value of 0.1. Moreover, if $\rho_{k+1} < p_{\text{min}}$, then $\rho_{k+1} = 0$, where $p_{\text{min}}$ is set with the command line argument -tao_nls_pmin and has a default value of $10^{-12}$.

Near a local minimizer to the unconstrained optimization problem, the Hessian matrix will be positive-semidefinite; the perturbation will shrink toward zero, and one would eventually observe a superlinear convergence rate.

When using nash, stcg, or gltr to solve the linear systems of equation, a trust-region radius needs to be initialized and updated. This trust-region radius simultaneously limits the size of the step computed and reduces the number of iterations of the conjugate gradient method. The method for initializing the trust-region radius is set with the command line argument -tao_nls_init_type <constant,direction,interpolation>: interpolation, which chooses an initial value based on the interpolation scheme found in [CGT00], is the default. This scheme performs a number of function and gradient evaluations to determine a radius such that the reduction predicted by the quadratic model along the gradient direction coincides with the actual reduction in the nonlinear function. The iterate obtaining the best objective function value is used as the starting point for the main line search algorithm. The constant method initializes the trust-region radius by using the value specified with the -tao_trust0 <real> command line argument, where the default value is 100. The direction technique solves the first quadratic optimization problem by using a standard conjugate gradient method and initializes the trust region to $\|s_0\|$.

The method for updating the trust-region radius is set with the command line argument -tao_nls_update_type <step,reduction,interpolation>: step is the default. The step method updates the trust-region radius based on the value of $\tau_k$. In particular,

$$\Delta_{k+1} = \left\{ \begin{array}{ll}
\omega_1 \min(\Delta_k, \|d_k\|) & \text{if } \tau_k \in [0, \nu_1) \\
\omega_2 \min(\Delta_k, \|d_k\|) & \text{if } \tau_k \in [\nu_1, \nu_2) \\
\omega_3 \Delta_k & \text{if } \tau_k \in [\nu_2, \nu_3) \\
\max(\Delta_k, \omega_4 \|d_k\|) & \text{if } \tau_k \in [\nu_3, \nu_4) \\
\max(\Delta_k, \omega_5 \|d_k\|) & \text{if } \tau_k \in [\nu_4, \infty), 
\end{array} \right.$$

where $0 < \omega_1 < \omega_2 < \omega_3 = 1 < \omega_4 < \omega_5$ and $0 < \nu_1 < \nu_2 < \nu_3 < \nu_4$ are constants. The reduction method computes the ratio of the actual reduction in the objective function to the reduction predicted by the quadratic model for the full step, $\kappa_k = \frac{f(x_k) - f(x_k + d_k)}{q(x_k) - q(x_k + d_k)}$, where $q_k$ is the quadratic model. The radius is then updated as

$$\Delta_{k+1} = \left\{ \begin{array}{ll}
\alpha_1 \min(\Delta_k, \|d_k\|) & \text{if } \kappa_k \in (-\infty, \eta_1) \\
\alpha_2 \min(\Delta_k, \|d_k\|) & \text{if } \kappa_k \in [\eta_1, \eta_2) \\
\alpha_3 \Delta_k & \text{if } \kappa_k \in [\eta_2, \eta_3) \\
\max(\Delta_k, \alpha_4 \|d_k\|) & \text{if } \kappa_k \in [\eta_3, \eta_4) \\
\max(\Delta_k, \alpha_5 \|d_k\|) & \text{if } \kappa_k \in [\eta_4, \infty), 
\end{array} \right.$$
where $0 < \alpha_1 < \alpha_2 < \alpha_3 = 1 < \alpha_4 < \alpha_5$ and $0 < \eta_1 < \eta_2 < \eta_3 < \eta_4$ are constants. The interpolation method uses the same interpolation mechanism as in the initialization to compute a new value for the trust-region radius.

This algorithm will be deprecated in the next version and replaced by the Bounded Newton Line Search (BNLS) algorithm that can solve both bound constrained and unconstrained problems.

**Newton Trust-Region Method (NTR)**

The Newton trust-region method solves the constrained quadratic programming problem

$$\min_d \frac{1}{2} d^T H_k d + g_k^T d \quad \text{subject to} \quad \|d\| \leq \Delta_k$$


to obtain a direction $d_k$, where $H_k$ is the Hessian of the objective function at $x_k$, $g_k$ is the gradient of the objective function at $x_k$, and $\Delta_k$ is the trust-region radius. If $x_k + d_k$ sufficiently reduces the nonlinear objective function, then the step is accepted, and the trust-region radius is updated. However, if $x_k + d_k$ does not sufficiently reduce the nonlinear objective function, then the step is rejected, the trust-region radius is reduced, and the quadratic program is re-solved by using the updated trust-region radius. The Newton trust-region method can be set by using the TAO solver `tao_ntr`. The options available for this solver are listed in Table 2.18. For the best efficiency, function and gradient evaluations should be performed separately when using this algorithm.

<table>
<thead>
<tr>
<th>Name <code>-tao_ntr_</code></th>
<th>Value</th>
<th>Default</th>
<th>Description</th>
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</tr>
<tr>
<td><code>pc_type</code></td>
<td>none, jacobi</td>
<td>luvm</td>
<td>PCType for linear system</td>
</tr>
<tr>
<td><code>init_type</code></td>
<td>constant, direction, interpolation</td>
<td>interpolation</td>
<td>Radius initialization method</td>
</tr>
<tr>
<td><code>mu1_i</code></td>
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<td>0.35</td>
<td>$\mu_1$ in interpolation init</td>
</tr>
<tr>
<td><code>mu2_i</code></td>
<td>real</td>
<td>0.50</td>
<td>$\mu_2$ in interpolation init</td>
</tr>
<tr>
<td><code>gamma1_i</code></td>
<td>real</td>
<td>0.0625</td>
<td>$\gamma_1$ in interpolation init</td>
</tr>
<tr>
<td><code>gamma2_i</code></td>
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<td>$\gamma_2$ in interpolation init</td>
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</tr>
<tr>
<td><code>gamma4_i</code></td>
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<td>5.00</td>
<td>$\gamma_4$ in interpolation init</td>
</tr>
<tr>
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<td>real</td>
<td>0.25</td>
<td>$\theta$ in interpolation init</td>
</tr>
<tr>
<td><code>update_type</code></td>
<td>step, reduction, interpolation</td>
<td>step</td>
<td>Radius update method</td>
</tr>
<tr>
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<td>$\mu_1$ in interpolation init</td>
</tr>
<tr>
<td><code>mu2_i</code></td>
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<td>$\mu_2$ in interpolation init</td>
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<td><code>gamma1_i</code></td>
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<td>$\gamma_1$ in interpolation init</td>
</tr>
<tr>
<td><code>gamma2_i</code></td>
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<td>0.50</td>
<td>$\gamma_2$ in interpolation init</td>
</tr>
<tr>
<td><code>gamma3_i</code></td>
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<td>$\gamma_3$ in interpolation init</td>
</tr>
<tr>
<td><code>gamma4_i</code></td>
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<td>5.00</td>
<td>$\gamma_4$ in interpolation init</td>
</tr>
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<td>$\theta$ in interpolation init</td>
</tr>
<tr>
<td><code>eta1</code></td>
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<td>:</td>
<td>$\eta_1$ in reduction update</td>
</tr>
<tr>
<td><code>eta2</code></td>
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<td>$\eta_2$ in reduction update</td>
</tr>
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<td><code>eta3</code></td>
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</tr>
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<td>$\eta_4$ in reduction update</td>
</tr>
<tr>
<td><code>alpha1</code></td>
<td>real</td>
<td>0.25</td>
<td>$\alpha_1$ in reduction update</td>
</tr>
<tr>
<td><code>alpha2</code></td>
<td>real</td>
<td>0.50</td>
<td>$\alpha_2$ in reduction update</td>
</tr>
<tr>
<td><code>alpha3</code></td>
<td>real</td>
<td>1.00</td>
<td>$\alpha_3$ in reduction update</td>
</tr>
<tr>
<td><code>alpha4</code></td>
<td>real</td>
<td>2.00</td>
<td>$\alpha_4$ in reduction update</td>
</tr>
<tr>
<td><code>alpha5</code></td>
<td>real</td>
<td>4.00</td>
<td>$\alpha_5$ in reduction update</td>
</tr>
</tbody>
</table>

continues on next page

170 Chapter 2. The Solvers in PETSc/TAO
The quadratic optimization problem is approximately solved by applying the Nash or Steihaug-Toint conjugate gradient methods or the generalized Lanczos method to the symmetric system of equations \( H_k d = -g_k \). The method used to solve the system of equations is specified with the command line argument `-tao_ntr_ksp_type <nash, stcg, gltr>;` stcg is the default. See the PETSc manual for further information on changing the behavior of these linear system solvers.

A good preconditioner reduces the number of iterations required to compute the direction. For the Nash and Steihaug-Toint conjugate gradient methods and generalized Lanczos method, this preconditioner must be symmetric and positive definite. The available options are to use no preconditioner, the absolute value of the diagonal of the Hessian matrix, a limited-memory BFGS approximation to the Hessian matrix, or one of the other preconditioners provided by the PETSc package. These preconditioners are specified by the command line argument `-tao_ntr_pc_type <none, jacobi, icc, ilu, lvm>;` respectively. The default is the lvm preconditioner. See the PETSc manual for further information on changing the behavior of the preconditioners.

The method for computing an initial trust-region radius is set with the command line arguments `-tao_ntr_init_type <constant, direction, interpolation>;` interpolation, which chooses an initial value based on the interpolation scheme found in [CGT06], is the default. This scheme performs a number of function and gradient evaluations to determine a radius such that the reduction predicted by the quadratic model along the gradient direction coincides with the actual reduction in the nonlinear function. The iterate obtaining the best objective function value is used as the starting point for the main trust-region algorithm. The constant method initializes the trust-region radius by using the value specified with the `-tao_trust0 <real>;` command line argument, where the default value is 100. The direction technique solves the first quadratic optimization problem by using a standard conjugate gradient method and initializes the trust region to \( \| s_0 \| \).

The method for updating the trust-region radius is set with the command line arguments `-tao_ntr_update_type <reduction, interpolation>;` reduction is the default. The reduction method computes the ratio of the actual reduction in the objective function to the reduction predicted by the quadratic model for the full step, \( \kappa_k = \frac{f(x_k) - f(x_k + d_k)}{q(x_k) - q(x_k + d_k)} \), where \( q_k \) is the quadratic model. The radius is then updated as

\[
\Delta_{k+1} = \begin{cases} 
\alpha_1 \min(\Delta_k, \|d_k\|) & \text{if } \kappa_k \in (-\infty, \eta_1) \\
\alpha_2 \min(\Delta_k, \|d_k\|) & \text{if } \kappa_k \in [\eta_1, \eta_2) \\
\alpha_3 \Delta_k & \text{if } \kappa_k \in [\eta_2, \eta_3) \\
\max(\Delta_k, \alpha_4 \|d_k\|) & \text{if } \kappa_k \in [\eta_3, \eta_4) \\
\max(\Delta_k, \alpha_5 \|d_k\|) & \text{if } \kappa_k \in [\eta_4, \infty) 
\end{cases}
\]

where \( 0 < \alpha_1 < \alpha_2 < \alpha_3 = 1 < \alpha_4 < \alpha_5 \) and \( 0 < \eta_1 < \eta_2 < \eta_3 < \eta_4 \) are constants. The interpolation method uses the same interpolation mechanism as in the initialization to compute a new value for the trust-region radius.

This algorithm will be deprecated in the next version and replaced by the Bounded Newton Trust Region (BNTR) algorithm that can solve both bound constrained and unconstrained problems.

<table>
<thead>
<tr>
<th>Name -tao_ntr_</th>
<th>Value</th>
<th>Default</th>
<th>Description</th>
</tr>
</thead>
<tbody>
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<td>mu1</td>
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<td>0.10</td>
<td>( \mu_1 ) in interpolation update</td>
</tr>
<tr>
<td>mu2</td>
<td>real</td>
<td>0.50</td>
<td>( \mu_2 ) in interpolation update</td>
</tr>
<tr>
<td>gamma1</td>
<td>real</td>
<td>0.25</td>
<td>( \gamma_1 ) in interpolation update</td>
</tr>
<tr>
<td>gamma2</td>
<td>real</td>
<td>0.50</td>
<td>( \gamma_2 ) in interpolation update</td>
</tr>
<tr>
<td>gamma3</td>
<td>real</td>
<td>2.00</td>
<td>( \gamma_3 ) in interpolation update</td>
</tr>
<tr>
<td>gamma4</td>
<td>real</td>
<td>4.00</td>
<td>( \gamma_4 ) in interpolation update</td>
</tr>
<tr>
<td>theta</td>
<td>real</td>
<td>0.05</td>
<td>( \theta ) in interpolation update</td>
</tr>
</tbody>
</table>
Newton Trust Region with Line Search (NTL)

NTL safeguards the trust-region globalization such that a line search is used in the event that the step is initially rejected by the predicted versus actual decrease comparison. If the line search fails to find a viable step length for the Newton step, it falls back onto a scaled gradient or a gradient descent step. The trust radius is then modified based on the line search step length.

This algorithm will be deprecated in the next version and replaced by the Bounded Newton Trust Region with Line Search (BNTL) algorithm that can solve both bound constrained and unconstrained problems.

Limited-Memory Variable-Metric Method (LMVM)

The limited-memory, variable-metric method (LMVM) computes a positive definite approximation to the Hessian matrix from a limited number of previous iterates and gradient evaluations. A direction is then obtained by solving the system of equations

\[ H_k d_k = -\nabla f(x_k), \]

where \( H_k \) is the Hessian approximation obtained by using the BFGS update formula. The inverse of \( H_k \) can readily be applied to obtain the direction \( d_k \). Having obtained the direction, a Moré-Thuente line search is applied to compute a step length, \( \tau_k \), that approximately solves the one-dimensional optimization problem

\[ \min_{\tau} f(x_k + \tau d_k). \]

The current iterate and Hessian approximation are updated, and the process is repeated until the method converges. This algorithm is the default unconstrained minimization solver and can be selected by using the TAO solver `tao_lmvm`. For best efficiency, function and gradient evaluations should be performed simultaneously when using this algorithm.

The primary factors determining the behavior of this algorithm are the type of Hessian approximation used, the number of vectors stored for the approximation and the initialization/scaling of the approximation. These options can be configured using the `-tao_lmvm_mat_lmvm` prefix. For further detail, we refer the reader to the `MATLMVM` matrix type definitions in the PETSc Manual.

The LMVM algorithm also allows the user to define a custom initial Hessian matrix \( H_{0,k} \) through the interface function `TaoLMVMSetH0()`. This user-provided initialization overrides any other scalar or diagonal initialization inherent to the LMVM approximation. The provided \( H_{0,k} \) must be a PETSc `Mat` type object that represents a positive-definite matrix. The approximation prefers `MatSolve()` if the provided matrix has `MATOP_SOLVE` implemented. Otherwise, `MatMult()` is used in a KSP solve to perform the inversion of the user-provided initial Hessian.

In applications where `TaoSolve()` on the LMVM algorithm is repeatedly called to solve similar or related problems, `-tao_lmvm_recycle` flag can be used to prevent resetting the LMVM approximation between subsequent solutions. This recycling also avoids one extra function and gradient evaluation, instead re-using the values already computed at the end of the previous solution.

This algorithm will be deprecated in the next version and replaced by the Bounded Quasi-Newton Line Search (BQNLS) algorithm that can solve both bound constrained and unconstrained problems.
**Nonlinear Conjugate Gradient Method (CG)**

The nonlinear conjugate gradient method can be viewed as an extension of the conjugate gradient method for solving symmetric, positive-definite linear systems of equations. This algorithm requires only function and gradient evaluations as well as a line search. The TAO implementation uses a Moré-Thuente line search to obtain the step length. The nonlinear conjugate gradient method can be selected by using the TAO solver \texttt{tao_cg}. For the best efficiency, function and gradient evaluations should be performed simultaneously when using this algorithm.

Five variations are currently supported by the TAO implementation: the Fletcher-Reeves method, the Polak-Ribiére method, the Polak-Ribiére-Plus method \cite{NW06}, the Hestenes-Stiefel method, and the Dai-Yuan method. These conjugate gradient methods can be specified by using the command line argument \texttt{-tao_cg_type <fr,pr,prp,hs,dy>}, respectively. The default value is \texttt{prp}.

The conjugate gradient method incorporates automatic restarts when successive gradients are not sufficiently orthogonal. TAO measures the orthogonality by dividing the inner product of the gradient at the current point and the gradient at the previous point by the square of the Euclidean norm of the gradient at the current point. When the absolute value of this ratio is greater than $\eta$, the algorithm restarts using the gradient direction. The parameter $\eta$ can be set by using the command line argument \texttt{-tao_cg_eta <real>}; 0.1 is the default value.

This algorithm will be deprecated in the next version and replaced by the Bounded Nonlinear Conjugate Gradient (BNCG) algorithm that can solve both bound constrained and unconstrained problems.

**Nelder-Mead Simplex Method (NM)**

The Nelder-Mead algorithm \cite{NM65} is a direct search method for finding a local minimum of a function $f(x)$. This algorithm does not require any gradient or Hessian information of $f$ and therefore has some expected advantages and disadvantages compared to the other TAO solvers. The obvious advantage is that it is easier to write an application when no derivatives need to be calculated. The downside is that this algorithm can be slow to converge or can even stagnate, and it performs poorly for large numbers of variables.

This solver keeps a set of $N + 1$ sorted vectors $x_1, x_2, \ldots, x_{N+1}$ and their corresponding objective function values $f_1 \leq f_2 \leq \ldots \leq f_{N+1}$. At each iteration, $x_{N+1}$ is removed from the set and replaced with

$$x(\mu) = (1 + \mu) \frac{1}{N} \sum_{i=1}^{N} x_i - \mu x_{N+1},$$

where $\mu$ can be one of $\mu_0, 2\mu_0, \frac{1}{2}\mu_0, -\frac{1}{2}\mu_0$ depending on the values of each possible $f(x(\mu))$.

The algorithm terminates when the residual $f_{N+1} - f_1$ becomes sufficiently small. Because of the way new vectors can be added to the sorted set, the minimum function value and/or the residual may not be impacted at each iteration.

Two options can be set specifically for the Nelder-Mead algorithm:

\texttt{-tao_nm_lambda <value>}

sets the initial set of vectors ($x_0$ plus \texttt{value} in each coordinate direction); the default value is 1.

\texttt{-tao_nm_mu <value>}

sets the value of $\mu_0$; the default is $\mu_0 = 1$. 

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Bound-Constrained Optimization

Bound-constrained optimization algorithms solve optimization problems of the form

$$\min_x f(x)$$

subject to $$l \leq x \leq u.$$ \hspace{1cm} (1)

These solvers use the bounds on the variables as well as objective function, gradient, and possibly Hessian information.

For any unbounded variables, the bound value for the associated index can be set to PETSC_INFINITY for the upper bound and PETSC_NINFINITY for the lower bound. If all bounds are set to infinity, then the bounded algorithms are equivalent to their unconstrained counterparts.

Before introducing specific methods, we will first define two projection operations used by all bound-constrained algorithms.

- **Gradient projection:**
  \[ P(g) = \begin{cases} 
  0 & \text{if } (x \leq l_i \land g_i > 0) \lor (x \geq u_i \land g_i < 0) \\
  g_i & \text{otherwise} 
  \end{cases} \]

- **Bound projection:**
  \[ B(x) = \begin{cases} 
  l_i & \text{if } x_i < l_i \\
  u_i & \text{if } x_i > u_i \\
  x_i & \text{otherwise} 
  \end{cases} \]

Bounded Newton-Krylov Methods

TAO features three bounded Newton-Krylov (BNK) class of algorithms, separated by their globalization methods: projected line search (BNLS), trust region (BNTR), and trust region with a projected line search fall-back (BNTL). They are available via the TAO solvers TAOBNLS, TAOBNTR and TAOBNTL, respectively, or the -tao_type bnls/bntr/bntl flag.

The BNK class of methods use an active-set approach to solve the symmetric system of equations,

$$H_k p_k = -g_k,$$

only for inactive variables in the interior of the bounds. The active-set estimation is based on Bertsekas [Ber82] with the following variable index categories:

- **lower bounded:** $$L(x) = \{ i : x_i \leq l_i + \epsilon \land g(x)_i > 0 \},$$
- **upper bounded:** $$U(x) = \{ i : x_i \geq u_i + \epsilon \land g(x)_i < 0 \},$$
- **fixed:** $$F(x) = \{ i : l_i = u_i \},$$
- **active-set:** $$A(x) = \{ L(x) \cup U(x) \cup F(x) \},$$
- **inactive-set:** $$I(x) = \{ 1, 2, \ldots, n \} \setminus A(x).$$

At each iteration, the bound tolerance is estimated as $$\epsilon_{k+1} = \min(\epsilon_k, \|w_k\|_2)$$ with $$w_k = x_k - B(x_k - \beta D_k g_k),$$ where the diagonal matrix $$D_k$$ is an approximation of the Hessian inverse $$H_k^{-1}$$. The initial bound tolerance $$\epsilon_0$$ and the step length $$\beta$$ have default values of 0.001 and can be adjusted using -tao_bnk_as_tol and -tao_bnk_as_step flags, respectively. The active-set estimation can be disabled using the option -tao_bnk_as_type none, in which case the algorithm simply uses the current iterate with no bound tolerances to determine which variables are actively bounded and which are free.

BNK algorithms invert the reduced Hessian using a Krylov iterative method. Trust-region conjugate gradient methods (KSPNASH, KSPSTCG, and KSPGLTR) are required for the BNTR and BNTL algorithms, and recommended for the BNLS algorithm. The preconditioner type can be changed using the -tao_bnk_pc_type flag.
none/ilu/icc/jacobi/lmvm. The \texttt{lmvm} option, which is also the default, preconditions the Krylov solution with a \texttt{MATLMVM} matrix. The remaining supported preconditioner types are default PETSc types. If Jacobi is selected, the diagonal values are safeguarded to be positive. \texttt{icc} and \texttt{ilu} options produce good results for problems with dense Hessians. The LMVM and Jacobi preconditioners are also used as the approximate inverse-Hessian in the active-set estimation. If neither are available, or if the Hessian matrix does not have \texttt{MATOP_GET_DIAGONAL} defined, then the active-set estimation falls back onto using an identity matrix in place of $D_k$ (this is equivalent to estimating the active-set using a gradient descent step).

A special option is available to \texttt{accelerate} the convergence of the BNK algorithms by taking a finite number of BNCG iterations at each Newton iteration. By default, the number of BNCG iterations is set to zero and the algorithms do not take any BNCG steps. This can be changed using the option flag \texttt{-tao_bnk_max_cg_its <i>}. While this reduces the number of Newton iterations, in practice it simply trades off the Hessian evaluations in the BNK solver for more function and gradient evaluations in the BNCG solver. However, it may be useful for certain types of problems where the Hessian evaluation is disproportionately more expensive than the objective function or its gradient.

**Bounded Newton Line Search (BNLS)**

BNLS safeguards the Newton step by falling back onto a BFGS, scaled gradient, or gradient steps based on descent direction verifications. For problems with indefinite Hessian matrices, the step direction is calculated using a perturbed system of equations,

$$(H_k + \rho_k I)p_k = -g_k,$$

where $\rho_k$ is a dynamically adjusted positive constant. The step is globalized using a projected Moré-Thuente line search. If a trust-region conjugate gradient method is used for the Hessian inversion, the trust radius is modified based on the line search step length.

**Bounded Newton Trust Region (BNTR)**

BNTR globalizes the Newton step using a trust region method based on the predicted versus actual reduction in the cost function. The trust radius is increased only if the accepted step is at the trust region boundary. The reduction check features a safeguard for numerical values below machine epsilon, scaled by the latest function value, where the full Newton step is accepted without modification.

**Bounded Newton Trust Region with Line Search (BNTL)**

BNTL safeguards the trust-region globalization such that a line search is used in the event that the step is initially rejected by the predicted versus actual decrease comparison. If the line search fails to find a viable step length for the Newton step, it falls back onto a scaled gradient or a gradient descent step. The trust radius is then modified based on the line search step length.
Bounded Quasi-Newton Line Search (BQNLS)

The BQNLS algorithm uses the BNLS infrastructure, but replaces the step calculation with a direct inverse application of the approximate Hessian based on quasi-Newton update formulas. No Krylov solver is used in the solution, and therefore the quasi-Newton method chosen must guarantee a positive-definite Hessian approximation. This algorithm is available via `tao_type bqnls`.

Bounded Quasi-Newton-Krylov

BQNK algorithms use the BNK infrastructure, but replace the exact Hessian with a quasi-Newton approximation. The matrix-free forward product operation based on quasi-Newton update formulas are used in conjunction with Krylov solvers to compute step directions. The quasi-Newton inverse application is used to precondition the Krylov solution, and typically helps converge to a step direction in $O(10)$ iterations. This approach is most useful with quasi-Newton update types such as Symmetric Rank-1 that cannot strictly guarantee positive-definiteness. The BNLS framework with Hessian shifting, or the BNTR framework with trust region safeguards, can successfully compensate for the Hessian approximation becoming indefinite.

Similar to the full Newton-Krylov counterpart, BQNK algorithms come in three forms separated by the globalization technique: line search (BQNKLS), trust region (BQNKTR) and trust region w/ line search fall-back (BQNKTL). These algorithms are available via `tao_type <bqnkls, bqnktr, bqnktl>`.

Bounded Nonlinear Conjugate Gradient (BNCG)

BNCG extends the unconstrained nonlinear conjugate gradient algorithm to bound constraints via gradient projections and a bounded Moré-Thuente line search.

Like its unconstrained counterpart, BNCG offers gradient descent and a variety of CG updates: Fletcher-Reeves, Polak-Ribiére, Polak-Ribiére-Plus, Hestenes-Stiefel, Dai-Yuan, Hager-Zhang, Dai-Kou, Kou-Dai, and the Self-Scaling Memoryless (SSML) BFGS, DFP, and Broyden methods. These methods can be specified by using the command line argument `-tao_bncg_type <gd,fr,pr,prp,hs,dy,hz,dk,ssml_bfgs,ssml_dfps,ssml_brkdn>`, respectively. The default value is `ssml_bfgs`. We have scalar preconditioning for these methods, and it is controlled by the flag `tao_bncg_alpha`. To disable rescaling, use $\alpha = -1.0$, otherwise $\alpha \in [0,1]$. BNCG is available via the TAO solver `TAOBNGC` or the `-tao_type bncg` flag.

Some individual methods also contain their own parameters. The Hager-Zhang and Dou-Kai methods have a parameter that determines the minimum amount of contribution the previous search direction gives to the next search direction. The flags are `-tao_bncg_hz_eta` and `-tao_bncg_dk_eta`, and by default are set to 0.4 and 0.5 respectively. The Kou-Dai method has multiple parameters. `-tao_bncg_zeta` serves the same purpose as the previous two; set to 0.1 by default. There is also a parameter to scale the contribution of $y_k \equiv \nabla f(x_k) - \nabla f(x_{k-1})$ in the search direction update. It is controlled by `-tao_bncg_xi`, and is equal to 1.0 by default. There are also times where we want to maximize the descent as measured by $\nabla f(x_k)^T d_k$, and that may be done by using a negative value of $\xi$; this achieves better performance when not using the diagonal preconditioner described next. This is enabled by default, and is controlled by `-tao_bncg_neg_xi`. Finally, the Broyden method has its convex combination parameter, set with `-tao_bncg_theta`. We have this as 1.0 by default, i.e. it is by default the BFGS method. One can also individually tweak the BFGS and DFP contributions using the multiplicative constants `-tao_bncg_scale`; both are set to 1 by default.

All methods can be scaled using the parameter `-tao_bncg_alpha`, which continuously varies in $[0,1]$. The default value is set depending on the method from initial testing.

BNCG also offers a special type of method scaling. It employs Broyden diagonal scaling as an option for its CG methods, turned on with the flag `-tao_bncg_diag_scaling`. Formulations for both the forward (reg-
ular) and inverse Broyden methods are developed, controlled by the flag `-tao_bncg_mat_lmvm_forward`. It is set to True by default. Whether one uses the forward or inverse formulations depends on the method being used. For example, in our preliminary computations, the forward formulation works better for the SSML_BFGS method, but the inverse formulation works better for the Hestenes-Stiefel method. The convex combination parameter for the Broyden scaling is controlled by `-tao_bncg_mat_lmvm_theta`, and is 0 by default. We also employ rescaling of the Broyden diagonal, which aids the linesearch immensely. The rescaling parameter is controlled by `-tao_bncg_mat_lmvm_alpha`, and should be $\in [0, 1]$. One can disable rescaling of the Broyden diagonal entirely by setting `-tao_bncg_mat_lmvm_sigma_hist 0`.

One can also supply their own preconditioner, serving as a Hessian initialization to the above diagonal scaling. The appropriate user function in the code is `TaoBNCGSetH0(tao, H0)` where $H0$ is the user-defined `Mat` object that serves as a preconditioner. For an example of similar usage, see `tao/tutorials/ex3.c`.

The active set estimation uses the Bertsekas-based method described in *Bounded Newton-Krylov Methods*, which can be deactivated using `-tao_bncg_as_type none`, in which case the algorithm will use the current iterate to determine the bounded variables with no tolerances and no look-ahead step. As in the BNK algorithm, the initial bound tolerance and estimator step length used in the Bertsekas method can be set via `-tao_bncg_as_tol` and `-tao_bncg_as_step`, respectively.

In addition to automatic scaled gradient descent restarts under certain local curvature conditions, we also employ restarts based on a check on descent direction such that $\nabla f(x_k)^T d_k \in [-10^{11}, -10^{-9}]$. Furthermore, we allow for a variety of alternative restart strategies, all disabled by default. The `-tao_bncg_unscaled_restart` flag allows one to disable rescaling of the gradient for gradient descent steps. The `-tao_bncg_spaced_restart` flag tells the solver to restart every $Mn$ iterations, where $n$ is the problem dimension and $M$ is a constant determined by `-tao_bncg_min_restart_num` and is 6 by default. We also have dynamic restart strategies based on checking if a function is locally quadratic; if so, go do a gradient descent step. The flag is `-tao_bncg_dynamic_restart`, disabled by default since the CG solver usually does better in those cases anyway. The minimum number of quadratic-like steps before a restart is set using `-tao_bncg_min_quad` and is 6 by default.

**Generally Constrained Solvers**

Constrained solvers solve optimization problems that incorporate either or both equality and inequality constraints, and may optionally include bounds on solution variables.

**Alternating Direction Method of Multipliers (ADMM)**

The TAOADMM algorithm is intended to blend the decomposability of dual ascent with the superior convergence properties of the method of multipliers. [BPC+11] The algorithm solves problems in the form

$$\min_x f(x) + g(z)$$

subject to $Ax + Bz = c$

where $x \in \mathbb{R}^n$, $z \in \mathbb{R}^m$, $A \in \mathbb{R}^{p \times n}$, $B \in \mathbb{R}^{p \times m}$, and $c \in \mathbb{R}^p$. Essentially, ADMM is a wrapper over two TAO solver, one for $f(x)$, and one for $g(z)$. With method of multipliers, one can form the augmented Lagrangian

$$L_\rho(x, z, y) = f(x) + g(z) + y^T(Ax + Bz - c) + (\rho/2)\|Ax + Bz - c\|^2_2$$

Then, ADMM consists of the iterations

$$x^{k+1} := \arg\min_{x} L_\rho(x, z^k, y^k)$$

$$z^{k+1} := \arg\min_{z} L_\rho(x^{k+1}, z, y^k)$$

$$y^{k+1} := y^k + \rho(Ax^{k+1} + Bz^{k+1} - c)$$
In certain formulation of ADMM, solution of $z^{k+1}$ may have closed-form solution. Currently ADMM provides one default implementation for $z^{k+1}$, which is soft-threshold. It can be used with either `TaoADMMSetRegularizerType_ADMM()` or `-tao_admm regularizer_type <regularizer_soft_thresh>`. User can also pass spectral penalty value, $\rho$, with either `TaoADMMSetSpectralPenalty()` or `-tao_admm spectral_penalty`. Currently, user can use

- `TaoADMMSetMisfitObjectiveAndGradientRoutine()`
- `TaoADMMSetRegularizerObjectiveAndGradientRoutine()`
- `TaoADMMSetMisfitHessianRoutine()`
- `TaoADMMSetRegularizerHessianRoutine()`

Any other combination of routines is currently not supported. Hessian matrices can either be constant or non-constant, of which fact can be set via `TaoADMMSetMisfitHessianChangeStatus()`, and `TaoADMMSetRegularizerHessianChangeStatus()`. Also, it may appear in certain cases where augmented Lagrangian’s Hessian may become nearly singular depending on the $\rho$, which may change in the case of `-tao_admm dual_update <update_adaptive>, <update_adaptive_relaxed>`. This issue can be prevented by `TaoADMMSetMinimumSpectralPenalty()`.

### Augmented Lagrangian Method of Multipliers (ALMM)

The TAOALMM method solves generally constrained problems of the form

$$\min_{x} \quad f(x)$$

subject to

$$g(x) = 0$$

$$h(x) \geq 0$$

$$l \leq x \leq u$$

where $g(x)$ are equality constraints, $h(x)$ are inequality constraints and $l$ and $u$ are lower and upper bounds on the optimization variables, respectively.

TAOALMM converts the above general constrained problem into a sequence of bound constrained problems at each outer iteration $k = 1, 2, \ldots$

$$\min_{x} \quad L(x, \lambda_k)$$

subject to

$$l \leq x \leq u$$

where $L(x, \lambda_k)$ is the augmented Lagrangian merit function and $\lambda_k$ is the Lagrange multiplier estimates at outer iteration $k$.

TAOALMM offers two versions of the augmented Lagrangian formulation: the canonical Hestenes-Powell augmented Lagrangian [Hes69] [Pow69] with inequality constrained converted to equality constraints via slack variables, and the slack-less Powell-Hestenes-Rockafellar formulation [Roc74] that utilizes a pointwise max() on the inequality constraints. For most applications, the canonical Hestenes-Powell formulation is likely to perform better. However, the PHR formulation may be desirable for problems featuring very large numbers of inequality constraints as it avoids inflating the dimension of the subproblem with slack variables.

The inner subproblem is solved using a nested bound-constrained first-order TAO solver. By default, TAOALM uses a quasi-Newton-Krylov trust-region method (TAOBQNKTR). Other first-order methods such as TAOBNCG and TAOBQNLS are also appropriate, but a trust-region globalization is strongly recommended for most applications.
Primal-Dual Interior-Point Method (PDIPM)

The TAOPDIPM method (-tao_type pdipm) implements a primal-dual interior point method for solving general nonlinear programming problems of the form

\[
\begin{align*}
\min_x & \quad f(x) \\
\text{subject to} & \quad g(x) = 0 \\
& \quad h(x) \geq 0 \\
& \quad x^- \leq x \leq x^+
\end{align*}
\]

Here, \(f(x)\) is the nonlinear objective function, \(g(x), h(x)\) are the equality and inequality constraints, and \(x^-\) and \(x^+\) are the lower and upper bounds on decision variables \(x\).

PDIPM converts the inequality constraints to equalities using slack variables \(z\) and a log-barrier term, which transforms (2.6) to

\[
\min f(x) - \mu \sum_{i=1}^{nci} \ln z_i \\
\text{s.t.} \\
\begin{align*}
cee(x) &= 0 \\
ci(x) - z &= 0
\end{align*}
\]

Here, \(ce(x)\) is set of equality constraints that include \(g(x)\) and fixed decision variables, i.e., \(x^- = x = x^+\). Similarly, \(ci(x)\) are inequality constraints including \(h(x)\) and lower/upper/box-constraints on \(x\). \(\mu\) is a parameter that is driven to zero as the optimization progresses.

The Lagrangian for (2.7) is

\[
L_\mu(x, \lambda_{ce}, \lambda_{ci}, z) = f(x) + \lambda_{ce}^T cce(x) - \lambda_{ci}^T (ci(x) - z) - \mu \sum_{i=1}^{nci} \ln z_i
\]

where, \(\lambda_{ce}\) and \(\lambda_{ci}\) are the Lagrangian multipliers for the equality and inequality constraints, respectively.

The first order KKT conditions for optimality are as follows

\[
\begin{bmatrix}
\nabla f(x) + \nabla cce(x)^T \lambda_{ce} - \nabla ci(x)^T \lambda_{ci} \\
cc(x) \\
cci(x) - z \\
Z \Lambda_{ce} e - \mu e
\end{bmatrix} = 0
\]

(2.9) is solved iteratively using Newton’s method using PETSc’s SNES object. After each Newton iteration, a line-search is performed to update \(x\) and enforce \(z, \lambda_{ci} \geq 0\). The barrier parameter \(\mu\) is also updated after each Newton iteration. The Newton update is obtained by solving the second-order KKT system \(Hd = -\nabla L_\mu\).

Here, \(H\) is the Hessian matrix of the KKT system. For interior-point methods such as PDIPM, the Hessian matrix tends to be ill-conditioned, thus necessitating the use of a direct solver. We recommend using LU preconditioner -pc_type lu and using direct linear solver packages such SuperLU_Dist or MUMPS.
PDE-Constrained Optimization

TAO solves PDE-constrained optimization problems of the form

$$\min_{u,v} f(u, v)$$
subject to $g(u, v) = 0$,

where the state variable $u$ is the solution to the discretized partial differential equation defined by $g$ and parametrized by the design variable $v$, and $f$ is an objective function. The Lagrange multipliers on the constraint are denoted by $y$. This method is set by using the linearly constrained augmented Lagrangian TAO solver `tao_lcl`.

We make two main assumptions when solving these problems: the objective function and PDE constraints have been discretized so that we can treat the optimization problem as finite dimensional and $\nabla_u g(u, v)$ is invertible for all $u$ and $v$.

Linearly-Constrained Augmented Lagrangian Method (LCL)

Given the current iterate $(u_k, v_k, y_k)$, the linearly constrained augmented Lagrangian method approximately solves the optimization problem

$$\min_{u,v} \tilde{f}_k(u, v)$$
subject to $A_k(u - u_k) + B_k(v - v_k) + g_k = 0$,

where $A_k = \nabla_u g(u_k, v_k)$, $B_k = \nabla_v g(u_k, v_k)$, and $g_k = g(u_k, v_k)$ and

$$\tilde{f}_k(u, v) = f(u, v) - g(u, v)^T y_k + \frac{\rho_k}{2} \|g(u, v)\|^2$$

is the augmented Lagrangian function. This optimization problem is solved in two stages. The first computes the Newton direction and finds a feasible point for the linear constraints. The second computes a reduced-space direction that maintains feasibility with respect to the linearized constraints and improves the augmented Lagrangian merit function.

Newton Step

The Newton direction is obtained by fixing the design variables at their current value and solving the linearized constraint for the state variables. In particular, we solve the system of equations

$$A_k du = -g_k$$

to obtain a direction $du$. We need a direction that provides sufficient descent for the merit function

$$\frac{1}{2} \|g(u, v)\|^2.$$ 

That is, we require $g_k^T A_k du < 0$.

If the Newton direction is a descent direction, then we choose a penalty parameter $\rho_k$ so that $du$ is also a sufficient descent direction for the augmented Lagrangian merit function. We then find $\alpha$ to approximately minimize the augmented Lagrangian merit function along the Newton direction.

$$\min_{\alpha \geq 0} \tilde{f}_k(u_k + \alpha u, v_k).$$
We can enforce either the sufficient decrease condition or the Wolfe conditions during the search procedure. The new point,

\[ u_{k+\frac{1}{2}} = u_k + \alpha_k du \]
\[ v_{k+\frac{1}{2}} = v_k, \]

satisfies the linear constraint

\[ A_k(u_{k+\frac{1}{2}} - u_k) + B_k(v_{k+\frac{1}{2}} - v_k) + \alpha_k g_k = 0. \]

If the Newton direction computed does not provide descent for the merit function, then we can use the steepest descent direction \( du = -A_k^T g_k \) during the search procedure. However, the implication that the intermediate point approximately satisfies the linear constraint is no longer true.

### Modified Reduced-Space Step

We are now ready to compute a reduced-space step for the modified optimization problem:

\[
\begin{align*}
\min_{u, v} & \quad \tilde{f}_k(u, v) \\
\text{subject to} & \quad A_k(u - u_k) + B_k(v - v_k) + \alpha_k g_k = 0.
\end{align*}
\]

We begin with the change of variables

\[
\begin{align*}
\min_{du, dv} & \quad \tilde{f}_k(u_k + du, v_k + dv) \\
\text{subject to} & \quad A_k du + B_k dv + \alpha_k g_k = 0
\end{align*}
\]

and make the substitution

\[ du = -A_k^{-1}(B_k dv + \alpha_k g_k). \]

Hence, the unconstrained optimization problem we need to solve is

\[ \min_{dv} \tilde{f}_k(u_k - A_k^{-1}(B_k dv + \alpha_k g_k), v_k + dv), \]

which is equivalent to

\[ \min_{dv} \tilde{f}_k(u_{k+\frac{1}{2}} - A_k^{-1}B_k dv, v_{k+\frac{1}{2}} + dv). \]

We apply one step of a limited-memory quasi-Newton method to this problem. The direction is obtained by solving the quadratic problem

\[ \min_{dv} \frac{1}{2} dv^T \tilde{H}_k dv + \tilde{g}_{k+\frac{1}{2}}^T dv, \]

where \( \tilde{H}_k \) is the limited-memory quasi-Newton approximation to the reduced Hessian matrix, a positive-definite matrix, and \( \tilde{g}_{k+\frac{1}{2}} \) is the reduced gradient.

\[ \tilde{g}_{k+\frac{1}{2}} = \nabla_v \tilde{f}_k(u_{k+\frac{1}{2}}, v_{k+\frac{1}{2}}) - \nabla_u \tilde{f}_k(u_{k+\frac{1}{2}}, v_{k+\frac{1}{2}}) A_k^{-1} B_k \]

\[ = d_{k+\frac{1}{2}} + c_{k+\frac{1}{2}} A_k^{-1} B_k. \]

The reduced gradient is obtained from one linearized adjoint solve

\[ y_{k+\frac{1}{2}} = A_k^T c_{k+\frac{1}{2}} \]

and some linear algebra

\[ \tilde{g}_{k+\frac{1}{2}} = d_{k+\frac{1}{2}} + y_{k+\frac{1}{2}}^T B_k. \]
Because the Hessian approximation is positive definite and we know its inverse, we obtain the direction

\[ dv = -H^{-1}_k g_k + \frac{1}{2} \]

and recover the full-space direction from one linearized forward solve,

\[ du = -A^{-1}_k B_k dv. \]

Having the full-space direction, which satisfies the linear constraint, we now approximately minimize the augmented Lagrangian merit function along the direction.

\[ \min_{\beta \geq 0} \tilde{f}_k(u_k + \frac{1}{2} + \beta dv, v_k + \frac{1}{2} + \beta dv) \]

We enforce the Wolfe conditions during the search procedure. The new point is

\[ u_{k+1} = u_k + \frac{1}{2} + \beta_k du \]
\[ v_{k+1} = v_k + \frac{1}{2} + \beta_k dv. \]

The reduced gradient at the new point is computed from

\[ y_{k+1} = A_k^{-T} c_{k+1} \]
\[ \tilde{g}_{k+1} = d_{k+1} - y_{k+1}^T B_k, \]

where \( c_{k+1} = \nabla_a \tilde{f}_k(u_{k+1}, v_{k+1}) \) and \( d_{k+1} = \nabla_v \tilde{f}_k(u_{k+1}, v_{k+1}) \). The multipliers \( y_{k+1} \) become the multipliers used in the next iteration of the code. The quantities \( v_{k+\frac{1}{2}}, v_{k}, \tilde{g}_{k+\frac{1}{2}}, \) and \( \tilde{g}_{k} \) are used to update \( H_k \) to obtain the limited-memory quasi-Newton approximation to the reduced Hessian matrix used in the next iteration of the code. The update is skipped if it cannot be performed.

**Nonlinear Least-Squares**

Given a function \( F : \mathbb{R}^n \to \mathbb{R}^m \), the nonlinear least-squares problem minimizes

\[ f(x) = \|F(x)\|_2^2 = \sum_{i=1}^{m} F_i(x)^2. \]  (2.10)

The nonlinear equations \( F \) should be specified with the function \texttt{TaoSetResidual()}.  

**Bound-constrained Regularized Gauss-Newton (BRGN)**

The TAOBRGN algorithms is a Gauss-Newton method is used to iteratively solve nonlinear least squares problem with the iterations

\[ x_{k+1} = x_k - \alpha_k (J_k^T J_k)^{-1} J_k^T r(x_k) \]

where \( r(x) \) is the least-squares residual vector, \( J_k = \partial r(x_k)/\partial x \) is the Jacobian of the residual, and \( \alpha_k \) is the step length parameter. In other words, the Gauss-Newton method approximates the Hessian of the objective as \( H_k \approx (J_k^T J_k) \) and the gradient of the objective as \( \tilde{g}_k \approx -J_k r(x_k) \). The least-squares Jacobian, \( J \), should be provided to Tao using \texttt{TaoSetJacobianResidual()} routine.

The BRGN (-tao_type brgn) implementation adds a regularization term \( \beta(x) \) such that

\[ \min_x \frac{1}{2} \|R(x)\|_2^2 + \lambda \beta(x), \]

where \( \lambda \) is the scalar weight of the regularizer. BRGN provides two default implementations for \( \beta(x) \):
- L2-norm - $\beta(x) = \frac{1}{2}||x||^2$
- L2-norm Proximal Point - $\beta(x) = \frac{1}{2}||x - x_{k-1}||^2$
- L1-norm with Dictionary - $\beta(x) = \sum_i \sqrt{y_i^2 + \epsilon^2} - \epsilon$ where $y = Dx$ and $\epsilon$ is the smooth approximation parameter.

The regularizer weight can be controlled with either TaoBRGNSetRegularizerWeight() or -tao_brgn_regularizer_weight command line option, while the smooth approximation parameter can be set with either TaoBRGNSetL1SmoothEpsilon() or -tao_brgn_l1_smooth_epsilon. For the L1-norm term, the user can supply a dictionary matrix with TaoBRGNSetDictionaryMatrix(). If no dictionary is provided, the dictionary is assumed to be an identity matrix and the regularizer reduces to a sparse solution term.

The regularization selection can be made using the command line option -tao_brgn_regularization_type <l2pure, l2prox, l1dict, user> where the user option allows the user to define a custom $C^2$-continuous regularization term. This custom term can be defined by using the interface functions:

- TaoBRGNSetRegularizerObjectiveAndGradientRoutine() - Provide user-call back for evaluating the function value and gradient evaluation for the regularization term.
- TaoBRGNSetRegularizerHessianRoutine() - Provide user call-back for evaluating the Hessian of the regularization term.

POUNDERS

One algorithm for solving the least squares problem ((2.10)) when the Jacobian of the residual vector $F$ is unavailable is the model-based POUNDERS (Practical Optimization Using No Derivatives for sums of Squares) algorithm (tao_pounders). POUNDERS employs a derivative-free trust-region framework as described in [CSV09] in order to converge to local minimizers. An example of this version of POUNDERS applied to a practical least-squares problem can be found in [KortelainenLesinskiMore+10].

Derivative-Free Trust-Region Algorithm

In each iteration $k$, the algorithm maintains a model $m_k(x)$, described below, of the nonlinear least squares function $f$ centered about the current iterate $x_k$.

If one assumes that the maximum number of function evaluations has not been reached and that $||\nabla m_k(x_k)||_2 > gtol$, the next point $x_+$ to be evaluated is obtained by solving the trust-region subproblem

$$\min \{ m_k(x) : ||x - x_k||_p \leq \Delta_k, \}, \quad (2.11)$$

where $\Delta_k$ is the current trust-region radius. By default we use a trust-region norm with $p = \infty$ and solve ((2.11)) with the BLMVM method described in Bound-constrained Limited-Memory Variable-Metric Method (BLMVM). While the subproblem is a bound-constrained quadratic program, it may not be convex and the BQPPIP and GPCG methods may not solve the subproblem. Therefore, a bounded Newton-Krylov Method should be used; the default is the BNTR algorithm. Note: BNTR uses its own internal trust region that may interfere with the infinity-norm trust region used in the model problem ((2.11)).

The residual vector is then evaluated to obtain $F(x_+)$ and hence $f(x_+)$. The ratio of actual decrease to predicted decrease,

$$\rho_k = \frac{f(x_k) - f(x_+)}{m_k(x_k) - m_k(x_+)}.$$
as well as an indicator, valid, on the model’s quality of approximation on the trust region is then used to update the iterate,

\[
x_{k+1} = \begin{cases} 
  x_+ & \text{if } \rho_k \geq \eta_1 \\
  x_+ & \text{if } 0 < \rho_k < \eta_1 \text{ and } \text{valid} = \text{true} \\
  x_k & \text{else,}
\end{cases}
\]

and trust-region radius,

\[
\Delta_{k+1} = \begin{cases} 
  \min(\gamma_1 \Delta_k, \Delta_{\text{max}}) & \text{if } \rho_k \geq \eta_1 \text{ and } \|x_+ - x_k\|_p \geq \omega_1 \Delta_k \\
  \gamma_0 \Delta_k & \text{if } \rho_k < \eta_1 \text{ and } \text{valid} = \text{true} \\
  \Delta_k & \text{else,}
\end{cases}
\]

where \(0 < \eta_1 < 1, 0 < \gamma_0 < 1 < \gamma_1, 0 < \omega_1 < 1\), and \(\Delta_{\text{max}}\) are constants.

If \(\rho_k \leq 0\) and valid is false, the iterate and trust-region radius remain unchanged after the above updates, and the algorithm tests whether the direction \(x_+ - x_k\) improves the model. If not, the algorithm performs an additional evaluation to obtain \(F(x_k + d_k)\), where \(d_k\) is a model-improving direction.

The iteration counter is then updated, and the next model \(m_k\) is obtained as described next.

**Forming the Trust-Region Model**

In each iteration, POUNDERS uses a subset of the available evaluated residual vectors \(\{F(y_1), F(y_2), \cdots\}\) to form an interpolatory quadratic model of each residual component. The \(m\) quadratic models

\[
q_k(i)(x) = F_i(x_k) + (x - x_k)^T g_k(i) + \frac{1}{2} (x - x_k)^T H_k(i)(x - x_k), \quad i = 1, \ldots, m
\]

thus satisfy the interpolation conditions

\[
q_k(i)(y_j) = F_i(y_j), \quad i = 1, \ldots, m; \quad j = 1, \ldots, l_k
\]

on a common interpolation set \(\{y_1, \cdots, y_{l_k}\}\) of size \(l_k \in [n + 1, n_{\text{pmax}}]\).

The gradients and Hessians of the models in (eq:eq_models) are then used to construct the main model,

\[
m_k(x) = f(x_k) + 2(x - x_k)^T \sum_{i=1}^{m} F_i(x_k) g_k(i) + (x - x_k)^T \sum_{i=1}^{m} \left( g_k(i)^T g_k(i) + F_i(x_k) H_k(i) \right) (x - x_k).
\]

The process of forming these models also computes the indicator valid of the model’s local quality.

**Parameters**

POUNDERS supports the following parameters that can be set from the command line or PETSc options file:

- **-tao_pounders_delta <delta>**
  The initial trust-region radius \((> 0, \text{real})\). This is used to determine the size of the initial neighborhood within which the algorithm should look.

- **-tao_pounders_npmax <npmax>**
  The maximum number of interpolation points used \((n + 2 \leq \text{npmax} \leq 0.5(n + 1)(n + 2))\). This input is made available to advanced users. We recommend the default value \((\text{npmax} = 2n + 1)\) be used by others.

- **-tao_pounders_gqt**
  Use the gqt algorithm to solve the subproblem ((2.11)) (uses \(p = 2\)) instead of BQPIP.
If the default BQPIP algorithm is used to solve the subproblem (2.11), the parameters of the subproblem solver can be accessed using the command line options prefix -pounders_subsolver_. For example,

- pounders_subsolver_tao_gatol 1.0e-5

sets the gradient tolerance of the subproblem solver to $10^{-5}$.

Additionally, the user provides an initial solution vector, a vector for storing the separable objective function, and a routine for evaluating the residual vector $F$. These are described in detail in **Objective Function and Gradient Routines** and **Nonlinear Least Squares**. Here we remark that because gradient information is not available for scaling purposes, it can be useful to ensure that the problem is reasonably well scaled. A simple way to do so is to rescale the decision variables $x$ so that their typical values are expected to lie within the unit hypercube $[0, 1]^n$.

**Convergence Notes**

Because the gradient function is not provided to POUNDERS, the norm of the gradient of the objective function is not available. Therefore, for convergence criteria, this norm is approximated by the norm of the model gradient and used only when the model gradient is deemed to be a reasonable approximation of the gradient of the objective. In practice, the typical grounds for termination for expensive derivative-free problems is the maximum number of function evaluations allowed.

**Complementarity**

Mixed complementarity problems, or box-constrained variational inequalities, are related to nonlinear systems of equations. They are defined by a continuously differentiable function, $F : \mathbb{R}^n \rightarrow \mathbb{R}^n$, and bounds, $\ell \in \{\mathbb{R} \cup \{-\infty\}\}^n$ and $u \in \{\mathbb{R} \cup \{\infty\}\}^n$, on the variables such that $\ell \leq u$. Given this information, $x^* \in [\ell, u]$ is a solution to MCP($F, \ell, u$) if for each $i \in \{1, \ldots, n\}$ we have at least one of the following:

- $F_i(x^*) \geq 0$ if $x^*_i = \ell_i$
- $F_i(x^*) = 0$ if $\ell_i < x^*_i < u_i$
- $F_i(x^*) \leq 0$ if $x^*_i = u_i$.

Note that when $\ell = \{-\infty\}^n$ and $u = \{\infty\}^n$, we have a nonlinear system of equations, and $\ell = \{0\}^n$ and $u = \{\infty\}^n$ correspond to the nonlinear complementarity problem [Cot64].

Simple complementarity conditions arise from the first-order optimality conditions from optimization [Kar39] [KT51]. In the simple bound-constrained optimization case, these conditions correspond to MCP($\nabla f, \ell, u$), where $f : \mathbb{R}^n \rightarrow \mathbb{R}$ is the objective function. In a one-dimensional setting these conditions are intuitive. If the solution is at the lower bound, then the function must be increasing and $\nabla f \geq 0$. If the solution is at the upper bound, then the function must be decreasing and $\nabla f \leq 0$. If the solution is strictly between the bounds, we must be at a stationary point and $\nabla f = 0$. Other complementarity problems arise in economics and engineering [FP97], game theory [Nas50], and finance [HP98].

Evaluation routines for $F$ and its Jacobian must be supplied prior to solving the application. The bounds, $[\ell, u]$, on the variables must also be provided. If no starting point is supplied, a default starting point of all zeros is used.
Semismooth Methods

TAO has two implementations of semismooth algorithms [MFF+01] [DeLucaFK96] [FFK97] for solving mixed complementarity problems. Both are based on a reformulation of the mixed complementarity problem as a nonsmooth system of equations using the Fischer-Burmeister function [FiSh92]. A nonsmooth Newton method is applied to the reformulated system to calculate a solution. The theoretical properties of such methods are detailed in the aforementioned references.

The Fischer-Burmeister function, $\phi : \mathbb{R}^2 \to \mathbb{R}$, is defined as

$$\phi(a, b) := \sqrt{a^2 + b^2} - a - b.$$ 

This function has the following key property,

$$\phi(a, b) = 0 \Leftrightarrow a \geq 0, \ b \geq 0, \ ab = 0,$$

used when reformulating the mixed complementarity problem as the system of equations $\Phi(x) = 0$, where $\Phi : \mathbb{R}^n \to \mathbb{R}^n$. The reformulation is defined componentwise as

$$\Phi_i(x) := \begin{cases} 
\phi(x_i - l_i, F_i(x)) & \text{if } -\infty < l_i < u_i = \infty, \\
-\phi(u_i - x_i, -F_i(x)) & \text{if } -\infty = l_i < u_i < \infty, \\
\phi(x_i - l_i, \phi(u_i - x_i, -F_i(x))) & \text{if } -\infty < l_i < u_i < \infty, \\
-F_i(x) & \text{if } -\infty = l_i < u_i = \infty, \\
l_i - x_i & \text{if } -\infty < l_i = u_i < \infty.
\end{cases}$$

We note that $\Phi$ is not differentiable everywhere but satisfies a semismoothness property [Mif77] [Qi93] [QS93]. Furthermore, the natural merit function, $\Psi(x) := \frac{1}{2} \|\Phi(x)\|^2_2$, is continuously differentiable.

The two semismooth TAO solvers both solve the system $\Phi(x) = 0$ by applying a nonsmooth Newton method with a line search. We calculate a direction, $d^k$, by solving the system $H^k d^k = -\Phi(x^k)$, where $H^k$ is an element of the B-subdifferential [QS93] of $\Phi$ at $x^k$. If the direction calculated does not satisfy a suitable descent condition, then we use the negative gradient of the merit function, $-\nabla \Psi(x^k)$, as the search direction. A standard Armijo search [Arm66] is used to find the new iteration. Nonmonotone searches [GLL86] are also available by setting appropriate runtime options. See Line Searches for further details.

The first semismooth algorithm available in TAO is not guaranteed to remain feasible with respect to the bounds, $[l, u]$, and is termed an infeasible semismooth method. This method can be specified by using the `tao_ssils` solver. In this case, the descent test used is that

$$\nabla \Psi(x^k)^T d^k \leq -\delta \|d^k\|^\rho.$$ 

Both $\delta > 0$ and $\rho > 2$ can be modified by using the runtime options `-tao_ssils_delta <delta>` and `-tao_ssils_rho <rho>`, respectively. By default, $\delta = 10^{-10}$ and $\rho = 2.1$.

An alternative is to remain feasible with respect to the bounds by using a projected Armijo line search. This method can be specified by using the `tao_ssfls` solver. The descent test used is the same as above where the direction in this case corresponds to the first part of the piecewise linear arc searched by the projected line search. Both $\delta > 0$ and $\rho > 2$ can be modified by using the runtime options `-tao_ssfls_delta <delta>` and `-tao_ssfls_rho <rho>` respectively. By default, $\delta = 10^{-10}$ and $\rho = 2.1$.

The recommended algorithm is the infeasible semismooth method, `tao_ssils`, because of its strong global and local convergence properties. However, if it is known that $F$ is not defined outside of the box, $[l, u]$, perhaps because of the presence of log functions, the feasibility-enforcing version of the algorithm, `tao_ssfls`, is a reasonable alternative.
Active-Set Methods

TAO also contained two active-set semismooth methods for solving complementarity problems. These methods solve a reduced system constructed by block elimination of active constraints. The subdifferential in these cases enables this block elimination.

The first active-set semismooth algorithm available in TAO is not guaranteed to remain feasible with respect to the bounds, $[l, u]$, and is termed an infeasible active-set semismooth method. This method can be specified by using the `tao_asils` solver.

An alternative is to remain feasible with respect to the bounds by using a projected Armijo line search. This method can be specified by using the `tao_asfls` solver.

Quadratic Solvers

Quadratic solvers solve optimization problems of the form

$$
\min_x \frac{1}{2} x^T Q x + c^T x
$$

subject to $l \geq x \geq u$

where the gradient and the Hessian of the objective are both constant.

Gradient Projection Conjugate Gradient Method (GPCG)

The GPCG [MoreT91] algorithm is much like the TRON algorithm, discussed in Section Trust-Region Newton Method (TRON), except that it assumes that the objective function is quadratic and convex. Therefore, it evaluates the function, gradient, and Hessian only once. Since the objective function is quadratic, the algorithm does not use a trust region. All the options that apply to TRON except for trust-region options also apply to GPCG. It can be set by using the TAO solver `tao_gpcg` or via the option flag `-tao_type gpcg`.

Interior-Point Newton’s Method (BQPIP)

The BQPIP algorithm is an interior-point method for bound constrained quadratic optimization. It can be set by using the TAO solver of `tao_bqpip` or via the option flag `-tao_type bqpip`. Since it assumes the objective function is quadratic, it evaluates the function, gradient, and Hessian only once. This method also requires the solution of systems of linear equations, whose solver can be accessed and modified with the command `TaoGetKSP()`.

Legacy and Contributed Solvers

Bundle Method for Regularized Risk Minimization (BMRM)

BMRM is a numerical approach to optimizing an unconstrained objective in the form of $f(x) + 0.5 \times \lambda \|x\|^2$. Here $f$ is a convex function that is finite on the whole space. $\lambda$ is a positive weight parameter, and $\|x\|$ is the Euclidean norm of $x$. The algorithm only requires a routine which, given an $x$, returns the value of $f(x)$ and the gradient of $f$ at $x$. 
Orthant-Wise Limited-memory Quasi-Newton (OWLQN)

OWLQN \cite{AG07} is a numerical approach to optimizing an unconstrained objective in the form of \(f(x) + \lambda \|x\|_1\). Here \(f\) is a convex and differentiable function, \(\lambda\) is a positive weight parameter, and \(\|x\|_1\) is the \(\ell_1\) norm of \(x: \sum_i |x_i|\). The algorithm only requires evaluating the value of \(f\) and its gradient.

Trust-Region Newton Method (TRON)

The TRON \cite{LM099} algorithm is an active-set method that uses a combination of gradient projections and a preconditioned conjugate gradient method to minimize an objective function. Each iteration of the TRON algorithm requires function, gradient, and Hessian evaluations. In each iteration, the algorithm first applies several conjugate gradient iterations. After these iterates, the TRON solver momentarily ignores the variables that equal one of its bounds and applies a preconditioned conjugate gradient method to a quadratic model of the remaining set of free variables.

The TRON algorithm solves a reduced linear system defined by the rows and columns corresponding to the variables that lie between the upper and lower bounds. The TRON algorithm applies a trust region to the conjugate gradients to ensure convergence. The initial trust-region radius can be set by using the command \texttt{TaoSetInitialTrustRegionRadius()}, and the current trust region size can be found by using the command \texttt{TaoGetCurrentTrustRegionRadius()}. The initial trust region can significantly alter the rate of convergence for the algorithm and should be tuned and adjusted for optimal performance.

This algorithm will be deprecated in the next version in favor of the Bounded Newton Trust Region (BNTR) algorithm.

Bound-constrained Limited-Memory Variable-Metric Method (BLMVM)

BLMVM is a limited-memory, variable-metric method and is the bound-constrained variant of the LMVM method for unconstrained optimization. It uses projected gradients to approximate the Hessian, eliminating the need for Hessian evaluations. The method can be set by using the TAO solver \texttt{tao_blmvm}. For more details, please see the LMVM section in the unconstrained algorithms as well as the LMVM matrix documentation in the PETSc manual.

This algorithm will be deprecated in the next version in favor of the Bounded Quasi-Newton Line Search (BQNLS) algorithm.

2.7.4 Advanced Options

This section discusses options and routines that apply to most TAO solvers and problem classes. In particular, we focus on linear solvers, convergence tests, and line searches.

Linear Solvers

One of the most computationally intensive phases of many optimization algorithms involves the solution of linear systems of equations. The performance of the linear solver may be critical to an efficient computation of the solution. Since linear equation solvers often have a wide variety of options associated with them, TAO allows the user to access the linear solver with the

\begin{verbatim}
TaoGetKSP(Tao, KSP *)
\end{verbatim}

command. With access to the KSP object, users can customize it for their application to achieve improved performance. Additional details on the KSP options in PETSc can be found in the \textit{User-Guide}.
**Monitors**

By default the TAO solvers run silently without displaying information about the iterations. The user can initiate monitoring with the command

```c
TaoSetMonitor(Tao, PetscErrorCode (*mon)(Tao, void*), void*);
```

The routine `mon` indicates a user-defined monitoring routine, and `void*` denotes an optional user-defined context for private data for the monitor routine.

The routine set by `TaoSetMonitor()` is called once during each iteration of the optimization solver. Hence, the user can employ this routine for any application-specific computations that should be done after the solution update.

**Convergence Tests**

Convergence of a solver can be defined in many ways. The methods TAO uses by default are mentioned in *Convergence*. These methods include absolute and relative convergence tolerances as well as a maximum number of iterations of function evaluations. If these choices are not sufficient, the user can specify a customized test.

Users can set their own customized convergence tests of the form

```c
PetscErrorCode conv(Tao, void*);
```

The second argument is a pointer to a structure defined by the user. Within this routine, the solver can be queried for the solution vector, gradient vector, or other statistic at the current iteration through routines such as `TaoGetSolutionStatus()` and `TaoGetTolerances()`.

To use this convergence test within a TAO solver, one uses the command

```c
TaoSetConvergenceTest(Tao, PetscErrorCode (*conv)(Tao, void*), void*);
```

The second argument of this command is the convergence routine, and the final argument of the convergence test routine denotes an optional user-defined context for private data. The convergence routine receives the TAO solver and this private data structure. The termination flag can be set by using the routine

```c
TaoSetConvergedReason(Tao, TaoConvergedReason);`
```

**Line Searches**

By using the command line option `-tao_ls_type`. Available line searches include Moré-Thuente [MoreT92], Armijo, gpcg, and unit.

The line search routines involve several parameters, which are set to defaults that are reasonable for many applications. The user can override the defaults by using the following options

- `-tao_ls_max_funcs <max>`
- `-tao_ls_stepmin <min>`
- `-tao_ls_stepmax <max>`
- `-tao_ls_ftol <ftol>`
- `-tao_ls_gtol <gtol>`
- `-tao_ls_rtol <rtol>`

---

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One should run a TAO program with the option `-help` for details. Users may write their own customized line search codes by modeling them after one of the defaults provided.

**Recycling History**

Some TAO algorithms can re-use information accumulated in the previous `TaoSolve()` call to hot-start the new solution. This can be enabled using the `-tao_recycle_history` flag, or in code via the `TaoSetRecycleHistory()` interface.

For the nonlinear conjugate gradient solver (`TAOBNCG`), this option re-uses the latest search direction from the previous `TaoSolve()` call to compute the initial search direction of a new `TaoSolve()`. By default, the feature is disabled and the algorithm sets the initial direction as the negative gradient.

For the quasi-Newton family of methods (`TAOBQNLS`, `TAOBQNKL`, `TAOBQNKR`, `TAOBQNKT`), this option re-uses the accumulated quasi-Newton Hessian approximation from the previous `TaoSolve()` call. By default, the feature is disabled and the algorithm will reset the quasi-Newton approximation to the identity matrix at the beginning of every new `TaoSolve()`.

The option flag has no effect on other TAO solvers.

### 2.7.5 Adding a Solver

One of the strengths of both TAO and PETSc is the ability to allow users to extend the built-in solvers with new user-defined algorithms. It is certainly possible to develop new optimization algorithms outside of TAO framework, but Using TAO to implement a solver has many advantages,

1. TAO includes other optimization solvers with an identical interface, so application problems may conveniently switch solvers to compare their effectiveness.

2. TAO provides support for function evaluations and derivative information. It allows for the direct evaluation of this information by the application developer, contains limited support for finite difference approximations, and allows the uses of matrix-free methods. The solvers can obtain this function and derivative information through a simple interface while the details of its computation are handled within the toolkit.

3. TAO provides line searches, convergence tests, monitoring routines, and other tools that are helpful in an optimization algorithm. The availability of these tools means that the developers of the optimization solver do not have to write these utilities.

4. PETSc offers vectors, matrices, index sets, and linear solvers that can be used by the solver. These objects are standard mathematical constructions that have many different implementations. The objects may be distributed over multiple processors, restricted to a single processor, have a dense representation, use a sparse data structure, or vary in many other ways. TAO solvers do not need to know how these objects are represented or how the operations defined on them have been implemented. Instead, the solvers apply these operations through an abstract interface that leaves the details to PETSc and external libraries. This abstraction allows solvers to work seamlessly with a variety of data structures while allowing application developers to select data structures tailored for their purposes.

5. PETSc provides the user a convenient method for setting options at runtime, performance profiling, and debugging.
Header File

TAO solver implementation files must include the TAO implementation file `taoimpl.h`:

```c
#include "petsc/private/taoimpl.h"
```

This file contains data elements that are generally kept hidden from application programmers, but may be necessary for solver implementations to access.

TAO Interface with Solvers

TAO solvers must be written in C or C++ and include several routines with a particular calling sequence. Two of these routines are mandatory: one that initializes the TAO structure with the appropriate information and one that applies the algorithm to a problem instance. Additional routines may be written to set options within the solver, view the solver, setup appropriate data structures, and destroy these data structures. In order to implement the conjugate gradient algorithm, for example, the following structure is useful.

```c
typedef struct{
    PetscReal beta;
    PetscReal eta;
    PetscInt ngradtseps;
    PetscInt nresetsteps;
    Vec X_old;
    Vec G_old;
} TAO_CG;
```

This structure contains two parameters, two counters, and two work vectors. Vectors for the solution and gradient are not needed here because the TAO structure has pointers to them.

Solver Routine

All TAO solvers have a routine that accepts a TAO structure and computes a solution. TAO will call this routine when the application program uses the routine `TaoSolve()` and will pass to the solver information about the objective function and constraints, pointers to the variable vector and gradient vector, and support for line searches, linear solvers, and convergence monitoring. As an example, consider the following code that solves an unconstrained minimization problem using the conjugate gradient method.

```c
PetscErrorCode TaoSolve_CG(Tao tao)
{
    TAO_CG *cg = (TAO_CG *) tao->data;
    Vec x = tao->solution;
    Vec g = tao->gradient;
    Vec s = tao->stepdirection;
    PetscInt iter=0;
    PetscReal gnormPrev,gdx,f,gnorm,steplength=0;
    TaoLineSearchConvergedReason lsflag=TAO_LINESEARCH_CONTINUE_ITERATING;
    TaoConvergedReason reason=TAO_CONTINUE_ITERATING;

    PetscFunctionBegin;
    PetscCall(TaoComputeObjectiveAndGradient(tao,x,&f,g));
    PetscCall(VecNorm(g,NORM_2,&gnorm));

    (continues on next page)
```

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PetscCall(VecSet(s, 0));

// Enter loop */
while (1){

/* Test for convergence */
PetscCall(TaoMonitor(tao, iter, f, gnorm, 0.0, step, &reason));
if (reason != TAO_CONTINUE_ITERATING) break;

cg->beta = (gnorm*gnorm)/(gnormPrev*gnormPrev);
PetscCall(VecScale(s, cg->beta));
PetscCall(VecAXPY(s, -1.0, g));

PetscCall(VecDot(s, g, &gdx));
if (gdx > 0){ /* If not a descent direction, use gradient */
  PetscCall(VecCopy(g, s));
  PetscCall(VecScale(s, -1.0));
  gdx = -gnorm*gnorm;
}

/* Line Search */
gnormPrev = gnorm; step = 1.0;
PetscCall(TaoLineSearchSetInitialStepLength(tao->linesearch, 1.0));
PetscCall(TaoLineSearchApply(tao->linesearch, x, &f, &g, &steplength, &lsflag));
PetscCall(TaoAddLineSearchCounts(tao));
PetscCall(VecNorm(g, NORM_2, &gnorm));
iter++;
}
PetscFunctionReturn(PETSC_SUCCESS);
}

The first line of this routine casts the second argument to a pointer to a TAO_CG data structure. This structure contains pointers to three vectors and a scalar that will be needed in the algorithm.

After declaring an initializing several variables, the solver lets TAO evaluate the function and gradient at the current point in the using the routine TaoComputeObjectiveAndGradient(). Other routines may be used to evaluate the Hessian matrix or evaluate constraints. TAO may obtain this information using direct evaluation or other means, but these details do not affect our implementation of the algorithm.

The norm of the gradient is a standard measure used by unconstrained minimization solvers to define convergence. This quantity is always nonnegative and equals zero at the solution. The solver will pass this quantity, the current function value, the current iteration number, and a measure of infeasibility to TAO with the routine

```
PetscErrorCode TaoMonitor(Tao tao, PetscInt iter, PetscReal f,
  PetscReal res, PetscReal cnorm, PetscReal steplength,
  TaoConvergedReason *reason);
```

Most optimization algorithms are iterative, and solvers should include this command somewhere in each iteration. This routine records this information, and applies any monitoring routines and convergence tests set by default or the user. In this routine, the second argument is the current iteration number, and the third argument is the current function value. The fourth argument is a nonnegative error measure associated
with the distance between the current solution and the optimal solution. Examples of this measure are the norm of the gradient or the square root of a duality gap. The fifth argument is a nonnegative error that usually represents a measure of the infeasibility such as the norm of the constraints or violation of bounds. This number should be zero for unconstrained solvers. The sixth argument is a nonnegative steplength, or the multiple of the step direction added to the previous iterate. The results of the convergence test are returned in the last argument. If the termination reason is TAO_CONTINUE_ITERATING, the algorithm should continue.

After this monitoring routine, the solver computes a step direction using the conjugate gradient algorithm and computations using Vec objects. These methods include adding vectors together and computing an inner product. A full list of these methods can be found in the manual pages.

Nonlinear conjugate gradient algorithms also require a line search. TAO provides several line searches and support for them. The routine

```c
TaoLineSearchApply(TaoLineSearch ls, Vec x, PetscReal *f, Vec g,
                   TaoVec *s, PetscReal *steplength,
                   TaoLineSearchConvergedReason *lsflag)
```

passes the current solution, gradient, and objective value to the line search and returns a new solution, gradient, and objective value. More details on line searches can be found in Line Searches. The details of the line search applied are specified elsewhere, when the line search is created.

TAO also includes support for linear solvers using PETSc KSP objects. Although this algorithm does not require one, linear solvers are an important part of many algorithms. Details on the use of these solvers can be found in the PETSc users manual.

### Creation Routine

The TAO solver is initialized for a particular algorithm in a separate routine. This routine sets default convergence tolerances, creates a line search or linear solver if needed, and creates structures needed by this solver. For example, the routine that creates the nonlinear conjugate gradient algorithm shown above can be implemented as follows.

```c
PETSC_EXTERN PetscErrorCode TaoCreate_CG(Tao tao)
{
    TAO_CG *cg = (TAO_CG*)tao->data;
    const char *morethuente_type = TAOLINESEARCH_MT;

    PetscFunctionBegin;

    PetscCall(PetscNew(&cg));
    tao->data = (void*)cg;
    cg->eta = 0.1;
    cg->delta_min = 1e-7;
    cg->delta_max = 100;
    cg->cg_type = CG_PolakRibierePlus;

    tao->max_it = 2000;
    tao->max_funcs = 4000;

    tao->ops->setup = TaoSetUp_CG;
    tao->ops->solve = TaoSolve_CG;
    tao->ops->view = TaoView_CG;
    tao->ops->setfromoptions = TaoSetFromOptions_CG;
    tao->ops->destroy = TaoDestroy_CG;

    (continues on next page)
}
This routine declares some variables and then allocates memory for the TAO_CG data structure. Notice that the Tao object now has a pointer to this data structure (tao->data) so it can be accessed by the other functions written for this solver implementation.

This routine also sets some default parameters particular to the conjugate gradient algorithm, sets default convergence tolerances, and creates a particular line search. These defaults could be specified in the routine that solves the problem, but specifying them here gives the user the opportunity to modify these parameters either by using direct calls setting parameters or by using options.

Finally, this solver passes to TAO the names of all the other routines used by the solver.

Note that the lines EXTERN_C_BEGIN and EXTERN_C_END surround this routine. These macros are required to preserve the name of this function without any name-mangling from the C++ compiler (if used).

### Destroy Routine

Another routine needed by most solvers destroys the data structures created by earlier routines. For the nonlinear conjugate gradient method discussed earlier, the following routine destroys the two work vectors and the TAO_CG structure.

```c
PetscErrorCode TaoDestroy_CG(TAO_SOLVER tao) {
    TAO_CG *cg = (TAO_CG *) tao->data;
    PetscFunctionBegin;
    PetscCall(VecDestroy(&cg->X_old));
    PetscCall(VecDestroy(&cg->G_old));
    PetscFree(tao->data);
    tao->data = NULL;
    PetscFunctionReturn(PETSC_SUCCESS);
}
```

This routine is called from within the TaoDestroy() routine. Only algorithm-specific data objects are destroyed in this routine; any objects indexed by TAO (tao->linesearch, tao->ksp, tao->gradient, etc.) will be destroyed by TAO immediately after the algorithm-specific destroy routine completes.
**SetUp Routine**

If the SetUp routine has been set by the initialization routine, TAO will call it during the execution of `TaoSolve()`. While this routine is optional, it is often provided to allocate the gradient vector, work vectors, and other data structures required by the solver. It should have the following form.

```c
PetscErrorCode TaoSetUp_CG(Tao tao) {
    TAO_CG *cg = (TAO_CG*)tao->data;
    PetscFunctionBegin;
    PetscCall(VecDuplicate(tao->solution,&tao->gradient));
    PetscCall(VecDuplicate(tao->solution,&tao->stepdirection));
    PetscCall(VecDuplicate(tao->solution,&cg->X_old));
    PetscCall(VecDuplicate(tao->solution,&cg->G_old));
    PetscFunctionReturn(PETSC_SUCCESS);
}
```

**SetFromOptions Routine**

The SetFromOptions routine should be used to check for any algorithm-specific options set by the user and will be called when the application makes a call to `TaoSetFromOptions()`. It should have the following form.

```c
PetscErrorCode TaoSetFromOptions_CG(Tao tao, void *solver) {
    TAO_CG *cg = (TAO_CG*)solver;
    PetscFunctionBegin;
    PetscCall(PetscOptionsReal("-tao_cg_eta", "restart tolerance", "", cg->eta,&cg->eta, 0));
    PetscCall(PetscOptionsReal("-tao_cg_delta_min", "minimum delta value", "", cg->delta_min,&cg->delta_min, 0));
    PetscCall(PetscOptionsReal("-tao_cg_delta_max", "maximum delta value", "", cg->delta_max,&cg->delta_max, 0));
    PetscFunctionReturn(PETSC_SUCCESS);
}
```

**View Routine**

The View routine should be used to output any algorithm-specific information or statistics at the end of a solve. This routine will be called when the application makes a call to `TaoView()` or when the command line option `-tao_view` is used. It should have the following form.

```c
PetscErrorCode TaoView_CG(Tao tao, PetscViewer viewer) {
    TAO_CG *cg = (TAO_CG*)tao->data;
    PetscFunctionBegin;
    PetscCall(PetscViewerASCIIPushTab(viewer));
    PetscCall(PetscViewerASCIIPrintf(viewer,"Grad. steps: %d\n",cg->ngradsteps));
    PetscCall(PetscViewerASCIIPrintf(viewer,"Reset steps: %d\n",cg->nresetsteps));
    PetscCall(PetscViewerASCIIPopTab(viewer));
}
```

(continues on next page)
Registering the Solver

Once a new solver is implemented, TAO needs to know the name of the solver and what function to use to create the solver. To this end, one can use the routine

```c
TaoRegister(const char *name,
            const char *path,
            const char *cname,
            PetscErrorCode (*create)(Tao));
```

where `name` is the name of the solver (i.e., `tao_blmvm`), `path` is the path to the library containing the solver, `cname` is the name of the routine that creates the solver (in our case, `TaoCreate_CG`), and `create` is a pointer to that creation routine. If one is using dynamic loading, then the fourth argument will be ignored.

Once the solver has been registered, the new solver can be selected either by using the `TaoSetType()` function or by using the `-tao_type` command line option.
DM: INTERFACING BETWEEN SOLVERS AND MODELS/DISCRETIZATIONS

3.1 DM Basics

The previous chapters have focused on the core numerical solvers in PETSc. However, numerical solvers without efficient ways (in both human and machine time) of connecting the solvers to the mathematical models and discretizations that people wish to build their simulations on, will not get widely used. Thus PETSc provides a set of abstractions represented by the DM object to provide a powerful, comprehensive mechanism for translating the problem specification of a model and its discretization to the language and API of solvers.

Some of the model classes DM currently supports are PDEs on structured and staggered grids with finite difference methods (DMDA and DMSTAG – DMSTAG: Staggered, Structured Grid), PDEs on unstructured grids with finite element and finite volume methods (DMPLEX – DMplex: Unstructured Grids), PDEs on quad and octree-grids (DMFOREST), models on networks (graphs) such as the power grid or river networks (DMNETWORK – Networks), and particle-in-cell simulations (DMSWARm).

In previous chapters, we have demonstrated some simple usage of DM to provide the input for the solvers. In this chapter, and those that follow, we will dive deep into the capabilities of DM.

It is possible to create a DM with

```c
DM dm;
DMCreate(MPI_Comm comm, DM *dm);
DMSetType(DM dm, DMType type);
```

but more commonly, a DM is created with a type-specific constructor; the construction process for each type of DM is discussed in the sections on each DMType. This chapter focuses on commonalities between all the DM so we assume the DM already exists and we wish to work with it.

As discussed earlier, a DM can construct vectors and matrices appropriate for a model and discretization and provide the mapping between the global and local vector representations.

```c
DMCreateLocalVector(DM dm,Vec *l);
DMCreateGlobalVector(DM dm,Vec *g);
DMGlobalToLocal(dm,g,l,INSERT_VALUES);
DMLocalToGlobal(dm,l,g,ADD_VALUES);
DMCreateMatrix(dm,Mat *m);
```

The matrices produced may support MatSetValuesLocal() allowing one to work with the local numbering on each MPI rank. For DMDA one can also use MatSetValuesStencil() and for DMSTAG with DMStagMatSetValuesStencil().
A given DM can be refined for certain DMTypes with DMRefine() or coarsened with DMCoarsen(). Mappings between DMs may be obtained with routines such as DMCreateInterpolation(), DMCreateRestriction() and DMCreateInjection().

One attaches a DM to a solver object with

```c
KSPSetDM(KSP ksp, DM dm);
SNESetDM(SNES snes, DM dm);
TSSetDM(TS ts, DM dm);
```

Once the DM is attached, the solver can utilize it to create and process much of the data that the solver needs to set up and implement its solve. For example, with PCMG simply providing a DM can allow it to create all the data structures needed to run geometric multigrid on your problem.

SNES Tutorial ex19 demonstrates how this may be done with DMDA.

### 3.2 DMPIlex: Unstructured Grids

This chapter introduces the DMPILEX subclass of DM, which allows the user to handle unstructured grids using the generic DM interface for hierarchy and multi-physics. DMPILEX was created to remedy a huge problem in all current PDE simulation codes, namely that the discretization was so closely tied to the data layout and solver that switching discretizations in the same code was not possible. Not only does this preclude the kind of comparison that is necessary for scientific investigation, but it makes library (as opposed to monolithic application) development impossible.

#### 3.2.1 Representing Unstructured Grids

The main advantage of DMPILEX in representing topology is that it treats all the different pieces of a mesh, e.g., cells, faces, edges, and vertices, in the same way. This allows the interface to be small and simple, while remaining flexible and general. This also allows “dimension independent programming”, which means that the same algorithm can be used unchanged for meshes of different shapes and dimensions.

All pieces of the mesh (vertices, edges, faces, and cells) are treated as points, which are each identified by a PetscInt. A mesh is built by relating points to other points, in particular specifying a “covering” relation among the points. For example, an edge is defined by being covered by two vertices, and a triangle can be defined by being covered by three edges (or even by three vertices). This structure is known as a Hasse Diagram, which is a Directed Acyclic Graph (DAG) representing a cell complex using the covering relation. The graph edges represent the relation, which also encodes a partially ordered set (poset).

For example, we can encode the doublet mesh as in Fig. 3.1, which can also be represented as the DAG in Fig. 3.2.

To use the PETSc API, we consecutively number the mesh pieces. The PETSc convention in 3 dimensions is to number first cells, then vertices, then faces, and then edges. In 2 dimensions the convention is to number in terms of the labels in Fig. 3.1, these numberings are

\[
\begin{align*}
f_0 \mapsto 0, f_1 \mapsto 1, \\
v_0 \mapsto 2, v_1 \mapsto 3, v_2 \mapsto 4, v_3 \mapsto 5, \\
e_0 \mapsto 6, e_1 \mapsto 7, e_2 \mapsto 8, e_3 \mapsto 9, e_4 \mapsto 10
\end{align*}
\]

First, we declare the set of points present in a mesh,

```c
DMPIlexSetChart(dm, 0, 11);
```
Fig. 3.1: A 2D doublet mesh, two triangles sharing an edge.

Fig. 3.2: The Hasse diagram for our 2D doublet mesh, expressed as a DAG.
Note that a *chart* here corresponds to a semi-closed interval (e.g. \([0, 11) = \{0, 1, \ldots, 10\}\)) specifying the range of indices we’d like to use to define points on the current rank. We then define the covering relation, which we call the *cone*, which are also the in-edges in the DAG. In order to preallocate correctly, we first provide sizes,

```c
/* DMPlexSetConeSize(dm, point, number of points that cover the point); */
DMPlexSetConeSize(dm, 0, 3);
DMPlexSetConeSize(dm, 1, 3);
DMPlexSetConeSize(dm, 6, 2);
DMPlexSetConeSize(dm, 7, 2);
DMPlexSetConeSize(dm, 8, 2);
DMPlexSetConeSize(dm, 9, 2);
DMPlexSetConeSize(dm, 10, 2);
DMSSetUp(dm);
```

and then point values (recall each point is an integer that represents a single geometric entity, a cell, face, edge, or vertex),

```c
/* DMPlexSetCone(dm, point, [points that cover the point]); */
DMPlexSetCone(dm, 0, [6, 7, 8]);
DMPlexSetCone(dm, 1, [7, 9, 10]);
DMPlexSetCone(dm, 6, [2, 3]);
DMPlexSetCone(dm, 7, [3, 4]);
DMPlexSetCone(dm, 8, [4, 2]);
DMPlexSetCone(dm, 9, [4, 5]);
DMPlexSetCone(dm, 10, [5, 3]);
```

There is also an API for providing the dual relation, using `DMPlexSetSupportSize()` and `DMPlexSetSupport()`, but this can be calculated automatically using the provided `DMPlexSetConeSize()` and `DMPlexSetCone()` information and then calling

```c
DMPlexSymmetrize(dm);
```

The “symmetrization” is in the sense of the DAG. Each point knows its covering (cone) and each point knows what it covers (support). Note that when using automatic symmetrization, cones will be ordered but supports will not. The user can enforce an ordering on supports by rewriting them after symmetrization using `DMPlexSetSupport()`. In order to support efficient queries, we construct fast search structures and indices for the different types of points using

```c
DMPlexStratify(dm);
```

### 3.2.2 Dealing with Periodicity

Plex allows you to represent periodic domains in two ways. Using the default scheme, periodic topology can be represented directly. This ensures that all topological queries can be satisfied, but then care must be taken in representing functions over the mesh, such as the coordinates. The second method is to use a non-periodic topology, but connect certain mesh points using the local-to-global map for that DM. This allows a more general set of mappings to be implemented, such as partial twists, but topological queries on the periodic boundary cease to function.

For the default scheme, a call to `DMLocalizeCoordinates()` (which usually happens automatically on mesh creation) creates a second, discontinuous coordinate field. These values can be accessed using `DMGetCellCoordinates()` and `DMGetCellCoordinatesLocal()`. Plex provides a convenience method, `DMPlexGetCellCo-`
ordinates(), that extracts cell coordinates correctly, depending on the periodicity of the mesh. An example of its use is shown below:

```c
const PetscScalar *array;
PetscScalar *coords = NULL;
PetscInt numCoords;
PetscBool isDG;
PetscCall(DMPlexGetCellCoordinates(dm, cell, &isDG, &numCoords, &array, &coords));
for (PetscInt cc = 0; cc < numCoords / dim; ++cc) {
    if (cc > 0) PetscCall(PetscPrintf(PETSC_COMM_SELF, " -- 
    PetscCall(PetscPrintf(PETSC_COMM_SELF, "("));
    for (PetscInt d = 0; d < dim; ++d) {
        if (d > 0) PetscCall(PetscPrintf(PETSC_COMM_SELF, ", 
        PetscCall(PetscPrintf(PETSC_COMM_SELF, "%g", (double)PetscRealPart(coords[cc * dim + d])));
    }
    PetscCall(PetscPrintf(PETSC_COMM_SELF, ")
    PetscCall(DMPlexRestoreCellCoordinates(dm, cell, &isDG, &numCoords, &array, &coords));
```

### 3.2.3 Connecting Data on Grids to its Location in arrays or Vec (PetscSection)

The strongest links between solvers and discretizations are

- the relationship between the layout of data (unknowns) over a mesh (or similar structure) and the data layout in arrays and Vec used for computation,
- data (unknowns) partitioning, and
- ordering of data (unknowns).

To enable modularity, we encode the operations above in simple data structures that can be understood by the linear algebra (Vec, Mat, KSP, PC, SNES), time integrator (TS), and optimization (Tao) engines in PETSc without explicit reference to the mesh (topology) or discretization (analysis).

#### Data Layout by Hand

Specific entries (or collections of entries) in a Vec (or a simple array) can be associated with a “location” on a mesh (or other types of data structure) using the PetscSection object. A point is a PetscInt that serves as an abstract “index” into arrays from iterable sets, such as points on a mesh.

**PetscSection** has two modes of operation.

Mode 1:

A PetscSection associates a set of degrees of freedom (dof), (a small space \( \{ e_k \} | 0 < k < d_p \) ), with every point. The number of dof and their meaning may be different for different points. For example, the dof on a cell point may represent pressure while a dof on a face point may represent velocity. Though points must be contiguously numbered, they can be in any range \([ pStart, pEnd ]\), which is called a chart. A PetscSection in mode 1 may be thought of as defining a two dimensional array indexed by point in the outer dimension with a variable length inner dimension indexed by the dof at that point, \( v[pStart <= point < pEnd][0 <= dof < d_p]^1 \).

---

1 A PetscSection can be thought of as a generalization of PetscLayout, in the same way that a fiber bundle is a
The sequence for constructing a `PetscSection` in mode 1 is the following:

1. Specify the range of points, or chart, with `PetscSectionSetChart()`.
2. Specify the number of dofs per point, with `PetscSectionSetDof()`. Any values not set will be zero.
3. Set up the `PetscSection` with `PetscSectionSetUp()`.

Below we demonstrate such a process used by DMPLEX but first we introduce the second mode for working with `PetscSection`.

Mode 2:

A `PetscSection` consists of one more fields each of which is represented (internally) by a `PetscSection`. A `PetscSection` in mode 2 may be thought of as defining a three dimensional array indexed by point and field in the outer dimensions with a variable length inner dimension indexed by the dof at that point. The actual order the values in the array are stored can be set with `PetscSectionSetPointMajor(PetscSection, PETSC_TRUE, PETSC_FALSE)`. In point major order all the degrees of freedom for each point for all fields are stored contiguously, otherwise all degrees of freedom for each field are stored are stored contiguously. With point major order the fields are said to be interlaced.

Consider a `PetscSection` with 2 fields and 3 points (from 0 to 2) with 1 dof for each point. In point major order the array has the storage (values for all the fields at point 0, values for all the fields at point 1, values for all the fields at point 2) while in field major order it is (values for all points in field 0, values for all points in field 1).

The sequence for constructing such a `PetscSection` is the following:

1. Specify the range of points, or chart, with `PetscSectionSetChart()`.
   All fields share the same chart.
2. Specify the number of fields with `PetscSectionSetNumFields()`.
3. Optionally provide a name for the fields with `PetscSectionSetFieldName()`.
4. Set the number of dof for each point on each field with `PetscSectionSetFieldDof()`. Again, values not set will be zero.
5. Set the total number of dof for each point with `PetscSectionSetDof()`. Thus value must be greater than or equal to the sum of the values set with `PetscSectionSetFieldDof()` at that point. Again, values not set will be zero.
6. Set up the `PetscSection` with `PetscSectionSetUp()`.

Once a `PetscSection` has been created one can use `PetscSectionGetStorageSize(PetscSection, PetscInt *)` to determine the total number of entries that can be stored in an array or `Vec` accessible by the `PetscSection`. The memory locations in the associated array are found using an offset which can be obtained with:

Mode 1:

```c
PetscSectionGetOffset(PetscSection, PetscInt point, PetscInt &offset);
```

Mode 2:

```c
PetscSectionGetFieldOffset(PetscSection, PetscInt point, PetscInt field, PetscInt &offset);
```

generalization of the normal Euclidean basis used in linear algebra. With `PetscLayout`, we associate a unit vector (e_i) with every point in the space, and just divide up points between processes.
The value in the array is then accessed with `array[offset]`. If there are multiple dof at a point (and field in mode 2) then `array[offset + 1]`, etc give access to each of those dof.

Using the mesh from Fig. 3.1, we provide an example of creating a PetscSection using mode 1. We can lay out data for a continuous Galerkin $P_3$ finite element method,

```c
PetscInt pStart, pEnd, cStart, cEnd, c, vStart, vEnd, v, eStart, eEnd, e;

DMPlexGetChart(dm, &pStart, &pEnd);
DMPlexGetHeightStratum(dm, 0, &cStart, &cEnd); // cells
DMPlexGetHeightStratum(dm, 1, &eStart, &eEnd); // edges
DMPlexGetHeightStratum(dm, 2, &vStart, &vEnd); // vertices, equivalent to,
    DMPlexGetDepthStratum(dm, 0, &vStart, &vEnd);
PetscSectionSetChart(s, pStart, pEnd);
for (c = cStart; c < cEnd; ++c)
    PetscSectionSetDof(s, c, 1);
for (v = vStart; v < vEnd; ++v)
    PetscSectionSetDof(s, v, 1);
for (e = eStart; e < eEnd; ++e)
    PetscSectionSetDof(s, e, 2); // two dof on each edge
PetcSectionSetUp(s);
```

DMPlexGetHeightStratum() returns all the points of the requested height in the DAG. Since this problem is in two dimensions the edges are at height 1 and the vertices at height 2 (the cells are always at height 0). One can also use DMPlexGetDepthStratum() to use the depth in the DAG to select the points. DMPlexGetDepth(dm,&depth) returns the depth of the DAG, hence DMPlexGetDepthStratum(dm, depth-1-h,) returns the same values as DMPlexGetHeightStratum(dm,h,).

For $P_3$ elements there is one degree of freedom at each vertex, 2 along each edge (resulting in a total of 4 degrees of freedom along each edge including the vertices, thus being able to reproduce a cubic function) and 1 degree of freedom within the cell (the bubble function which is zero along all edges).

Now a PETSc local vector can be created manually using this layout,

```c
PetscSectionGetStorageSize(s, &n);
VecSetSizes(localVec, n, PETSC_DETERMINE);
VecSetFromOptions(localVec);
```

When working with DMPLEX and PetscFE (see below) one can simply get the sections (and related vectors) with

```c
DMSetLocalSection(dm, s);
DMGetLocalVector(dm, &localVec);
DMGetGlobalVector(dm, &globalVec);
```

A global vector is missing both the shared dofs which are not owned by this process, as well as constrained dofs. These constraints represent essential (Dirichlet) boundary conditions. They are dofs that have a given fixed value, so they are present in local vectors for assembly purposes, but absent from global vectors since they are never solved for during algebraic solves.

We can indicate constraints in a local section using PetscSectionSetConstraintDof(), to set the number of constrained dofs for a given point, and PetscSectionSetConstraintIndices() which indicates which dofs on the given point are constrained. Once we have this information, a global section can be created using PetscSectionCreateGlobalSection(), and this is done automatically by the DM. A global section returns $-(dof + 1)$ for the number of dofs on an unowned point, and $-(off + 1)$ for its offset on the owning process. This can be used to create global vectors, just as the local section is used to create local vectors.
Data Layout using DMPLEX and PetscFE

A DM can automatically create the local section if given a description of the discretization, for example using a PetscFE object. We demonstrate this by creating a PetscFE that can be configured from the command line. It is a single, scalar field, and is added to the DM using DMSSetField(). When a local or global vector is requested, the DM builds the local and global sections automatically.

```c
DMIsSimplex(dm, &simplex);
PetscFECreateDefault(PETSC_COMM_SELF, dim, 1, simplex, NULL, -1, &fe);
DMSSetField(dm, 0, NULL, (PetscObject) fe);
DMCreateDS(dm);
```

Here the call to DMSSetField() declares the discretization will have one field with the integer label 0 that has one degree of freedom at each point on the DMPlex. To get the $P_3$ section above, we can either give the option -petscspace_degree 3, or call PetscFECreateLagrange() and set the degree directly.

Partitioning and Ordering

In the same way as MatPartitioning or MatGetOrdering(), give the results of a partitioning or ordering of a graph defined by a sparse matrix, PetscPartitionerDMPlexPartition or DMPlexPermute are encoded in an IS. However, the graph is not the adjacency graph of the matrix but the mesh itself. Once the mesh is partitioned and reordered, the data layout from a PetscSection can be used to automatically derive a problem partitioning/ordering.

Influence of Variables on One Another

The Jacobian of a problem represents the influence of some variable on other variables in the problem. Very often, this influence pattern is determined jointly by the computational mesh and discretization. DMCreateMatrix() must compute this pattern when it automatically creates the properly preallocated Jacobian matrix. In DMDA the influence pattern, or what we will call variable adjacency, depends only on the stencil since the topology is Cartesian and the discretization is implicitly finite difference.

In DMPLEX, we allow the user to specify the adjacency topologically, while maintaining good defaults. The pattern is controlled by two flags. The first flag, useCone, indicates whether variables couple first to their boundary and then to neighboring entities, or the reverse. For example, in finite elements, the variables couple to the set of neighboring cells containing the mesh point, and we set the flag to useCone = PETSC_FALSE. By contrast, in finite volumes, cell variables first couple to the cell boundary, and then to the neighbors, so we set the flag to useCone = PETSC_TRUE. The second flag, useClosure, indicates whether we consider the transitive closure of the neighbor relation above, or just a single level. For example, in finite elements, the entire boundary of any cell couples to the interior, and we set the flag to useClosure = PETSC_TRUE. By contrast, in most finite volume methods, cells couple only across faces, and not through vertices, so we set the flag to useClosure = PETSC_FALSE. However, the power of this method is its flexibility. If we wanted a finite volume method that coupled all cells around a vertex, we could easily prescribe that by changing to useClosure = PETSC_TRUE.

---

2 The boundary of a cell is its faces, the boundary of a face is its edges and the boundary of an edge is the two vertices.
3.2.4 Evaluating Residuals

The evaluation of a residual or Jacobian, for most discretizations has the following general form:

- Traverse the mesh, picking out pieces (which in general overlap),
- Extract some values from the current solution vector, associated with this piece,
- Calculate some values for the piece, and
- Insert these values into the residual vector

DMPlex separates these different concerns by passing sets of points from mesh traversal routines to data extraction routines and back. In this way, the PetscSection which structures the data inside a Vec does not need to know anything about the mesh inside a DMPLEX.

The most common mesh traversal is the transitive closure of a point, which is exactly the transitive closure of a point in the DAG using the covering relation. In other words, the transitive closure consists of all points that cover the given point (generally a cell) plus all points that cover those points, etc. So in 2d the transitive closure for a cell consists of edges and vertices while in 3d it consists of faces, edges, and vertices. Note that this closure can be calculated orienting the arrows in either direction. For example, in a finite element calculation, we calculate an integral over each element, and then sum up the contributions to the basis function coefficients. The closure of the element can be expressed discretely as the transitive closure of the element point in the mesh DAG, where each point also has an orientation. Then we can retrieve the data using PetscSection methods,

```c
PetscScalar *a;
PetscInt numPoints, *points = NULL, p;
VecGetArrayRead(u, &a);
DMPlexGetTransitiveClosure(dm, cell, PETSC_TRUE, &numPoints, &points);
for (p = 0; p < numPoints*2; p += 2) {
    PetscInt dof, off, d;
    PetscSectionGetDof(section, points[p], &dof);
    PetscSectionGetOffset(section, points[p], &off);
    for (d = 0; d < dof; ++d) {
        myfunc(a[off+d]);
    }
}
DMPlexRestoreTransitiveClosure(dm, cell, PETSC_TRUE, &numPoints, &points);
VecRestoreArrayRead(u, &a);
```

This operation is so common that we have built a convenience method around it which returns the values in a contiguous array, correctly taking into account the orientations of various mesh points:

```c
const PetscScalar *values;
PetscInt csize;
DMPlexVecGetClosure(dm, section, u, cell, &csize, &values);
// Do integral in quadrature loop putting the result into r[]
DMPlexVecRestoreClosure(dm, section, u, cell, &csize, &values);
DMPlexVecSetClosure(dm, section, residual, cell, &r, ADD_VALUES);
```

A simple example of this kind of calculation is in DMPlexComputeL2Diff_Plex() (source). Note that there is no restriction on the type of cell or dimension of the mesh in the code above, so it will work for polyhedral cells, hybrid meshes, and meshes of any dimension, without change. We can also reverse the covering relation, so that the code works for finite volume methods where we want the data from neighboring cells for each face:
This kind of calculation is used in TS Tutorial ex11.

### 3.2.5 Saving and Loading DMPIlex Data with HDF5

PETSc allows users to save/load DMPIlexs representing meshes, PetscSections representing data layouts on the meshes, and Vecs defined on the data layouts to/from an HDF5 file in parallel, where one can use different number of processes for saving and for loading.

#### Saving

The simplest way to save DM data is to use options for configuration. This requires only the code

```c
DMViewFromOptions(dm, NULL, "-dm_view");
VecViewFromOptions(vec, NULL, "-vec_view")
```

along with the command line options

```
$ ./myprog -dm_view hdf5:myprog.h5 -vec_view hdf5:myprog.h5::append
```

Options prefixes can be used to separately control the saving and loading of various fields. However, the user can have fine grained control by explicitly creating the PETSc objects involved. To save data to "example.h5" file, we can first create a PetscViewer of type PETSCVIEWERHDF5 in FILE_MODE_WRITE mode as:

```c
PetscViewer viewer;
PetscViewerHDF5Open(PETSC_COMM_WORLD, "example.h5", FILE_MODE_WRITE, &viewer);
```

As dm is a DMPIex object representing a mesh, we first give it a mesh name, “plexA”, and save it as:

```c
PetscObjectSetName((PetscObject)dm, "plexA");
PetscViewerPushFormat(viewer, PETSC_VIEWER_HDF5_PETSC);
DMView(dm, viewer);
PetscViewerPopFormat(viewer);
```

The DMView() call is shorthand for the following sequence

```c
DMplexTopologyView(dm, viewer);
DMplexCoordinatesView(dm, viewer);
DMplexLabelsView(dm, viewer);
```
If the *mesh name* is not explicitly set, the default name is used. In the above PETSC_VIEWER_HDF5>PETSC format was used to save the entire representation of the mesh. This format also saves global point numbers attached to the mesh points. In this example the set of all global point numbers is \(X = [0, 11]\).

The data layout, \(s\), needs to be wrapped in a DM object for it to be saved. Here, we create the wrapping DM, \(sdm\), with DMClone(), give it a *dm name*, “dmA”, attach \(s\) to \(sdm\), and save it as:

```c
DMClone(dm, &sdm);
PetscObjectSetName((PetscObject)sdm, "dmA");
DMSetLocalSection(sdm, s);
DMPlexSectionView(dm, viewer, sdm);
```

If the *dm name* is not explicitly set, the default name is to be used. In the above, instead of using DMClone(), one could also create a new DMSHELL object to attach \(s\) to. The first argument of DMPlexSectionView() is a DMPLEX object that represents the mesh, and the third argument is a DM object that carries the data layout that we would like to save. They are, in general, two different objects, and the former carries a *mesh name*, while the latter carries a *dm name*. These names are used to construct a group structure in the HDF5 file. Note that the data layout points are associated with the mesh points, so each of them can also be tagged with a global point number in \(X\): DMPlexSectionView() saves these tags along with the data layout itself, so that, when the mesh and the data layout are loaded separately later, one can associate the points in the former with those in the latter by comparing their global point numbers.

We now create a local vector associated with \(sdm\), e.g., as:

```c
Vec vec;
DMGetLocalVector(sdm, &vec);
```

After setting values of \(vec\), we name it “vecA” and save it as:

```c
PetscObjectSetName((PetscObject)vec, "vecA");
DMPlexLocalVectorView(dm, viewer, sdm, vec);
```

A global vector can be saved in the exact same way with trivial changes.

After saving, we destroy the PetscViewer with:

```c
PetscViewerDestroy(&viewer);
```

The output file “example.h5” now looks like the following:

```bash
$ h5dump --contents example.h5
HDF5 "example.h5" {
  FILE_CONTEXTS {
      group /
      group /topologies
      group /topologies/plexA
      group /topologies/plexA/dms
      group /topologies/plexA/dms/dmA
data/dataset /topologies/plexA/dms/dmA/order
data/dataset /topologies/plexA/dms/dmA/section
      group /topologies/plexA/dms/dmA/section/atlasDof
data/dataset /topologies/plexA/dms/dmA/section/atlasOff
data/dataset /topologies/plexA/dms/dmA/vecs
      group /topologies/plexA/dms/dmA/vecs/vecA
data/dataset /topologies/plexA/dms/dmA/vecs/vecA/vecA
      group /topologies/plexA/labels
      group /topologies/plexA/topology
  }
(continues on next page)
}
```

3.2. DMPlex: Unstructured Grids
Saving in the new parallel HDF5 format

Since PETSc 3.19, we offer a new format which supports parallel loading. To write in this format, you currently need to specify it explicitly using the option

```
-dm_plex_view_hdf5_storage_version 3.0.0
```

The storage version is stored in the file and set automatically when loading (described below). You can check the storage version of the HDF5 file with

```
$ h5dump -a /dmplex_storage_version example.h5
```

To allow for simple and efficient implementation, and good load balancing, the 3.0.0 format changes the way the mesh topology is stored. Different strata (sets of mesh entities with an equal dimension; or vertices, edges, faces, and cells) are now stored separately. The new structure of `/topologies/<mesh_name>/topology` is following:

```
$ h5dump --contents example.h5
HDF5 "example.h5" {
FILE_CONTENTS {
... 
  group /topologies/plexA/topology 
  dataset /topologies/plexA/topology/permutation 
  group /topologies/plexA/topology/strata 
  dataset /topologies/plexA/topology/strata/0/cone_sizes 
  dataset /topologies/plexA/topology/strata/0/cones 
  dataset /topologies/plexA/topology/strata/0/orientations 
  group /topologies/plexA/topology/strata/1 
  dataset /topologies/plexA/topology/strata/1/cone_sizes 
  dataset /topologies/plexA/topology/strata/1/cones 
  dataset /topologies/plexA/topology/strata/1/orientations 
  group /topologies/plexA/topology/strata/2 
  dataset /topologies/plexA/topology/strata/2/cone_sizes 
  dataset /topologies/plexA/topology/strata/2/cones 
  dataset /topologies/plexA/topology/strata/2/orientations 
  group /topologies/plexA/topology/strata/3 
  dataset /topologies/plexA/topology/strata/3/cone_sizes 
  dataset /topologies/plexA/topology/strata/3/cones 
  dataset /topologies/plexA/topology/strata/3/orientations 
}
```

Group `/topologies/<mesh_name>/topology/strata` contains a subgroup for each stratum depth (dimension; 0 for vertices up to 3 for cells). DAG points (mesh entities) have an implicit global numbering, given by the position in `orientations` (or `cone_sizes`) plus the stratum offset. The stratum offset is given by a sum of lengths of all previous strata with respect to the order stored in `/topologies/`.
<mesh_name>/topology/permutation. This global numbering is compatible with the explicit numbering in dataset topology/order of previous versions.

For a DAG point with index \(i\) at depth \(s\), \(\text{cone_sizes}[i]\) gives a size of this point’s cone (set of adjacent entities with depth \(s-1\)). Let \(o = \text{sum(\text{cone_sizes}[0:i])}\) (in Python syntax). Points forming the cone are then given by \(\text{cones}[o:o+\text{cone_sizes}[i]]\) (in numbering relative to the depth \(s-1\)). The orientation of the cone with respect to point \(i\) is then stored in \(\text{orientations}[i]\).

### Loading

To load data from “example.h5” file, we create a PetscViewer of type PETSCVIEWERHDF5 in FILE_MODE_READ mode as:

```c
PetscViewerHDF5Open(PETSC_COMM_WORLD, "example.h5", FILE_MODE_READ, &viewer);
```

We then create a DMPLEX object, give it a mesh name, “plexA”, and load the mesh as:

```c
DMCreate(PETSC_COMM_WORLD, &dm);
DMSerialize(dm, DMPLEX);
PetscObjectSetName((PetscObject)dm, "plexA");
PetscViewerPushFormat(viewer, PETSC_VIEWER_HDF5_PETSC);
DMLoad(dm, viewer);
PetscViewerPopFormat(viewer);
```

where PETSC_VIEWER_HDF5_PETSC format was again used. The user can have more control by replace the single load call with

```c
PetscSF sfO;
DMCreate(PETSC_COMM_WORLD, &dm);
DMSerialize(dm, DMPLEX);
PetscObjectSetName((PetscObject)dm, "plexA");
PetscViewerPushFormat(viewer, PETSC_VIEWER_HDF5_PETSC);
DMPlexTopologyLoad(dm, viewer, &sfO);
DMPlexCoordinatesLoad(dm, viewer, sfO);
PetscViewerPopFormat(viewer);
```

The object returned by DMPlexTopologyLoad(), sfO, is a PetscSF that pushes forward \(X\) to the loaded mesh, dm; this PetscSF is constructed with the global point number tags that we saved along with the mesh points.

As the DMPLEX mesh just loaded might not have a desired distribution, it is common to redistribute the mesh for a better distribution using DMPlexDistribute(), e.g., as:

```c
DM distributedDM;
PetscInt overlap = 1;
PetscSF sfDist, sf;
DMPlexDistribute(dm, overlap, &sfDist, &distributedDM);
if (distributedDM) {
    DMDestroy(&dm);
    dm = distributedDM;
    PetscObjectSetName((PetscObject)dm, "plexA");
}
PetscSFCompose(sfO, sfDist, &sf);
PetscSFDestroy(&sfO);
PetscSFDestroy(&sfDist);
```
Note that the new `DMPLEX` does not automatically inherit the mesh name, so we need to name it “plexA” once again. `sfDist` is a `PetscSF` that pushes forward the loaded mesh to the redistributed mesh, so, composed with `sf0`, it makes the `PetscSF` that pushes forward `X` directly to the redistributed mesh, which we call `sf`.

We then create a new `DM`, `sdm`, with `DMClone()`, give it a `dm name`, “dmA”, and load the on-disk data layout into `sdm` as:

```c
PetscSF globalDataSF, localDataSF;
DMClone(dm, &sdm);
PetscObjectSetName((PetscObject)sdm, "dmA");
DMPlexSectionLoad(dm, viewer, sdm, sf, &globalDataSF, &localDataSF);
```

where we could also create a new `DMSHELL` object instead of using `DMClone()`. Each point in the on-disk data layout being tagged with a global point number in `X`, `DMPlexSectionLoad()` internally constructs a `PetscSF` that pushes forward the on-disk data layout to `X`. Composing this with `sf`, `DMPlexSectionLoad()` internally constructs another `PetscSF` that pushes forward the on-disk data layout directly to the redistributed mesh. It then reconstructs the data layout `s` on the redistributed mesh and attaches it to `sdm`. The objects returned by this function, `globalDataSF` and `localDataSF`, are `PetscSFs` that can be used to migrate the on-disk vector data into local and global `Vecs` defined on `sdm`.

We now create a local vector associated with `sdm`, e.g., as:

```c
Vec vec;
DMGetLocalVector(sdm, &vec);
```

We then name `vec` “vecA” and load the on-disk vector into `vec` as:

```c
PetscObjectSetName((PetscObject)vec, "vecA");
DMPlexLocalVectorLoad(dm, viewer, sdm, localDataSF, localVec);
```

where `localDataSF` knows how to migrate the on-disk vector data into a local `Vec` defined on `sdm`. The on-disk vector can be loaded into a global vector associated with `sdm` in the exact same way with trivial changes.

After loading, we destroy the `PetscViewer` with:

```c
PetscViewerDestroy(&viewer);
```

The above infrastructure works seamlessly in distributed-memory parallel settings, in which one can even use different number of processes for saving and for loading; a more comprehensive example is found in `DMPlex` Tutorial ex12.

### 3.2.6 Metric-based mesh adaptation

`DMPlex` supports mesh adaptation using the Riemannian metric framework. The idea is to use a Riemannian metric space within the mesher. The metric space dictates how mesh resolution should be distributed across the domain. Using this information, the remesher transforms the mesh such that it is a unit mesh when viewed in the metric space. That is, the image of each of its elements under the mapping from Euclidean space into the metric space has edges of unit length.

One of the main advantages of metric-based mesh adaptation is that it allows for fully anisotropic remeshing. That is, it provides a means of controlling the shape and orientation of elements in the adapted mesh, as well as their size. This can be particularly useful for advection-dominated and directionally-dependent problems.
The two main ingredients for metric-based mesh adaptation are an input mesh (i.e. the `DMPLEX`) and a Riemannian metric. The implementation in PETSc assumes that the metric is piecewise linear and continuous across elemental boundaries. Such an object can be created using the routine

```
DMPlexMetricCreate(DM dm, PetscInt field, Vec *metric);
```

A metric must be symmetric positive-definite, so that distances may be properly defined. This can be checked using

```
DMPlexMetricEnforceSPD(DM dm, Vec metricIn, PetscBool restrictSizes, PetscBool restrictAnisotropy, Vec metricOut, Vec determinant);
```

This routine may also be used to enforce minimum and maximum tolerated metric magnitudes (i.e. cell sizes), as well as maximum anisotropy. These quantities can be specified using

```
DMPlexMetricSetMinimumMagnitude(DM dm, PetscReal h_min);
DMPlexMetricSetMaximumMagnitude(DM dm, PetscReal h_max);
DMPlexMetricSetMaximumAnisotropy(DM dm, PetscReal a_max);
```

or the command line arguments

```
-dm_plex_metric_h_min <h_min>
-dm_plex_metric_h_max <h_max>
-dm_plex_metric_a_max <a_max>
```

One simple way to combine two metrics is by simply averaging them entry-by-entry. Another is to intersect them, which amounts to choosing the greatest level of refinement in each direction. These operations are available in PETSc through the routines

```
DMPlexMetricAverage(DM dm, PetscInt numMetrics, PetscReal weights[], Vec metrics[], Vec metricAvg);
DMPlexMetricIntersection(DM dm, PetscInt numMetrics, Vec metrics[], Vec metricInt);
```

However, before combining metrics, it is important that they are scaled in the same way. Scaling also allows the user to control the number of vertices in the adapted mesh (in an approximate sense). This is achieved using the $L^p$ normalization framework, with the routine

```
DMPlexMetricNormalize(DM dm, Vec metricIn, PetscBool restrictSizes, PetscBool restrictAnisotropy, Vec metricOut, Vec determinant);
```

There are two important parameters for the normalization: the normalization order $p$ and the target metric complexity, which is analogous to the vertex count. They are controlled using

```
DMPlexMetricSetNormalizationOrder(DM dm, PetscReal p);
DMPlexMetricSetTargetComplexity(DM dm, PetscReal target);
```

or the command line arguments

```
-dm_plex_metric_p <p>
-dm_plex_metric_target_complexity <target>
```

Two different metric-based mesh adaptation tools are available in PETSc:

- Pragmatic;
- Mmg/ParMmg.
Mmg is a serial package, whereas ParMmg is the MPI version. Note that surface meshing is not currently supported and that ParMmg works only in three dimensions. Mmg can be used for both two and three dimensional problems. Pragmatic, Mmg and ParMmg may be specified by the command line arguments

\begin{verbatim}
-dm_adaptor pragmatic
-dm_adaptor mmg
-dm_adaptor parmmg
\end{verbatim}

Once a metric has been constructed, it can be used to perform metric-based mesh adaptation using the routine

\begin{verbatim}
DMAdaptMetric(DM dm, Vec metric, DMLabel bdLabel, DMLabel rgLabel, DM dmAdapt);
\end{verbatim}

where \texttt{bdLabel} and \texttt{rgLabel} are boundary and interior tags to be preserved under adaptation, respectively.

### 3.3 DMSTAG: Staggered, Structured Grid

For structured (aka “regular”) grids with staggered data (living on elements, faces, edges, and/or vertices), the \texttt{DMSTAG} object is available. This can be useful for problems in many domains, including fluid flow, MHD, and seismology.

It is possible, though cumbersome, to implement a staggered-grid code using multiple \texttt{DMDA} objects, or a single multi-component \texttt{DMDA} object where some degrees of freedom are unused. \texttt{DMSTAG} was developed for two main purposes:

1. To help manage some of the burden of choosing and adhering to the complex indexing conventions needed for staggered grids (in parallel)
2. To provide a uniform abstraction for which scalable solvers and preconditioners may be developed (in particular, using \texttt{PCFIELDSPLIT} and \texttt{PCMG}).

\texttt{DMSTAG} is design to behave much like \texttt{DMDA}, with a couple of important distinctions, and borrows some terminology from \texttt{DMPLEX}.

#### 3.3.1 Terminology

Like a \texttt{DMPLEX} object, a \texttt{DMSTAG} represents a cell complex, distributed in parallel over the ranks of an MPI \texttt{Comm}. It is, however, a very regular complex, consisting of a structured grid of \(d\)-dimensional cells, with \(d \in \{1, 2, 3\}\), which are referred to as elements, \(d - 1\) dimensional cells defining boundaries between these elements, and the boundaries of the domain, and in 2 or more dimensions, boundaries of these cells, all the way down to 0 dimensional cells referred to as vertices. In 2 dimensions, the 1-dimensional element boundaries are referred to as edges or faces. In 3 dimensions, the 2-dimensional element boundaries are referred to as faces and the 1-dimensional boundaries between faces are referred to as edges. The set of cells of a given dimension is referred to as a stratum (which one can think of as a level in DAG representation of the mesh); a \texttt{DMSTAG} object of dimension \(d\) represents a complete cell complex with \(d + 1\) strata (levels).

In the description of any ch\texttt{unstructured} the cells at each level are referred to as points. Thus we adopt that terminology uniformly in PETSc and so furthermore in this document, point will refer to a cell.

Each stratum has a constant number of unknowns (which may be zero) associated with each point (cell) on that level. The distinct unknowns associated with each point are referred to as components.

The structured grid, is like with \texttt{DMDA}, decomposed via a Cartesian product of decompositions in each dimension, giving a rectangular local subdomain on each rank. This is extended by an element-wise stencil width of ghost elements to create an atlas of overlapping patches.
3.3.2 Working with vectors and operators (matrices)

**DMSTAG** allows the user to reason almost entirely about a global indexing of elements. Element indices are simply 1-3 **PetscInt** values, starting at 0, in the back, bottom, left corner of the domain. For instance, element (1, 2, 3), in 3D, is the element second from the left, third from the bottom, and fourth from the back (regardless of how many MPI ranks are used).

To refer to points (elements, faces, edges, and vertices), a value of **DMStagStencilLocation** is used, relative to the element index. The element itself is referred to with **DMSTAG_ELEMENT**, the top right vertex (in 2D) or the top right edge (in 3D) with **DMSTAG_UP_RIGHT**, the back bottom left corner in 3D with **DMSTAG_BACK_DOWN_LEFT**, and so on.

Fig. 3.3 gives a few examples in 2D.

Fig. 3.3: Locations in **DMSTAG** are indexed according to global element indices (here, two in 2D) and a location name. Elements have unique names but other locations can be referred to in more than one way. Element colors correspond to a parallel decomposition, but locations on the grid have names which are invariant to this. Note that the face on the top right can be referred to as being to the left of a “dummy” element (3, 3) outside the physical domain.

Crucially, this global indexing scheme does not include any “ghost” or “padding” unknowns outside the physical domain. This is useful for higher-level operations such as computing norms or developing physics-based solvers. However (unlike **DMDA**), this implies that the global **Vec** do not have a natural block structure, as different strata have different numbers of points (e.g. in 1D there is an “extra” vertex on the right). This regular block structure is, however, very useful for the local representation of the data, so in that case dummy DOF are included, drawn as grey in Fig. 3.4.

For working with **Vec** data, this approach is used to allow direct access to a multi-dimensional, regular-
Fig. 3.4: Local and global representations for a 2D DMSTAG object, 3 by 4 elements, with one degree of freedom on each of the three strata: element (squares), faces (triangles), and vertices (circles). The cell complex is parallelized across 4 MPI ranks. In the global representation, the colors correspond to which rank holds the native representation of the unknown. The 4 local representations are shown, with an (elementwise) stencil “box” stencil width of 1. Unknowns are color by their native rank. Dummy unknowns, which correspond to no global degree of freedom, are colored grey. Note that the local representations have a natural block size of 4, and the global representation has no natural block size.
blocked array. To avoid the user having to know about the internal numbering conventions used, helper functions are used to produce the proper final integer index for a given location and component, referred to as a “slot”. Similarly to DMDAVecGetArrayDOF(), this uses a d + 1 dimensional array in d dimensions. The following snippet give an example of this usage.

```c
/* Set the second component of all vertex dof to 2.0 */
PetscCall(DMStagGetCorners(dm, &s_x, &s_y, &s_z, &n_x, &n_y, &n_z, &n_e_x, &n_e_y, &n_e_z));
PetscCall(DMStagGetLocationSlot(dm, location_vertex, 1, &slot_vertex_2));
PetscCall(DMStagVecGetArray(dm, x, &x_array));
for (PetscInt k = s_z; k < s_z + n_z + n_e_z; ++k) {
  for (PetscInt j = s_y; j < s_y + n_y + n_e_y; ++j) {
    for (PetscInt i = s_x; i < s_x + n_x + n_e_x; ++i) x_array[k][j][i][slot_vertex_2] = 2.0;
  }
}
PetscCall(DMStagVecRestoreArray(dm, x, &x_array));
```

DMSTAG provides a stencil-based method for getting and setting entries of Mat and Vec objects. The follow excerpt from DMSTAG Tutorial ex1 demonstrates the idea. For more, see the manual page for DMStagMatSetValuesStencil().

```c
/* Velocity is either a BC or an interior point */
if (isFirstRank && e == start) {
  DMStagStencil row;
  PetscScalar val;
  row.i = e;
  row.loc = LEFT;
  row.c = 0;
  val = 1.0;
  PetscCall(DMStagMatSetValuesStencil(dmSol, A, 1, &row, 1, &row, &val, INSERT_VALUES));
}
```

The array-based approach for Vec is likely to be more efficient than the stencil-based method just introduced above.

### 3.3.3 Coordinates

DMSTAG, unlike DMDA, supports two approaches to defining coordinates. This is captured by which type of DM is used to represent the coordinates. No default is imposed, so the user must directly or indirectly call DMStagSetCoordinateDMType().

If a second DMSTAG object is used to represent coordinates in “explicit” form, behavior is much like with DMDA - the coordinate DM has d DOF on each stratum corresponding to coordinates associated with each point.

If DMPRODUCT is used instead, coordinates are represented by a DMPRODUCT object referring to a Cartesian product of 1D DMSTAG objects, each of which features explicit coordinates as just mentioned.

Navigating these nested DM in DMPRODUCT can be tedious, but note the existence of helper functions like DMStagSetUniformCoordinatesProduct() and DMStagGetProductCoordinateArrays().
3.3.4 Numberings and internal data layout

While DMSTAG aims to hide the details of its internal data layout, for debugging, optimization, and customization purposes, it can be important to know how DMSTAG internally numbers unknowns.

Internally, each point is canonically associated with an element (top-level point (cell)). For purposes of local, regular-blocked storage, an element is grouped with lower-dimensional points left of, below (“down”), and behind (“back”) it. This means that “canonical” values of DMtagStencilLocation are DMTAG_ELEMENT, plus all entries consisting only of “LEFT”, “DOWN”, and “BACK”. In general, these are the most efficient values to use, unless convenience dictates otherwise, as they are the ones used internally.

When creating the decomposition of the domain to local ranks, and extending these local domains to handle overlapping halo regions and boundary ghost unknowns, this same per-element association is used. This has the advantage of maintaining a regular blocking, but may not be optimal in some situations in terms of data movement.

Numberings are, like DMDA, based on a local “x-fastest, z-slowest” or “PETSc” ordering of elements (see Application Orderings), with ordering of locations canonically associated with each element decided by considering unknowns on each point to be located at the center of their point, and using a nested ordering of the same style. Thus, in 3-D, the ordering of the 8 canonical DMtagStencilLocation values associated with an element is

<table>
<thead>
<tr>
<th>DMTAG_BACK_DOWN_LEFT</th>
<th>DMTAG_BACK_DOWN</th>
</tr>
</thead>
<tbody>
<tr>
<td>DMTAG_BACK_LEFT</td>
<td>DMTAG_BACK</td>
</tr>
<tr>
<td>DMTAG_DOWN_LEFT</td>
<td>DMTAG_DOWN</td>
</tr>
<tr>
<td>DMTAG_LEFT</td>
<td>DMTAG_ELEMENT</td>
</tr>
</tbody>
</table>

Multiple DOF associated with a given point are stored sequentially (as with DMDA).

For local Vecs, this gives a regular-blocked numbering, with the same number of unknowns associated with each element, including some “dummy” unknowns which to not correspond to any (local or global) unknown in the global representation. See Fig. 3.6 for an example.

In the global representation, only physical unknowns are numbered (using the same “Z” ordering for unknowns which are present), giving irregular numbers of unknowns, depending on whether a domain boundary is present. See Fig. 3.5 for an example.

It should be noted that this is an interlaced (AoS) representation. If a segregated (SoA) representation is required, one should use DMCOMPOSITE collecting several DMSTAG objects, perhaps using DMtagCreate-CompatibleDMStag() to quickly create additional DMSTAG objects from an initial one.

3.4 Networks

The DMNETWORK class provides abstractions for representing general unstructured networks such as communication networks, power grid, computer networks, transportation networks, electrical circuits, graphs, and others.
Fig. 3.5: Global numbering scheme for a 2D DMSTAG object with one DOF per stratum. Note that the numbering depends on the parallel decomposition (over 4 ranks, here).
3.4.1 Application flow

The general flow of an application code using DMNETWORK is as follows:

1. Create a network object.

   ```c
   DMNetworkCreate(MPI_Comm comm, DM *dm);
   ```

2. Create components and register them with the network. A “component” is specific application data at a vertex/edge of the network required for its residual evaluation. For example, components could be resistor/inductor data for circuit applications, edge weights for graph problems, or generator/transmission line data for power grids. Components are registered by calling

   ```c
   DMNetworkRegisterComponent(DM dm, const char *name, size_t size, PetscInt *compkey);
   ```

   Here, name is the component name, size is the size of component data, and compkey is an integer key that can be used for setting/getting the component at a vertex or an edge.

3. A DMNETWORK can consist of one or more physical subnetworks. Each subnetwork has its own mathematical model. When multiple subnetworks are used one can (optionally) provide coupling information between subnetworks. That is vertices that are shared between multiple subnetworks; edges can only belong to a single subnetwork. The number of subnetwork is set by calling

   ```c
   DMNetworkSetNumSubNetworks(DM dm, PetscInt nsubnet, PetscInt Nsubnet);
   ```

   Here, nsubnet and Nsubnet are the local and global number of subnetworks.

4. A subnetwork is added to the network by calling

---

Fig. 3.6: Local numbering scheme on rank 1 (Cf. Fig. 3.4) for a 2D DMSTAG object with one DOF per stratum. Note that dummy locations (grey) are used to give a regular block size (here, 4).
DMNetworkAddSubnetwork(dm, const char* name, PetscInt ne, PetscInt edgelist[], &Petscit *netnum);

Here \textit{name} is the subnetwork name, \textit{ne} is the number of local edges on the subnetwork, and \textit{edgelist} is the connectivity for the subnetwork. The output \textit{netnum} is the global numbering of the subnetwork in the network. Each element of \textit{edgelist} is an integer array of size $2 \times ne$ containing the edge connectivity for the subnetwork.

As an example, consider a network comprised of 2 subnetworks that are coupled. The topological information for the network is as follows:

- subnetwork 0: $v_0 \rightarrow v_1 \rightarrow v_2 \rightarrow v_3$
- subnetwork 1: $v_1 \rightarrow v_2 \rightarrow v_0$

The two subnetworks are coupled by merging vertex 0 from subnetwork 0 with vertex 2 from subnetwork 1.

The \textit{edgelist} of this network is

- $\text{edgelist}[0] = \{0,1,1,2,2,3\}$
- $\text{edgelist}[1] = \{1,2,2,0\}$

The coupling is done by calling

DMNetworkAddSharedVertices(dm, anet, bnet, nsv, asv[], bsv[]);

Here \textit{anet} and \textit{bnet} are the first and second subnetwork global numberings returned by \textbf{DMNetworkAddSubnetwork()}, \textit{nsv} is the number of vertices shared by the two subnetworks, \textit{asv} and \textit{bsv} are the vertex indices in the subnetwork \textit{anet} and \textit{bnet}.

5. The next step is to have \textbf{DMNETWORK} create a bare layout (graph) of the network by calling

DMNetworkLayoutSetUp(dm);

6. After completing the previous steps, the network graph is set up, but no physics is associated yet. This is done by adding the components and setting the number of variables to the vertices and edges.

A component and number of variables are added to a vertex/edge by calling

DMNetworkAddComponent(dm, p, compkey, compdata, nvar)

where \textit{p} is the network vertex/edge point in the range obtained by either \textbf{DMNetworkGetVertexRange()}/\textbf{DMNetworkGetEdgeRange()}. \textbf{DMNetworkGetSubnetwork()}, or \textbf{DMNetworkGetSharedVertices()}: \textit{compkey} is the component key returned when registering the component (\textbf{DMNetworkRegisterComponent()}); \textit{compdata} holds the data for the component; and \textit{nvar} is the number of variables associated to the added component at this network point. \textbf{DMNETWORK} currently requires the owner process of the vertex adds all the components and number of variables. \textbf{DMNETWORK} currently assumes the component data to be stored in a contiguous chunk of memory. As such, it does not do any packing/unpacking before/after the component data gets distributed. Any such serialization (packing/unpacking) should be done by the application.

7. Set up network internal data structures.
PETSc/TAO Users Manual, Release 3.20.3

8. Distribute the network (also moves components attached with vertices/edges) to multiple processors.

\[
\text{DMNetworkDistribute(DM } \text{dm, const char partitioner[], PetscInt overlap, DM } \_\text{distDM);} \\
\]

9. Associate the DM with a PETSc solver:

\[
\text{KSPSetDM(KSP ksp, DM } \text{dm) or SNESetDM(SNES snes, DM } \text{dm) or TSSetDM(TS ts, DM } \text{dm).} \\
\]

3.4.2 Utility functions

DMNETWORK provides several utility functions for operations on the network. The most commonly used functions are:

- obtaining iterators for vertices/edges,

\[
\text{DMNetworkGetEdgeRange(DM } \text{dm, PetscInt *eStart, PetscInt *eEnd);} \\
\]

- obtaining iterators for vertices/edges,

\[
\text{DMNetworkGetVertexRange(DM } \text{dm, PetscInt *vStart, PetscInt *vEnd);} \\
\]

- obtaining subnetworks with vertices/edges,

\[
\text{DMNetworkGetSubnetwork(DM } \text{dm, PetscInt netnum, PetscInt *nv, PetscInt *ne, const PetscInt **vtx, const PetscInt **edge);} \\
\]

- checking the status of a vertex,

\[
\text{DMNetworkIsGhostVertex(DM } \text{dm, PetscInt p, PetscBool *isghost);} \\
\]

- checking the status of a vertex,

\[
\text{DMNetworkIsSharedVertex(DM } \text{dm, PetscInt p, PetscBool *isshared);} \\
\]

and retrieving local/global indices of vertex/edge component variables for inserting elements in vectors/matrices,

\[
\text{DMNetworkGetLocalVecOffset(DM } \text{dm, PetscInt p, PetscInt compnum, PetscInt *offset);} \\
\]

\[
\text{DMNetworkGetGlobalVecOffset(DM } \text{dm, PetscInt p, PetscInt compnum, PetscInt *offsetg);} \\
\]

In network applications, one frequently needs to find the supporting edges for a vertex or the connecting vertices covering an edge. These can be obtained by the following two routines.

- finding supporting edges for a vertex,

\[
\text{DMNetworkGetSupportingEdges(DM } \text{dm, PetscInt vertex, PetscInt *nedges, const PetscInt **edges[]);} \\
\]

- connecting vertices covering an edge,

\[
\text{DMNetworkGetConnectedVertices(DM } \text{dm, PetscInt edge, const PetscInt *vertices[]);} \\
\]
3.4.3 Retrieving components and number of variables

The components and the corresponding number of variables set at a vertex/edge can be accessed by

```c
DMNetworkGetComponent(DM dm, PetscInt p, PetscInt compnum, PetscInt *compkey, void **component, PetscInt *nvar)
```

input `compnum` is the component number, output `compkey` is the key set by `DMNetworkRegisterComponent()`. An example of accessing and retrieving the components and number of variables at vertices is:

```c
PetscInt Start, End, numcomps, key, v, compnum;
void *component;
DMNetworkGetVertexRange(dm, &Start, &End);
for (v = Start; v < End; v++) {
    DMNetworkGetNumComponents(dm, v, &numcomps);
    for (compnum = 0; compnum < numcomps; compnum++) {
        DMNetworkGetComponent(dm, v, compnum, &key, &component, &nvar);
        compdata = (UserCompDataType)(component);
    }
}
```

The above example does not explicitly use the component key. It is used when different component types are set at different vertices. In this case, `compkey` is used to differentiate the component type.

3.5 PetscDT: Discretization Technology in PETSc

This chapter discusses the low-level infrastructure which supports higher-level discretizations in PETSc, which includes things such as quadrature and probability distributions.

3.5.1 Quadrature

3.5.2 Probability Distributions

A probability distribution function (PDF) returns the probability density at a given location $P(x)$, so that the probability for an event at location in $[x, x + dx]$ is $P(x) dx$. This means that we must have the normalization condition,

$$\int_{\Omega} P(x) dx = 1.$$  

where :math:`\Omega` is the domain for $x$. This requires that the PDF must have units which are the inverse of the volume form $dx$, meaning that it is homogeneous of order $d$ under scaling

$$pdf(x) = \lambda^d P(\lambda x).$$
We can check this using the normalization condition,

\[
\int_{\Omega} P(x)dx = \int_{\Omega} P(\lambda s)\lambda^d ds \\
= \int_{\Omega} P(s)\lambda^{-d}\lambda^d ds \\
= \int_{\Omega} P(s) ds \\
= 1
\]

The cumulative distribution function (CDF) is the incomplete integral of the PDF,

\[
C(x) = \int_{x_{\text{min}}}^{x} P(s) ds
\]

where \(x_{\text{min}}\) is the lower limit of our domain. We can work out the effect of scaling on the CDF using this definition,

\[
C(\lambda x) = \int_{x_{\text{min}}}^{\lambda x} P(s) ds \\
= \int_{x_{\text{min}}}^{x} \lambda^d P(\lambda t) dt \\
= \int_{x_{\text{min}}}^{x} P(t) dt \\
= C(x)
\]

so the CDF itself is scale invariant and unitless.

We do not add a scale argument to the PDF in PETSc, since all variables are assuming to be dimensionless. This means that inputs to the PDF and CDF should be scaled by the appropriate factor for the units of \(x\), and the output can be rescaled if it is used outside the library.

### 3.6 PetscFE: Finite Element Infrastructure in PETSc

This chapter introduces the **PetscFE** class, and related subclasses **PetscSpace** and **PetscDualSpace**, which are used to represent finite element discretizations. It details there interaction with the **DMPLEX** class to assemble functions and operators over computational meshes, and produce optimal solvers by constructing multilevel iterations, for example using **PCPATCH**. The idea behind these classes is not to encompass all of computational finite elements, but rather to establish an interface and infrastructure that will allow PETSc to leverage the excellent work done in packages such as Firedrake, FEniCS, LibMesh, and Deal.II.

#### 3.6.1 Using Pointwise Functions to Specify Finite Element Problems

See the paper about **Unified Residual Evaluation**, which explains the use of pointwise evaluation functions to describe weak forms.
3.6.2 Describing a particular finite element problem to PETSc

A finite element problem is presented to PETSc in a series of steps. This is both to facilitate automation, and to allow multiple entry points for user code and external packages since so much finite element software already exists. First, we tell the DM, usually a DMPLEX or DMFOREST, that we have a set of finite element fields which we intended to solve for in our problem, using

```
DMAddField(dm, NULL, presDisc);
DMAddField(dm, channelLabel, velDisc);
```

The second argument is a DMLabel object indicating the support of the field on the mesh, with NULL indicating the entire domain. Once we have a set of fields, we calls

```
DMCreateDS(dm);
```

This divides the computational domain into subdomains, called regions in PETSc, each with a unique set of fields supported on it. These subdomain are identified by labels, and each one has a PetscDS object describing the discrete system on that subdomain. There are query functions to get the set of PetscDS objects for the DM, but it is usually easiest to get the proper PetscDS for a given cell using

```
DMGetCellDS(dm, cell, &ds, NULL);
```

Each PetscDS object has a set of fields, each with a PetscFE or PetscFV discretization. This allows it to calculate the size of the local discrete approximation, as well as allocate scratch space for all the associated computations. The final thing needed is to specify the actual equations to be enforced on each region. The PetscDS contains a PetscWeakForm object that holds callback function pointers that define the equations. A simplified, top-level interface through PetscDS allows users to quickly define problems for a single region. For example, in SNES Tutorial ex13, we define the Poisson problem using

```
DMLabel label;
PetscInt f = 0, id = 1;
PetscDSSetResidual(ds, f, f0_trig_inhomogeneous_u, f1_u);
PetscDSSetJacobian(ds, f, f, NULL, NULL, NULL, g3_uu);
PetscDSSetExactSolution(ds, f, trig_inhomogeneous_u, user);
DMGetLabel(dm, "marker", &label);
DMAddBoundary(dm, DM_BC_ESSENTIAL, "wall", label, 1, &id, f, 0, NULL, (void (*)(void)) ex, NULL, user, NULL);
```

where the pointwise functions are

```
static PetscErrorCode trig_inhomogeneous_u(PetscInt dim, PetscReal time, const PetscReal x[], PetscInt Nc, PetscScalar *u, void *ctx)
{
    PetscInt d;
    *u = 0.0;
    for (d = 0; d < dim; ++d) *u += PetscSinReal(2.0*PETSC_PI*x[d]);
    return 0;
}

static void f0_trig_inhomogeneous_u(PetscInt dim, PetscInt Nf, PetscInt NfAux, const PetscInt uOff[], const PetscInt uOff_x[], const PetscScalar u[], const PetscScalar u_t[], const PetscScalar u_x[],
                const PetscInt aOff[], const PetscInt aOff_x[], const PetscScalar a[],
                const PetscScalar a_t[], const PetscScalar a_x[],
                PetscReal t, const PetscReal x[], PetscInt numConstants, const PetscInt aOFF[],
```

(continues on next page)
Notice that we set boundary conditions using `DMAddBoundary`, which will be described later in this chapter. Also we set an exact solution for the field. This can be used to automatically calculate mesh convergence using the `PetscConvEst` object described later in this chapter.

For more complex cases with multiple regions, we need to use the `PetscWeakForm` interface directly. The weak form object allows you to set any number of functions for a given field, and also allows functions to be associated with particular subsets of the mesh using labels and label values. We can reproduce the above problem using the `SetIndex` variants which only set a single function at the specified index, rather than a list of functions. We use a `NULL` label and value, meaning that the entire domain is used.

```c
PetscInt f = 0, val = 0;

PetscDSGetWeakForm(ds, &wf);
PetscWeakFormSetIndexResidual(ds, NULL, val, f, 0, 0, f0_quad_u, 0, f1_u);
PetscWeakFormSetIndexJacobian(ds, NULL, val, f, f, 0, NULL, 0, NULL, 0, NULL, 0, g3_uu);
```

In `SNES Tutorial ex23`, we define the Poisson problem over the entire domain, but in the top half we also define a pressure. The entire problem can be specified as follows

```c
DMGetRegionNumDS(dm, 0, &label, NULL, &ds, NULL);
PetscDSGetWeakForm(ds, &wf);
PetscWeakFormSetIndexResidual(wf, label, 1, 0, 0, 0, f0_quad_u, 0, f1_u);
PetscWeakFormSetIndexJacobian(wf, label, 1, 0, 0, 0, NULL, 0, NULL, 0, NULL, 0, g3_uu);
```
PetscDSSetExactSolution(ds, 0, quad_u, user);
DMGetRegionNumDS(dm, 1, &label, NULL, &ds, NULL);
PetscDSGetWeakForm(dm, &wf);
PetscWeakFormSetIndexResidual(wf, label, 1, 0, 0, 0, f0_quad_u, 0, f1_u);
PetscWeakFormSetIndexJacobian(wf, label, 1, 0, 0, 0, NULL, 0, NULL, 0, NULL, 0, g3__uu);
PetscWeakFormSetIndexResidual(wf, label, 1, 1, 0, 0, f0_quad_p, 0, NULL);
PetscWeakFormSetIndexJacobian(wf, label, 1, 1, 0, 0, g0_pp, 0, NULL, 0, NULL, 0, w_1__NULL);
PetscDSSetExactSolution(ds, 0, quad_u, user);
PetscDSSetExactSolution(ds, 1, quad_p, user);
DMGetLabel(dm, "marker", &label);
DMAddBoundary(dm, DM_BC_ESSENTIAL, "wall", label, 1, &id, 0, 0, NULL, (void (*)(void)) quad_u, NULL, user, NULL);

In the PyLith software we use this capability to combine bulk elasticity with a fault constitutive model integrated over the embedded manifolds corresponding to earthquake faults.

### 3.6.3 Assembling finite element residuals and Jacobians

Once the pointwise functions are set in each PetscDS, mesh traversals can be automatically determined from the DMLabel and value specifications in the keys. This default traversal strategy can be activated by attaching the DM and default callbacks to a solver

```
SNESSetDM(snes, dm);
DMPlexSetSNESLocalFEM(dm, &user, &user, &user);
TSSetDM(ts, dm);
DMTSSetBoundaryLocal(dm, DMPlexTSComputeBoundary, &user);
DMTSSetIFunctionLocal(dm, DMPlexTSComputeIFunctionFEM, &user);
DMTSSetIJacobianLocal(dm, DMPlexTSComputeIJacobianFEM, &user);
```
4.1 PETSc for Fortran Users

Most of the functionality of PETSc can be obtained from Fortran programs. Make sure the suffix of all your Fortran files is .F90, not .f or .f90.

4.1.1 Modules and Include Files

To use PETSc with Fortran you must use both PETSc include files and modules. At the beginning of every function and module definition you need something like

```
#include "petsc/finclude/petscXXX.h"
use petscXXX
```

You can declare PETSc object variables using either of the following:

```
XXX variablename
```

```
tXXX) variablename
```

For example,

```
#include "petsc/finclude/petscvec.h"
use petscvec

Vec b
type(tVec) x
```

PETSc types like PetscInt and PetscReal are simply aliases for basic Fortran types and cannot be written as type(tPetscInt)

The Fortran include files for PETSc are located in the directory $PETSC_DIR/include/petsc/finclude and the module files are located in $PETSC_DIR/$PETSC_ARCH/include

Most Fortran routines have the same names as the corresponding C versions, and PETSc command line options are fully supported. The routine arguments follow the usual Fortran conventions; the user need not worry about passing pointers or values. The calling sequences for the Fortran version are in most cases identical to the C version, except for the error checking variable discussed in Error Checking and a few routines listed in Routines with Different Fortran Interfaces.

When passing floating point numbers into PETSc Fortran subroutines, always make sure you have them marked as double precision (e.g., pass in 10.d0 instead of 10.0 or declare them as PETSc variables, e.g. PetscScalar one = 1.0). Otherwise, the compiler interprets the input as a single precision number,
which can cause crashes or other mysterious problems. We **highly** recommend using the `implicit none` option at the beginning of each Fortran subroutine and declare all variables.

### Error Checking

In the Fortran version, each PETSc routine has as its final argument an integer error variable. The error code is set to be nonzero if an error has been detected; otherwise, it is zero. For example, the Fortran and C variants of `KSPSolve()` are given, respectively, below, where `ierr` denotes the `PetscErrorCode` error variable:

```fortran
    call KSPSolve(ksp, b, x, ierr) ! Fortran
    ierr = KSPSolve(ksp, b, x);   // C
```

For proper error handling one should not use the above syntax instead one should use

```fortran
    PetscCall(KSPSolve(ksp, b, x, ierr)) ! Fortran subroutines
    PetscCallA(KSPSolve(ksp, b, x, ierr)) ! Fortran main program
    PetscCall(KSPSolve(ksp, b, x))        // C
```

### Calling Fortran Routines from C (and C Routines from Fortran)

Different compilers have different methods of naming Fortran routines called from C (or C routines called from Fortran). Most Fortran compilers change all the capital letters in Fortran routines to all lowercase. With some compilers, the Fortran compiler appends an underscore to the end of each Fortran routine name; for example, the Fortran routine `Dabsc()` would be called from C with `dabsc_()`. Other compilers change all the letters in Fortran routine names to capitals.

PETSc provides two macros (defined in C/C++) to help write portable code that mixes C/C++ and Fortran. They are `PETSC_HAVE_FORTRAN_UNDERSCORE` and `PETSC_HAVE_FORTRAN_CAPS`, which will be defined in the file `$PETSC_DIR/$PETSC_ARCH/include/petscconf.h` based on the compilers conventions. The macros are used, for example, as follows:

```c
    #if defined(PETSC_HAVE_FORTRAN_CAPS)
    #define dabsc_ DABSC
    #elif !defined(PETSC_HAVE_FORTRAN_UNDERSCORE)
    #define dabsc_ dabsc
    #endif

    ....
    dabsc_( &n,x,y); /* call the Fortran function */
```

### Passing Null Pointers

Many PETSc C functions have the option of passing a `NULL` argument (for example, the fifth argument of `MatCreateSeqAIJ()`). From Fortran, users must pass `PETSC_NULL_XXX` to indicate a null argument (where XXX is `INTEGER`, `DOUBLE`, `CHARACTER`, `SCALAR`, `VEC`, `MAT`, etc depending on the argument type); passing a literal 0 from Fortran in this case will crash the code. For example, when no options prefix is desired in the routine `PetscOptionsGetInt()`, one must use the following command in Fortran:

```fortran
    PetscCall(PetscOptionsGetInt(PETSC_NULL_OPTIONS, PETSC_NULL_CHARACTER, PETSC_NULL_CHARACTER, '-name', N, flg, ierr))
```
Matrix, Vector and IS Indices

All matrices, vectors and IS in PETSc use zero-based indexing, regardless of whether C or Fortran is being used. For example, MatSetValues() and VecSetValues() always use zero indexing. See Basic Matrix Operations for further details.

Setting Routines

When a function pointer (declared as external in Fortran) is passed as an argument to a PETSc function, such as the test function in KSPSetConvergenceTest(), it is assumed that this function references a routine written in the same language as the PETSc interface function that was called. For instance, if KSPSetConvergenceTest() is called from C, the test function must be a C function. Likewise, if it is called from Fortran, the test function must be (a subroutine) written in Fortran.

Compiling and Linking Fortran Programs

See Writing C/C++ or Fortran Applications.

Routines with Different Fortran Interfaces

The following Fortran routines differ slightly from their C counterparts; see the manual pages and previous discussion in this chapter for details:

```c
PetscInitialize()
PetscOptionsGetString()
```

The following functions are not supported in Fortran:

```c
PetscFClose(), PetscFOpen(), PetscFPrintf(), PetscPrintf(),
PetscPopErrorHandler(), PetscPushErrorHandler(), PetscInfo(),
PetscSetDebugger(), VecGetArrays(), VecRestoreArrays(),
PetscViewerASCIIGetPointer(), PetscViewerBinaryGetDescriptor(),
PetscViewerStringOpen(), PetscViewerStringSPrintf(),
PetscOptionsGetStringArray()
```

Duplicating Multiple Vectors

The Fortran interface to VecDuplicateVecs() differs slightly from the C/C++ variant. To create \( n \) vectors of the same format as an existing vector, the user must declare a vector array, \( \text{v\_new} \) of size \( n \). Then, after VecDuplicateVecs() has been called, \( \text{v\_new} \) will contain (pointers to) the new PETSc vector objects. When finished with the vectors, the user should destroy them by calling VecDestroyVecs(). For example, the following code fragment duplicates \( \text{v\_old} \) to form two new vectors, \( \text{v\_new}(1) \) and \( \text{v\_new}(2) \).

```c
Vec v_old, v_new(2)
PetscInt ierr
PetscScalar alpha
....
PetscCall(VecDuplicateVecs(v_old, 2, v_new, ierr))
alpha = 4.3
PetscCall(VecSet(v_new(1), alpha, ierr))
alpha = 6.0
PetscCall(VecSet(v_new(2), alpha, ierr))
```

(continues on next page)
Routines that Return Fortran Allocatable Arrays

Many PETSc functions that return an array of values in C in an argument (such as ISGetIndices()) return an allocatable array in Fortran. The Fortran function names for these are suffixed with F90 as indicated below. A few routines, such as VecDuplicateVecs() discussed above, do not return a Fortran allocatable array; a large enough array must be explicitly declared before use.

<table>
<thead>
<tr>
<th>C-API</th>
<th>Fortran-API</th>
</tr>
</thead>
<tbody>
<tr>
<td>ISGetIndices()</td>
<td>ISGetIndicesF90()</td>
</tr>
<tr>
<td>ISRestoreIndices()</td>
<td>ISRestoreIndicesF90()</td>
</tr>
<tr>
<td>ISLocalToGlobalMappingGetIndices()</td>
<td>ISLocalToGlobalMappingGetIndicesF90()</td>
</tr>
<tr>
<td>ISLocalToGlobalMappingRestoreIndices()</td>
<td>ISLocalToGlobalMappingRestoreIndicesF90()</td>
</tr>
<tr>
<td>VecGetArray()</td>
<td>VecGetArrayF90()</td>
</tr>
<tr>
<td>VecRestoreArray()</td>
<td>VecRestoreArrayF90()</td>
</tr>
<tr>
<td>VecGetArrayRead()</td>
<td>VecGetArrayReadF90()</td>
</tr>
<tr>
<td>VecRestoreArrayRead()</td>
<td>VecRestoreArrayReadF90()</td>
</tr>
<tr>
<td>VecDuplicateVecs()</td>
<td>VecDuplicateVecsF90()</td>
</tr>
<tr>
<td>VecDestroyVecs()</td>
<td>VecDestroyVecsF90()</td>
</tr>
<tr>
<td>DMDAVecGetArray()</td>
<td>DMDAVecGetArrayF90()</td>
</tr>
<tr>
<td>DMDAVecRestoreArray()</td>
<td>DMDAVecRestoreArrayF90()</td>
</tr>
<tr>
<td>DMDAVecGetArrayRead()</td>
<td>DMDAVecGetArrayReadF90()</td>
</tr>
<tr>
<td>DMDAVecRestoreArrayRead()</td>
<td>DMDAVecRestoreArrayReadF90()</td>
</tr>
<tr>
<td>DMDAVecGetArrayWrite()</td>
<td>DMDAVecGetArrayWriteF90()</td>
</tr>
<tr>
<td>DMDAVecRestoreArrayWrite()</td>
<td>DMDAVecRestoreArrayWriteF90()</td>
</tr>
<tr>
<td>MatGetRowIJ()</td>
<td>MatGetRowIJF90()</td>
</tr>
<tr>
<td>MatRestoreRowIJ()</td>
<td>MatRestoreRowIJF90()</td>
</tr>
<tr>
<td>MatSeqAIJGetArray()</td>
<td>MatSeqAIJGetArrayF90()</td>
</tr>
<tr>
<td>MatSeqAIJRestoreArray()</td>
<td>MatSeqAIJRestoreArrayF90()</td>
</tr>
<tr>
<td>MatMPIAIJGetSeqAIJ()</td>
<td>MatMPIAIJGetSeqAIJF90()</td>
</tr>
<tr>
<td>MatDenseGetArray()</td>
<td>MatDenseGetArrayF90()</td>
</tr>
<tr>
<td>MatDenseRestoreArray()</td>
<td>MatDenseRestoreArrayF90()</td>
</tr>
</tbody>
</table>

The array arguments to these Fortran functions should be declared with forms such as

```
PetscScalar, pointer :: x(:)
PetscInt, pointer :: idx(:)
```

See the manual pages for details and pointers to example programs.
4.1.2 Sample Fortran Programs

Sample programs that illustrate the PETSc interface for Fortran are given below, corresponding to Vec Test ex19f, Vec Tutorial ex4f, Draw Test ex5f, and SNES Tutorial ex1f, respectively. We also refer Fortran programmers to the C examples listed throughout the manual, since PETSc usage within the two languages differs only slightly.

Listing: src/vec/vec/tests/ex19f.F90

```fortran
!/ 
! program main
#include <petsc/finclude/petscvec.h>
use petscvec
implicit none
!/ This example demonstrates basic use of the PETSc Fortran interface
!/ to vectors.
!/ 
PetscInt  n
PetscErrorCode ierr
PetscBool  flg
PetscScalar one,two,three,dot
PetscReal  norm,rdot
Vec        x,y,w
PetscOptions options

n  = 20
one = 1.0
two = 2.0
three = 3.0

PetscCallA(PetscInitialize(ierr))
PetscCallA(PetscOptionsCreate(options,ierr))
PetscCallA(PetscOptionsGetInt(options,PETSC_NULL_CHARACTER,'-n',n,flg,ierr))
PetscCallA(PetscOptionsDestroy(options,ierr))

! Create a vector, then duplicate it
PetscCallA(VecCreate(PETSC_COMM_WORLD,x,ierr))
PetscCallA(VecSetSizes(x,PETSC_DECIDE,n,ierr))
PetscCallA(VecSetFromOptions(x,ierr))
PetscCallA(VecDuplicate(x,y,ierr))
PetscCallA(VecDuplicate(x,w,ierr))

PetscCallA(VecSet(x,one,ierr))
PetscCallA(VecSet(y,two,ierr))

PetscCallA(VecDot(x,y,dot,ierr))

format('Result of inner product ',f10.4)
write(6,100) rdot

PetscCallA(VecScale(x,two,ierr))
PetscCallA(VecNorm(x,NORM_2,norm,ierr))
write(6,110) norm

format('Result of scaling ',f10.4)
```

(continues on next page)
Listing: src/vec/vec/tutorials/ex4f.F90

```
! Description: Illustrates the use of VecSetValues() to set ! multiple values at once; demonstrates VecGetArrayF90().
!
program main

#include <petsc/finclude/petscvec.h>
use petscvec
implicit none
!
! Beginning of program

PetscScalar xwork(6)
PetscScalar, pointer :: xx_v(:,), yy_v(:)
PetscInt i, n, loc(6), isix
PetscErrorCode ierr
Vec x, y

PetscCallA(PetscInitialize(ierr))

n = 6
isix = 6
!
! Create initial vector and duplicate it

PetscCallA(VecCreateSeq(PETSC_COMM_SELF,n,x,ierr))
PetscCallA(VecDuplicate(x,y,ierr))
!
! Fill work arrays with vector entries and locations. Note that ! the vector indices are 0-based in PETSc (for both Fortran and ! C vectors)

do 10 i=1,n

PetscCallA(VecCopy(x,w,ierr))
PetscCallA(VecNorm(w,NORM_2,norm,ierr))
write(6,120) norm
120 format('Result of copy ',f10.4)
PetscCallA(VecAXPY(y,three,x,ierr))
PetscCallA(VecNorm(y,NORM_2,norm,ierr))
write(6,130) norm
130 format('Result of axpy ',f10.4)
PetscCallA(VecDestroy(x,ierr))
PetscCallA(VecDestroy(y,ierr))
PetscCallA(VecDestroy(w,ierr))
PetscCallA(PetscFinalize(ierr))
end
```

(continues on previous page)
loc(i) = i-1
xwork(i) = 10.0*real(i)
10 continue

! Set vector values. Note that we set multiple entries at once.
! Of course, usually one would create a work array that is the
! natural size for a particular problem (not one that is as long
! as the full vector).

PetscCallA(VecSetValues(x,isix,loc,xwork,INSERT_VALUES,ierr))

! Assemble vector
PetscCallA(VecAssemblyBegin(x,ierr))
PetcCallA(VecAssemblyEnd(x,ierr))

! View vector
PetscCallA(PetscObjectSetName(x,'initial vector:',ierr))
PetcCallA(VecView(x,PETSC_VIEWER_STDOUT_SELF,ierr))
PetcCallA(VecCopy(x,y,ierr))

! Get a pointer to vector data.
! - For default PETSc vectors, VecArray() returns a pointer to
!   the data array. Otherwise, the routine is implementation dependent.
! - You MUST call VecRestoreArray() when you no longer need access to
!   the array.
! - Note that the Fortran interface to VecArray() differs from the
!   C version. See the users manual for details.

PetscCallA(VecGetArrayF90(x,xx_v,ierr))
PetcCallA(VecGetArrayF90(y,yy_v,ierr))

! Modify vector data
   do 30 i=1,n
      xx_v(i) = 100.0*real(i)
      yy_v(i) = 1000.0*real(i)
   30 continue

! Restore vectors
PetscCallA(VecRestoreArrayF90(x,xx_v,ierr))
PetcCallA(VecRestoreArrayF90(y,yy_v,ierr))

! View vectors
PetscCallA(PetscObjectSetName(x,'new vector 1:',ierr))
PetcCallA(VecView(x,PETSC_VIEWER_STDOUT_SELF,ierr))

PetscCallA(PetscObjectSetName(y,'new vector 2:',ierr))
PetcCallA(VecView(y,PETSC_VIEWER_STDOUT_SELF,ierr))

! Free work space. All PETSc objects should be destroyed when they
! are no longer needed.

PetscCallA(VecDestroy(x,ierr))
PetcCallA(VecDestroy(y,ierr))
PetscCallA(PetscFinalize(ierr))

end

Listing: src/sys/classes/draw/tests/ex5f.F90

! This example demonstrates basic use of the Fortran interface for PetscDraw routines.
!
PetscDraw          draw
PetscDrawLG         lg
PetscDrawAxis       axis
PetscErrorCode      ierr
PetscBool           flg
integer             x,y,width,height
PetscScalar         xd,yd
PetscReal           ten
PetscInt            i,n,w,h
PetscInt            one

n    = 15
x    = 0
y    = 0
w    = 400
h    = 300
ten  = 10.0
one  = 1

PetscCallA(PetscInitialize(ierr))

/* GetInt requires a PetscInt so have to do this ugly setting */
PetscCallA(PetscOptionsGetInt(PETSC_NULL_OPTIONS,PETSC_NULL_CHARACTER,'-width',width,flg,ierr))
PetscCallA(PetscOptionsGetInt(PETSC_NULL_OPTIONS,PETSC_NULL_CHARACTER,'-height',height,flg,ierr))
PetscCallA(PetscDrawCreate(PETSC_COMM_WORLD,PETSC_NULL_CHARACTER,x,y,width,height,draw,ierr))
PetscCallA(PetscDrawSetFromOptions(draw,ierr))
PetscCallA(PetscDrawLGCreate(draw,one,lg,ierr))
PetscCallA(PetscDrawLGGetAxis(lg,axis,ierr))
PetscCallA(PetscDrawAxisSetColors(axis,PETSC_DRAW_BLACK,PETSC_DRAW_RED,PETSC_DRAW_GREEN,PETSC_DRAW_BLUE))

(continues on next page)
PetscCallA(PetscDrawAxisSetLabels(axis,'toplabel','xlabel','ylabel',ierr))

\begin{verbatim}
do 10, i=0,n-1
   xd = real(i) - 5.0
   yd = xd*xd
   PetscCallA(PetscDrawLGAddPoint(lg,xd,yd,ierr))
10 continue
\end{verbatim}

PetscCallA(PetscDrawLGSetUseMarkers(lg,PETSC_TRUE,ierr))
PetscCallA(PetscDrawLGDraw(lg,ierr))

PetscCallA(PetscSleep(ten,ierr))

PetscCallA(PetscDrawLGDestroy(lg,ierr))
PetscCallA(PetscDrawDestroy(draw,ierr))
PetscCallA(PetscFinalize(ierr))
end

Listing: src/snes/tutorials/ex1f.F90

! ! Description: Uses the Newton method to solve a two-variable system.
!
program main
#include <petsc/finclude/petsc.h>
use petsc
  implicit none
/
! - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
! Variable declarations
/
! - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
/
! Variables:
!    snes - nonlinear solver
!    ksp - linear solver
!    pc - preconditioner context
!    ksp - Krylov subspace method context
!    x, r - solution, residual vectors
!    J - Jacobian matrix
!    its - iterations for convergence
/
SNES  snes
PC    pc
KSP   ksp
Vec   x,r
Mat   J
SNESLineSearch linesearch
PetscErrorCode ierr
PetscInt its,i2,i20
PetscMPIInt size,rank
PetscScalar pfive
PetscReal tol
PetscBool setls
#if defined(PETSC_USE_LOG)
PetscViewer viewer
#endif
double precision threshold, oldthreshold

! Note: Any user-defined Fortran routines (such as FormJacobian)
! MUST be declared as external.

external FormFunction, FormJacobian, MyLineSearch

! - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
! Beginning of program
! - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -

PetscCallA(PetscInitialize(ierr))
PetscCallA(PetscLogNestedBegin(ierr))
threshold = 1.0
PetscCallA(PetscLogSetThreshold(threshold, oldthreshold, ierr))
PetscCallMPIA(MPI_Comm_size(PETSC_COMM_WORLD, size, ierr))
PetscCallMPIA(MPI_Comm_rank(PETSC_COMM_WORLD, rank, ierr))
PetscCheckA(size .eq. 1, PETSC_COMM_SELF, PETSC_ERR_WRONG_MPI_SIZE, 'Uniprocessor example')

  i2 = 2
  i20 = 20

! Create nonlinear solver context
! - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -

PetscCallA(SNESCreate(PETSC_COMM_WORLD, snes, ierr))

! Create matrix and vector data structures; set corresponding routines
! - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -

! Create vectors for solution and nonlinear function

    PetscCallA(VecCreateSeq(PETSC_COMM_SELF, i2, x, ierr))
    PetscCallA(VecDuplicate(x, r, ierr))

! Create Jacobian matrix data structure

    PetscCallA(MatCreate(PETSC_COMM_SELF, J, ierr))
    PetscCallA(MatSetSizes(J, PETSC_DECIDE, PETSC_DECIDE, i2, i2, ierr))
    PetscCallA(MatSetFromOptions(J, ierr))
    PetscCallA(MatSetUp(J, ierr))

! Set function evaluation routine and vector

    PetscCallA(SNESSetFunction(snes, r, FormFunction, 0, ierr))

! Set Jacobian matrix data structure and Jacobian evaluation routine

    PetscCallA(SNESSetJacobian(snes, J, J, FormJacobian, 0, ierr))
Customize nonlinear solver; set runtime options

Set linear solver defaults for this problem. By extracting the
KSP, KSP, and PC contexts from the SNES context, we can then
directly call any KSP, KSP, and PC routines to set various options.

```fortran
PetscCallA(SNESGetKSP(snes, ksp, ierr))
PetscCallA(KSPGetPC(ksp, pc, ierr))
PetscCallA(PCSetType(pc, PCNONE, ierr))
tol = 1.e-4
PetscCallA(KSPSetTolerances(ksp, tol, PETSC_DEFAULT_REAL, PETSC_DEFAULT_REAL, i20, ierr))
```

Set SNES/KSP/KSP/PC runtime options, e.g.,
- `snes_view` -snes_monitor -ksp_type <ksp> -pc_type <pc>
These options will override those specified above as long as
SNESetFromOptions() is called _after_ any other customization
routines.

```fortran
PetscCallA(SNNESSetFromOptions(snes, ierr))
PetscCallA(PetscOptionsHasName(PETSC_NULL_OPTIONS, PETSC_NULL_CHARACTER, '-setls', setls, ierr))
```

if (setls)
  PetscCallA(SNESGetLineSearch(snes, linesearch, ierr))
PetscCallA(SNESLineSearchSetType(linesearch, 'shell', ierr))
PetscCallA(SNESLineSearchShellSetUserFunc(linesearch, MyLineSearch, 0, ierr))
endif

Evaluate initial guess; then solve nonlinear system

Note: The user should initialize the vector, x, with the initial guess
for the nonlinear solver prior to calling SNESolve(). In particular,
to employ an initial guess of zero, the user should explicitly set
this vector to zero by calling VecSet().

```fortran
pfive = 0.5
PetscCallA(VecSet(x, pfive, ierr))
PetscCallA(SNESSolve(snes, PETSC_NULL_VEC, x, ierr))
```

View solver converged reason; we could instead use the option `-snes_converged_reason`
```fortran
PetscCallA(SNESConvergedReasonView(snes, PETSC_VIEWER_STDOUT_WORLD, ierr))
PetscCallA(SNESGetIterationNumber(snes, its, ierr))
if (rank .eq. 0) then
  write(6,100) its
endif
100 format('Number of SNES iterations = ',i5)
```
Free work space. All PETSc objects should be destroyed when they
are no longer needed.

PetscCall(VecDestroy(x,ierr))
PetscCall(VecDestroy(r,ierr))
PetscCall(MatDestroy(J,ierr))
PetscCall(SNESDestroy(snes,ierr))

#if defined(PETSC_USE_LOG)
   PetscCall(PetscViewerASCIIOpen(PETSC_COMM_WORLD,'filename.xml',viewer,ierr))
   PetscCall(PetscViewerPushFormat(viewer,PETSC_VIEWER_ASCII_XML,ierr))
   PetscCall(PetscLogView(viewer,ierr))
   PetscCall(PetscViewerDestroy(viewer,ierr))
#endif

PetscCall(PetscFinalize(ierr))
end

FormFunction - Evaluates nonlinear function, F(x).

Input Parameters:
- snes - the SNES context
- x - input vector
- dummy - optional user-defined context (not used here)

Output Parameter:
- f - function vector

subroutine FormFunction(snes,x,f,dummy,ierr)
   use petscsnes
   implicit none
   SNES   snes
   Vec    x,f
   PetscErrorCode ierr
   integer   dummy(*)

   ! Declarations for use with local arrays
   PetscScalar,pointer :: lx_v(:),lf_v(:)

   ! Get pointers to vector data.
   ! - VecGetArrayF90() returns a pointer to the data array.
   ! - You MUST call VecRestoreArrayF90() when you no longer need access to
     the array.
   PetscCall(VecGetArrayReadF90(x,lx_v,ierr))
   PetscCall(VecGetArrayF90(f,lf_v,ierr))

   ! Compute function
   lf_v(1) = lx_v(1)*lx_v(1) + lx_v(1)*lx_v(2) - 3.0
   lf_v(2) = lx_v(1)*lx_v(2) + lx_v(2)*lx_v(2) - 6.0

   ! Restore vectors
```
PetscCall(VecRestoreArrayReadF90(x,lx_v,ierr))
PetscCall(VecRestoreArrayF90(f,lf_v,ierr))

return
end

! ---------------------------------------------------------------------
!
! FormJacobian - Evaluates Jacobian matrix.
!
! Input Parameters:
! snes - the SNES context
! x - input vector
! dummy - optional user-defined context (not used here)
!
! Output Parameters:
! A - Jacobian matrix
! B - optionally different preconditioning matrix
!
subroutine FormJacobian(snes,X,jac,B,dummy,ierr)
use petscsnes
implicit none
SNES snes
Vec X
Mat jac,B
PetscScalar A(4)
PetscErrorCode ierr
PetscInt idx(2),i2
integer dummy(*)

! Declarations for use with local arrays
PetscScalar,pointer :: lx_v(:)

! Get pointer to vector data
i2 = 2
PetscCall(VecGetArrayReadF90(x,lx_v,ierr))

! Compute Jacobian entries and insert into matrix.
! - Since this is such a small problem, we set all entries for
! the matrix at once.
! - Note that MatSetValues() uses 0-based row and column numbers
! in Fortran as well as in C (as set here in the array idx).

idx(1) = 0
idx(2) = 1
A(1) = 2.0*lx_v(1) + lx_v(2)
A(2) = lx_v(1)
A(3) = lx_v(2)
A(4) = lx_v(1) + 2.0*lx_v(2)
PetscCall(MatSetValues(B,i2,idx,i2,idx,A,INSERT_VALUES,ierr))

! Restore vector
```
PetscCall(VecRestoreArrayReadF90(x, lx_v, ierr))

! Assemble matrix

PetscCall(MatAssemblyBegin(B, MAT_FINAL_ASSEMBLY, ierr))
PetscCall(MatAssemblyEnd(B, MAT_FINAL_ASSEMBLY, ierr))
if (B .ne. jac) then
  PetscCall(MatAssemblyBegin(jac, MAT_FINAL_ASSEMBLY, ierr))
  PetscCall(MatAssemblyEnd(jac, MAT_FINAL_ASSEMBLY, ierr))
endif
return
end

subroutine MyLineSearch(linesearch, lctx, ierr)
use petscsnes
implicit none
SNESLineSearch   linesearch
SNES            snes
integer         lctx
Vec             x, f, g, y, w
PetscReal       ynorm, gnorm, xnorm
PetscErrorCode  ierr
PetscScalar     mone

mone = -1.0
PetscCall(SNESLineSearchGetSNES(linesearch, snes, ierr))
PetscCall(SNESLineSearchGetVecs(linesearch, x, f, y, w, g, ierr))
PetscCall(VecNorm(y, NORM_2, ynorm, ierr))
PetscCall(VecAXPY(x, mone, y, ierr))
PetscCall(SNESComputeFunction(snes, x, f, ierr))
PetscCall(VecNorm(f, NORM_2, gnorm, ierr))
PetscCall(VecNorm(x, NORM_2, xnorm, ierr))
PetscCall(VecNorm(y, NORM_2, ynorm, ierr))
PetscCall(SNESLineSearchSetNorms(linesearch, xnorm, gnorm, ynorm, ierr))
return
end

4.2 Using MATLAB with PETSc

There are three basic ways to use MATLAB with PETSc:

1. Dumping Data for MATLAB into files to be read into MATLAB,
2. Sending Data to an Interactive MATLAB Session from a running PETSc program to a MATLAB process where you may interactively type MATLAB commands (or run scripts), and
3. Using the MATLAB Compute Engine to send data back and forth between PETSc and MATLAB where MATLAB commands are issued, not interactively, but from a script or the PETSc program (this uses the MATLAB Engine).
For the latter two approaches one must ./configure PETSc with the argument `--with-matlab` `--with-matlab-dir=matlab_root_directory`.

## 4.2.1 Dumping Data for MATLAB

### Dumping ASCII MATLAB data

One can dump PETSc matrices and vectors to the screen in an ASCII format that MATLAB can read in directly. This is done with the command line options `-vec_view ::ascii_matlab` or `-mat_view ::ascii_matlab`. To write a file, use `-vec_view :filename.m:ascii_matlab` or `-mat_view :filename.m:ascii_matlab`.

This causes the PETSc program to print the vectors and matrices every time `VecAssemblyEnd()` or `MatAssemblyEnd()` are called. To provide finer control over when and what vectors and matrices are dumped one can use the `VecView()` and `MatView()` functions with a viewer type of `PETSCVIEWERASCII` (see `PetscViewerASCIIOpen()`, `PETSC_VIEWER_STDOUT_WORLD`, `PETSC_VIEWER_STDOUT_SELF`, or `PETSC_VIEWER_STDOUT_(MPI_Comm)`). Before calling the viewer set the output type with, for example,

```c
PetscViewerPushFormat(PETSC_VIEWER_STDOUT_WORLD,PETSC_VIEWER_ASCII_MATLAB);
VecView(A,PETSC_VIEWER_STDOUT_WORLD);
PetscViewerPopFormat(PETSC_VIEWER_STDOUT_WORLD);
```

The name of each PETSc variable printed for MATLAB may be set with

```c
PetscObjectSetName((PetscObject)A,"name");
```

If no name is specified, the object is given a default name using `PetscObjectName()`.

### Dumping Binary Data for MATLAB

One can also read PETSc binary files (see Viewers: Looking at PETSc Objects) directly into MATLAB via the scripts available in `$PETSC_DIR/share/petsc/matlab`. This requires less disk space and is recommended for all but the smallest data sizes. One can also use

```c
PetscViewerPushFormat(viewer,PETSC_VIEWER_BINARY_MATLAB)
```

to dump both a PETSc binary file and a corresponding `.info` file which `PetscReadBinaryMatlab.m` will use to format the binary file in more complex cases, such as using a DMDA. For an example, see DM Tutorial ex7. In MATLAB one may then generate a useful structure. For example:

```matlab
setenv('PETSC_DIR','~/petsc');
setenv('PETSC_ARCH','arch-darwin-double-debug');
addpath('~/petsc/share/petsc/matlab');
gridData=PetscReadBinaryMatlab('output_file');
```
4.2.2 Sending Data to an Interactive MATLAB Session

One creates a viewer to MATLAB via

```c
PetscViewerSocketOpen(MPI_Comm, char *machine, int port, PetscViewer *v);
```

(port is usually set to PETSC_DEFAULT; use NULL for the machine if the MATLAB interactive session is running on the same machine as the PETSc program) and then sends matrices or vectors via

```c
VecView(Vec A, v);
MatView(Mat B, v);
```

See Viewers: Looking at PETSc Objects for more on PETSc viewers. One may start the MATLAB program manually or use the PETSc command `PetscStartMatlab(MPI_Comm, char *machine, char *script, FILE **fp)`; where `machine` and `script` may be NULL. It is also possible to start your PETSc program from MATLAB via `launch()`.

To receive the objects in MATLAB, make sure that `$PETSC_DIR/$PETSC_ARCH/lib/petsc/matlab` and `$PETSC_DIR/share/petsc/matlab` are in the MATLAB path. Use `p = PetscOpenSocket();` (or `p = PetscOpenSocket(portnum)` if you provided a port number in your call to `PetscViewerSocketOpen()`), and then `a = PetscBinaryRead(p);` returns the object passed from PETSc. `PetscBinaryRead()` may be called any number of times. Each call should correspond on the PETSc side with viewing a single vector or matrix. `close()` closes the connection from MATLAB. On the PETSc side, one should destroy the viewer object with `PetscViewerDestroy()`.

For an example, which includes sending data back to PETSc, see Vec Tutorial `ex42` and the associated `.m` file.

4.2.3 Using the MATLAB Compute Engine

One creates access to the MATLAB engine via

```c
PetscMatlabEngineCreate(MPI_Comm comm, char *machine, PetscMatlabEngine *e);
```

where `machine` is the name of the machine hosting MATLAB (NULL may be used for localhost). One can send objects to MATLAB via

```c
PetscMatlabEnginePut(PetscMatlabEngine e, PetscObject obj);
```

One can get objects via

```c
PetscMatlabEngineGet(PetscMatlabEngine e, PetscObject obj);
```

Similarly, one can send arrays via

```c
PetscMatlabEnginePutArray(PetscMatlabEngine e, int m, int n, PetscScalar *array, char _*name);
```

and get them back via

```c
PetscMatlabEngineGetArray(PetscMatlabEngine e, int m, int n, PetscScalar *array, char _*name);
```

One cannot use MATLAB interactively in this mode but one can send MATLAB commands via
PetscMatlabEngineEvaluate(PetscMatlabEngine,"format",...);

where *format* has the usual *printf()* format. For example,

PetscMatlabEngineEvaluate(PetscMatlabEngine,"x = \%g *y + z;",avalue);

The name of each PETSc variable passed to MATLAB may be set with

PetscObjectSetName((PetscObject)A,"name");

Text responses can be returned from MATLAB via

PetscMatlabEngineGetOutput(PetscMatlabEngine,char **);

or

PetscMatlabEnginePrintOutput(PetscMatlabEngine,FILE*).

There is a short-cut to starting the MATLAB engine with PETSC_MATLAB_ENGINE_(MPI_Comm).

If you are running PETSc on a cluster (or machine) that does not have a license for MATLAB, you might be able to run MATLAB on the head node of the cluster or some other machine accessible to the cluster using the -matlab_engine_host hostname option.

### 4.2.4 Licensing the MATLAB Compute Engine on a cluster

To activate MATLAB on head node which does not have access to the internet.\(^1\)

First ssh into the head node using the command: ssh node_name


Login to your MathWorks Account from a computer which has internet access. You will see the available license that your account has. Select a license from the list.

Then, select Install and Activate option and select the Activate to Retrieve License File option.

Enter the information and click Continue.

An option to download the License file will appear. Download it and copy the license file to the cluster (your home directory). Now, launch MATLAB where you have sshed into your head node.

Select the Activate manually without the internet option and click Next >. Browse and locate the license file.

MATLAB is activated and ready to use.

---


Manually Activate Software on a Computer

Select software release.

**Release**  R2020b

Describe computer on which software will be activated.

**Operating System**  Linux

**Host ID**

How do I find my computer's host id?

**Computer Login Name**

How do I find my Computer Login Name?

Provide a label so you can easily identify this computer later.

**Activation Label**

[Buttons: Cancel, Continue]
Activate MathWorks Software

Activation is a process that verifies licensed use of MathWorks products. This process validates the license and ensures that it is not used on more systems than allowed by the license option you have acquired.

- Activate automatically using the internet (recommended)
- Activate manually without the internet

Activate without an Internet connection

- Enter the full path to your license file, including the file name:
- I do not have a license file. Help me with the next steps.
4.3 Profiling

PETSc includes a consistent, lightweight scheme for profiling application programs. The PETSc routines automatically log performance data if certain options are specified at runtime. The user can also log information about application codes for a complete picture of performance. In addition, as described in Interpreting -log_view Output: The Basics, PETSc provides a mechanism for printing informative messages about computations. Basic Profiling Information introduces the various profiling options in PETSc, while the remainder of the chapter focuses on details such as monitoring application codes and tips for accurate profiling.

4.3.1 Basic Profiling Information

If an application code and the PETSc libraries have been configured with -with-log=1, the default, then various kinds of profiling of code between calls to PetscInitialize() and PetscFinalize() can be activated at runtime. The profiling options include the following:

- **-log_view [:filename]** - Prints an ASCII version of performance data at program’s conclusion. These statistics are comprehensive and concise and require little overhead; thus, -log_view is intended as the primary means of monitoring the performance of PETSc codes. See Interpreting -log_view Output: The Basics

- **-info [infofile]** - Prints verbose information about code to stdout or an optional file. This option provides details about algorithms, data structures, etc. Since the overhead of printing such output slows a code, this option should not be used when evaluating a program’s performance. See Interpreting -info Output: Informative Messages

- **-log_trace [logfile]** - Traces the beginning and ending of all PETSc events. This option, which can be used in conjunction with -info, is useful to see where a program is hanging without running in the debugger. See PetscLogTraceBegin().

As discussed in Using -log_mpe with Jumpshot, additional profiling can be done with MPE.

Interpreting -log_view Output: The Basics

As shown in listing in Profiling Programs, the option -log_view [:filename] activates printing of profile data to standard output or an ASCII file at the conclusion of a program. See PetscLogView() for all the possible output options.

We print performance data for each routine, organized by PETSc libraries, followed by any user-defined events (discussed in Profiling Application Codes). For each routine, the output data include the maximum time and floating point operation (flop) rate over all processes. Information about parallel performance is also included, as discussed in the following section.

For the purpose of PETSc floating point operation counting, we define one flop as one operation of any of the following types: multiplication, division, addition, or subtraction. For example, one VecAXPY() operation, which computes \( y = \alpha x + y \) for vectors of length \( N \), requires \( 2N \) flop (consisting of \( N \) additions and \( N \) multiplications). Bear in mind that flop rates present only a limited view of performance, since memory loads and stores are the real performance barrier.

For simplicity, the remainder of this discussion focuses on interpreting profile data for the KSP library, which provides the linear solvers at the heart of the PETSc package. Recall the hierarchical organization of the PETSc library, as shown in Numerical Libraries in PETSc. Each KSP solver is composed of a PC (preconditioner) and a KSP (Krylov subspace) part, which are in turn built on top of the Mat (matrix) and Vec (vector) modules. Thus, operations in the KSP module are composed of lower-level operations in these
packages. Note also that the nonlinear solvers library, \texttt{SNES}, is built on top of the \texttt{KSP} module, and the timestepping library, \texttt{TS}, is in turn built on top of \texttt{SNES}.

We briefly discuss interpretation of the sample output in listing, which was generated by solving a linear system on one process using restarted GMRES and ILU preconditioning. The linear solvers in \texttt{KSP} consist of two basic phases, \texttt{KSPSetUp()} and \texttt{KSPSolve()}, each of which consists of a variety of actions, depending on the particular solution technique. For the case of using the \texttt{PCILU} preconditioner and \texttt{KSPGMRES} Krylov subspace method, the breakdown of PETSc routines is listed below. As indicated by the levels of indentation, the operations in \texttt{KSPSetUp()} include all of the operations within \texttt{PCSetUp()}, which in turn include \texttt{MatILUFactor()}, and so on.

- \texttt{KSPSetUp} - Set up linear solver
  - \texttt{PCSetUp} - Set up preconditioner
    - \texttt{MatILUFactor} - Factor preconditioning matrix
      - \texttt{MatILUFactorSymbolic} - Symbolic factorization phase
      - \texttt{MatILUFactorNumeric} - Numeric factorization phase

- \texttt{KSPSolve} - Solve linear system
  - \texttt{PCApply} - Apply preconditioner
    - \texttt{MatSolve} - Forward/backward triangular solves
  - \texttt{KSPGMRESOrthog} - Orthogonalization in GMRES
    - \texttt{VecDot} or \texttt{VecMDot} - Inner products
    - \texttt{VecAXPY} or \texttt{VecMAXPY} - vector updates
  - \texttt{MatMult} - Matrix-vector product
  - \texttt{MatMultAdd} - Matrix-vector product + vector addition
    - \texttt{VecScale}, \texttt{VecNorm}, \texttt{VecAXPY}, \texttt{VecCopy}, ...

The summaries printed via \texttt{-log_view} reflect this routine hierarchy. For example, the performance summaries for a particular high-level routine such as \texttt{KSPSolve()} include all of the operations accumulated in the lower-level components that make up the routine.

The output produced with \texttt{-log_view} is flat, meaning that the hierarchy of PETSc operations is not completely clear. For a particular problem, the user should generally have an idea of the basic operations that are required for its implementation (e.g., which operations are performed when using GMRES and ILU, as described above), so that interpreting the \texttt{-log_view} data should be relatively straightforward. If this is problematic then it is also possible to examine the profiling information in a nested format. For more information see \textit{Profiling Nested Events}.

**Interpreting \texttt{-log_view} Output: Parallel Performance**

We next discuss performance summaries for parallel programs, as shown within listing and listing, which presents the output generated by the \texttt{-log_view} option. The program that generated this data is KSP Tutorial \texttt{ex10}. The code loads a matrix and right-hand-side vector from a binary file and then solves the resulting linear system; the program then repeats this process for a second linear system. This particular case was run on four processors of an Intel x86_64 Linux cluster, using restarted GMRES and the block Jacobi preconditioner, where each block was solved with ILU. The two input files \texttt{medium} and \texttt{arco6} can be obtained from datafiles, see \texttt{petsc_repositories}.

The first listing presents an overall performance summary, including times, floating-point operations, computational rates, and message-passing activity (such as the number and size of messages sent and collective
operations). Summaries for various user-defined stages of monitoring (as discussed in Profiling Multiple Sections of Code) are also given. Information about the various phases of computation then follow (as shown separately here in the second listing). Finally, a summary of object creation and destruction is presented.

```
mpiexec -n 4 ./ex10 -f0 medium -f1 arco6 -ksp_gmres_classicalgramschmidt -log_view -
    -mat_type baij \
    -matload_block_size 3 -pc_type bjacobi -options_left
```

Number of iterations = 19
Residual norm 1.088292e-05
Number of iterations = 59
Residual norm 3.871022e-02

PETSc Performance Summary: 

```
./ex10 on a intel-bdw-opt named beboplogin4 with 4 processors, by jczhang Mon Apr 23
13:36:54 2018
Using Petsc Development GIT revision: v3.9-163-gbe3efd42 GIT Date: 2018-04-16
10:45:40 -0500
```

<table>
<thead>
<tr>
<th>Time (sec):</th>
<th>Max</th>
<th>Max/Min</th>
<th>Avg</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>Objects:</td>
<td>1.060e+02</td>
<td>1.00000</td>
<td>1.060e+02</td>
<td></td>
</tr>
<tr>
<td>Flops:</td>
<td>2.361e+08</td>
<td>1.00684</td>
<td>2.353e+08</td>
<td>9.413e+08</td>
</tr>
<tr>
<td>Flops/sec:</td>
<td>1.277e+09</td>
<td>1.00685</td>
<td>1.273e+09</td>
<td>5.091e+09</td>
</tr>
<tr>
<td>MPI Msg Count:</td>
<td>2.360e+02</td>
<td>1.34857</td>
<td>2.061e+02</td>
<td>8.245e+02</td>
</tr>
<tr>
<td>MPI Msg Len (bytes):</td>
<td>1.256e+07</td>
<td>2.24620</td>
<td>4.071e+04</td>
<td>3.357e+07</td>
</tr>
<tr>
<td>MPI Reductions:</td>
<td>2.160e+02</td>
<td>1.00000</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Summary of Stages: 

```
<table>
<thead>
<tr>
<th>Message Lengths --</th>
<th>-- Reductions --</th>
</tr>
</thead>
<tbody>
<tr>
<td>%Total counts</td>
<td>%Total Avg</td>
</tr>
<tr>
<td>%Total Avg</td>
<td>%Total counts</td>
</tr>
</tbody>
</table>
```

Object Type Creations Destinations (Reports information only for process 0.)

(continues on next page)

4.3. Profiling 249
We next focus on the summaries for the various phases of the computation, as given in the table within the following listing. The summary for each phase presents the maximum times and flop rates over all processes, as well as the ratio of maximum to minimum times and flop rates for all processes. A ratio of approximately 1 indicates that computations within a given phase are well balanced among the processes; as the ratio increases, the balance becomes increasingly poor. Also, the total computational rate (in units of MFlop/sec) is given for each phase in the final column of the phase summary table.

Total MFlop/sec = \(10^{-6} \times \text{(sum of flop over all processors)}/\text{(max time over all processors)}\)

Note: Total computational rates \(<1\) MFlop are listed as 0 in this column of the phase summary table. Additional statistics for each phase include the total number of messages sent, the average message length, and the number of global reductions.

```
mpiexec -n 4 ./ex10 -f0 medium -f1 arco6 -ksp_gmres_classicalgramschmidt -log_view -
-mat_type baij \n-matload_block_size 3 -pc_type bjacobi -options_left
```

--- [Overall summary, see part I] ---

--- Phase summary info: ---
Count: number of times phase was executed
Time and Flop/sec: Max - maximum over all processors
Ratio - ratio of maximum to minimum over all processors
Mess: number of messages sent
AvgLen: average message length
Reduct: number of global reductions
Global: entire computation
Stage: optional user-defined stages of a computation. Set stages with

   PetscLogStagePush() and PetscLogStagePop().

   %T - percent time in this phase
   %F - percent flop in this phase
   %M - percent messages in this phase
   %L - percent message lengths in this phase
   %R - percent reductions in this phase

Total MFlop/s: \(10^{-6} \times \text{(sum of flop over all processors)}/\text{(max time over all processors)}\)
As discussed in the preceding section, the performance summaries for higher-level PETSc routines include
the statistics for the lower levels of which they are made up. For example, the communication within matrix-vector products MatMult() consists of vector scatter operations, as given by the routines VecScatterBegin() and VecScatterEnd().

The final data presented are the percentages of the various statistics (time (%T), flop/sec (%F), messages(%M), average message length (%L), and reductions (%R)) for each event relative to the total computation and to any user-defined stages (discussed in Profiling Multiple Sections of Code). These statistics can aid in optimizing performance, since they indicate the sections of code that could benefit from various kinds of tuning. Hints for Performance Tuning gives suggestions about achieving good performance with PETSc codes.

The additional option -log_view_memory causes the display of additional columns of information about how much memory was allocated and freed during each logged event. This is useful to understand what phases of a computation require the most memory.

**Using -log_mpe with Jumpshot**

It is also possible to use the Jumpshot package [HL91] to visualize PETSc events. This package comes with the MPE software, which is part of the MPICH [Getal] implementation of MPI. The option

```
-log_mpe [logfile]
```

creates a logfile of events appropriate for viewing with Jumpshot. The user can either use the default logging file or specify a name via logfile. Events can be deactivated as described in Restricting Event Logging.

The user can also log MPI events. To do this, simply consider the PETSc application as any MPI application, and follow the MPI implementation’s instructions for logging MPI calls. For example, when using MPICH, this merely required adding -llmpich to the library list before -lmpich.

**Profiling Nested Events**

It is possible to output the PETSc logging information in a nested format where the hierarchy of events is explicit. This output can be generated either as an XML file or as a text file in a format suitable for viewing as a flame graph.

One can generate the XML output by passing the option -log_view :[logfile]:ascii_xml. It can be viewed by copying ${PETSC_DIR}/share/petsc/xml/performance_xml2html.xsl into the current directory, then opening the logfile in your browser.

The flame graph output can be generated with the option -log_view :[logfile]:ascii_flamegraph. It can then be visualised with either FlameGraph or speedscope. A flamegraph can be visualized directly from stdout using, for example, ImageMagick’s display utility <https://imagemagick.org/script/display.php>:

```
cd $PETSC_DIR/src/sys/tests
make ex30
mpiexec -n 2 ./ex30 -log_view ::ascii_flamegraph | flamegraph | display
```

Note that user-defined stages (see Profiling Multiple Sections of Code) will be ignored when using this nested format.
4.3.2 Profiling Application Codes

PETSc automatically logs object creation, times, and floating-point counts for the library routines. Users can easily supplement this information by profiling their application codes as well. The basic steps involved in logging a user-defined portion of code, called an event, are shown in the code fragment below:

```
PetscLogEvent USER_EVENT;
PetscClassId classid;
PetscLogDouble user_event_flops;

PetscClassIdRegister("class name", &classid);
PetscLogEventRegister("User event name", classid, &USER_EVENT);
/* code segment to monitor */
PetscLogFlops(user_event_flops);
PetscLogEventEnd(USER_EVENT, 0, 0, 0, 0);
```

One must register the event by calling `PetscLogEventRegister()`, which assigns a unique integer to identify the event for profiling purposes:

```
PetscLogEventRegister(const char string[], PetscClassId classid, PetscLogEvent *e);
```

Here `string` is a user-defined event name, and `color` is an optional user-defined event color (for use with `Jumpshot` logging; see `Using -log_mpe with Jumpshot`); one should see the manual page for details. The argument returned in `e` should then be passed to the `PetscLogEventBegin()` and `PetscLogEventEnd()` routines.

Events are logged by using the pair

```
PetscLogEventBegin(int event, PetscObject o1, PetscObject o2, PetscObject o3, PetscObject o4);
PetscLogEventEnd(int event, PetscObject o1, PetscObject o2, PetscObject o3, PetscObject o4);
```

The four objects are the PETSc objects that are most closely associated with the event. For instance, in a matrix-vector product they would be the matrix and the two vectors. These objects can be omitted by specifying 0 for `o1` - `o4`. The code between these two routine calls will be automatically timed and logged as part of the specified event.

The user can log the number of floating-point operations for this segment of code by calling

```
PetscLogFlops(number of flop for this code segment);
```

between the calls to `PetscLogEventBegin()` and `PetscLogEventEnd()`. This value will automatically be added to the global flop counter for the entire program.

4.3.3 Profiling Multiple Sections of Code

By default, the profiling produces a single set of statistics for all code between the `PetscInitialize()` and `PetscFinalize()` calls within a program. One can independently monitor several “stages” of code by switching among the various stages with the commands

```
PetscLogStagePush(PetscLogStage stage);
PetscLogStagePop();
```

see the manual pages for details. The command
The function `PetscLogStageRegister(const char *name, PetscLogStage *stage)` allows one to associate a name with a stage; these names are printed whenever summaries are generated with `-log_view`. The following code fragment uses three profiling stages within an program.

```c
PetscInitialize(int argc, char ***args, 0, 0);
/* stage 0 of code here */
PetscLogStageRegister("Stage 0 of Code", &stagenum0);
for (i=0; i<ntimes; i++) {
    PetscLogStageRegister("Stage 1 of Code", &stagenum1);
    PetscLogStagePush(stagenum1);
    /* stage 1 of code here */
    PetscLogStagePop();
    PetscLogStageRegister("Stage 2 of Code", &stagenum2);
    PetscLogStagePush(stagenum2);
    /* stage 2 of code here */
    PetscLogStagePop();
}
PetscFinalize();
```

The listings above show output generated by `-log_view` for a program that employs several profiling stages. In particular, this program is subdivided into six stages: loading a matrix and right-hand-side vector from a binary file, setting up the preconditioner, and solving the linear system; this sequence is then repeated for a second linear system. For simplicity, the second listing contains output only for stages 5 and 6 (linear solve of the second system), which comprise the part of this computation of most interest to us in terms of performance monitoring. This code organization (solving a small linear system followed by a larger system) enables generation of more accurate profiling statistics for the second system by overcoming the often considerable overhead of paging, as discussed in *Accurate Profiling and Paging Overheads*.

### 4.3.4 Restricting Event Logging

By default, all PETSc operations are logged. To enable or disable the PETSc logging of individual events, one uses the commands

```c
PetscLogEventActivate(PetscLogEvent event);
PetscLogEventDeactivate(PetscLogEvent event);
```

The `event` may be either a predefined PETSc event (as listed in the file `$PETSC_DIR/include/petsclog.h`) or one obtained with `PetscLogEventRegister()` (as described in *Profiling Application Codes*).

PETSc also provides routines that deactivate (or activate) logging for entire components of the library. Currently, the components that support such logging (de)activation are `Mat` (matrices), `Vec` (vectors), `KSP` (linear solvers, including `KSP` and `PC`), and `SNES` (nonlinear solvers):

```c
PetscLogEventDeactivateClass(MAT_CLASSID);
PetscLogEventDeactivateClass(KSP_CLASSID); /* includes PC and KSP */
PetscLogEventDeactivateClass(VEC_CLASSID);
PetscLogEventDeactivateClass(SNES_CLASSID);
```

and

```c
PetscLogEventActivateClass(MAT_CLASSID);
PetscLogEventActivateClass(KSP_CLASSID); /* includes PC and KSP */
PetscLogEventActivateClass(VEC_CLASSID);
PetscLogEventActivateClass(SNES_CLASSID);
```
Users can activate the printing of verbose information about algorithms, data structures, etc. to the screen by using the option `-info` or by calling `PetscInfoAllow(PETSC_TRUE)`. Such logging, which is used throughout the PETSc libraries, can aid the user in understanding algorithms and tuning program performance. For example, as discussed in *Sparse Matrices*, `-info` activates the printing of information about memory allocation during matrix assembly.

One can selectively turn off informative messages about any of the basic PETSc objects (e.g., `Mat`, `SNES`) with the command

```c
PetscInfoDeactivateClass(int object_classid)
```

where `object_classid` is one of `MAT_CLASSID`, `SNES_CLASSID`, etc. Messages can be reactivated with the command

```c
PetscInfoActivateClass(int object_classid)
```

Such deactivation can be useful when one wishes to view information about higher-level PETSc libraries (e.g., `TS` and `SNES`) without seeing all lower level data as well (e.g., `Mat`).

One can turn on or off logging for particular classes at runtime

```c
-info [filename][::<list,of,classnames>[:~]<list,self>]
```

The `list,of,classnames` is a list, separated by commas with no spaces, of classes one wishes to view the information on. For example `vec,ksp`. Information on all other classes will not be displayed. The ~ indicates to not display the list of classes but rather to display all other classes.

`self` indicates to display information on objects that are associated with `PETSC_COMM_SELF` while `~self` indicates to display information only for parallel objects.

See `PetscInfo()` for links to all the info operations that are available.

Application programmers can log their own messages, as well, by using the routine

```c
PetscInfo(void* obj, char* message,...)
```

where `obj` is the PETSc object associated most closely with the logging statement, `message`. For example, in the line search Newton methods, we use a statement such as

```c
PetscInfo(snes,"Cubic step, lambda %g\n",lambda);
```

### 4.3.6 Time

PETSc application programmers can access the wall clock time directly with the command

```c
PetscLogDouble time;
PetscCall(PetscTime(&time));
```

which returns the current time in seconds since the epoch, and is commonly implemented with `MPI_Wtime`. A floating point number is returned in order to express fractions of a second. In addition, as discussed in *Profiling Application Codes*, PETSc can automatically profile user-defined segments of code.
4.3.7 Saving Output to a File

All output from PETSc programs (including informative messages, profiling information, and convergence data) can be saved to a file by using the command line option -history [filename]. If no file name is specified, the output is stored in the file `${HOME}/.petschistory`. Note that this option only saves output printed with the PetscPrintf() and PetscFPrintf() commands, not the standard printf() and fprintf() statements.

4.3.8 Accurate Profiling and Paging Overheads

One factor that often plays a significant role in profiling a code is paging by the operating system. Generally, when running a program, only a few pages required to start it are loaded into memory rather than the entire executable. When the execution proceeds to code segments that are not in memory, a pagefault occurs, prompting the required pages to be loaded from the disk (a very slow process). This activity distorts the results significantly. (The paging effects are noticeable in the log files generated by -log_mpe, which is described in Using -log_mpe with Jumpshot.)

To eliminate the effects of paging when profiling the performance of a program, we have found an effective procedure is to run the exact same code on a small dummy problem before running it on the actual problem of interest. We thus ensure that all code required by a solver is loaded into memory during solution of the small problem. When the code proceeds to the actual (larger) problem of interest, all required pages have already been loaded into main memory, so that the performance numbers are not distorted.

When this procedure is used in conjunction with the user-defined stages of profiling described in Profiling Multiple Sections of Code, we can focus easily on the problem of interest. For example, we used this technique in the program KSP Tutorial ex10 to generate the timings within listing and listing. In this case, the profiled code of interest (solving the linear system for the larger problem) occurs within event stages 5 and 6. Interpreting -log_view Output: Parallel Performance provides details about interpreting such profiling data.

In particular, the macros

\begin{verbatim}
PetscPreLoadBegin(PetscBool flag, char* stagename)
PetscPreLoadStage(char *stagename)
\end{verbatim}

and

\begin{verbatim}
PetscPreLoadEnd()
\end{verbatim}

can be used to easily convert a regular PETSc program to one that uses preloading. The command line options -preload true and -preload false may be used to turn on and off preloading at run time for PETSc programs that use these macros.

4.3.9 NVIDIA Nsight Systems profiling

Nsight Systems will generate profiling data with a CUDA executable with the command nsys. For example, in serial

\begin{verbatim}
nsys profile -t nvtx,cuda -o file --stats=true --force-overwrite true ./a.out
\end{verbatim}

will generate a file file.qdstrm with performance data that is annotated with PETSc events (methods) and Kokkos device kernel names. The Nsight Systems GUI, nsys-ui, can be used to navigate this file (https://developer.nvidia.com/nsight-systems). The Nsight Systems GUI lets you see a timeline of code performance information like kernels, memory mallocs and frees, CPU-GPU communication, and high-level data like time, sizes of memory copies, and more, in a popup window when the mouse hovers over the section.
To view the data, start `nsys-ui` without any arguments and then Import the `.qdstrm` file in the GUI. A side effect of this viewing process is the generation of a file `file.nsys-rep`, which can be viewed directly with `nsys-ui` in the future.

For an MPI parallel job, only one process can call `nsys`, say have rank zero output `nsys` data and have all other ranks call the executable directly. For example with MPICH or Open MPI - we can run a parallel job on 4 MPI tasks as:

```bash
mpiexec -n 1 nsys profile -t nvtx,cuda -o file_name --stats=true --force-overwrite, _true ./a.out : -n 3 ./a.out
```

Note: The Nsight GUI can open profiling reports from elsewhere. For example, a report from a compute node can be analyzed on your local machine, but care should be taken to use the exact same versions of Nsight Systems that generated the report. To check the version of Nsight on the compute node run `nsys-ui` and note the version number at the top of the window.

### 4.3.10 Using TAU

TAU profiles can be generated without the need for instrumentation through the use of the perfstubs package. PETSc by default is configured with `--with-tau-perfstubs`. To generate profiles with TAU, first setup TAU:

```bash
wget http://tau.uoregon.edu/tau.tgz
./configure -cc=mpicc -c++=mpicxx -mpi -bfd=download -unwind=download && make install
export PATH=/tau_dir/x86_64/bin:$PATH
```

For more information on configuring TAU, see [http://tau.uoregon.edu](http://tau.uoregon.edu). Next, run your program with TAU. For instance, to profile `ex56`,

```bash
cd $PETSC_DIR/src/snes/tutorials
make ex56
mpirun -n 4 tau_exec -T mpi ./ex56 -log_perfstubs <args>
```

This should produce four `profile.*` files with profile data that can be viewed with `paraprof/pprof`:

```
Reading Profile files in profile.*

NODE 0; CONTEXT 0; THREAD 0:

<table>
<thead>
<tr>
<th></th>
<th>Exclusive</th>
<th>Inclusive</th>
<th>#Call</th>
<th>#Subrs</th>
<th>Inclusive</th>
</tr>
</thead>
<tbody>
<tr>
<td>%Time</td>
<td>msec</td>
<td>total msec</td>
<td></td>
<td>usec/call</td>
<td>Name</td>
</tr>
<tr>
<td>---</td>
<td>---</td>
<td>---</td>
<td>---</td>
<td>---</td>
<td>---</td>
</tr>
<tr>
<td>100.0</td>
<td>26</td>
<td>1,838</td>
<td>1</td>
<td>41322</td>
<td>1838424 .TAU application</td>
</tr>
<tr>
<td>73.2</td>
<td>1</td>
<td>1,345</td>
<td>2</td>
<td>168</td>
<td>672950 SNESolve</td>
</tr>
<tr>
<td>62.2</td>
<td>3</td>
<td>1,142</td>
<td>2</td>
<td>1282</td>
<td>571442 SNESJacobianEval</td>
</tr>
<tr>
<td>62.0</td>
<td>1,136</td>
<td>1,138</td>
<td>2</td>
<td>76</td>
<td>569494 DMPlexJacobianFE</td>
</tr>
<tr>
<td>60.1</td>
<td>0.046</td>
<td>1,105</td>
<td>1</td>
<td>32</td>
<td>1105041 Solve 1</td>
</tr>
<tr>
<td>15.2</td>
<td>87</td>
<td>279</td>
<td>5</td>
<td>11102</td>
<td>55943 Mesh Setup</td>
</tr>
<tr>
<td>13.2</td>
<td>0.315</td>
<td>241</td>
<td>1</td>
<td>32</td>
<td>241765 Solve 0</td>
</tr>
<tr>
<td>7.8</td>
<td>80</td>
<td>144</td>
<td>38785</td>
<td>38785</td>
<td>4 MPI_Allreduce()</td>
</tr>
<tr>
<td>7.0</td>
<td>69</td>
<td>128</td>
<td>6</td>
<td>43386</td>
<td>21491 DualSpaceSetUp</td>
</tr>
<tr>
<td>6.2</td>
<td>1</td>
<td>114</td>
<td>4</td>
<td>54</td>
<td>28536 PCSetUp</td>
</tr>
<tr>
<td>6.0</td>
<td>12</td>
<td>110</td>
<td>2</td>
<td>892</td>
<td>55407 PCSetUp_GAMG+</td>
</tr>
<tr>
<td>3.9</td>
<td>70</td>
<td>70</td>
<td>1</td>
<td>0</td>
<td>70888 MPI_Init_thread()</td>
</tr>
</tbody>
</table>
```

(continues on next page)
4.4 Hints for Performance Tuning

This chapter provides hints on how to get to achieve best performance with PETSc, particularly on distributed-memory machines with multiple CPU sockets per node. We focus on machine-related performance optimization here; algorithmic aspects like preconditioner selection are not the focus of this section.

4.4.1 Maximizing Memory Bandwidth

Most operations in PETSc deal with large datasets (typically vectors and sparse matrices) and perform relatively few arithmetic operations for each byte loaded or stored from global memory. Therefore, the arithmetic intensity expressed as the ratio of floating point operations to the number of bytes loaded and stored is usually well below unity for typical PETSc operations. On the other hand, modern CPUs are able to execute on the order of 10 floating point operations for each byte loaded or stored. As a consequence, almost all PETSc operations are limited by the rate at which data can be loaded or stored (memory bandwidth limited) rather than by the rate of floating point operations.

This section discusses ways to maximize the memory bandwidth achieved by applications based on PETSc. Where appropriate, we include benchmark results in order to provide quantitative results on typical performance gains one can achieve through parallelization, both on a single compute node and across nodes. In particular, we start with the answer to the common question of why performance generally does not increase 20-fold with a 20-core CPU.

Memory Bandwidth vs. Processes

Consider the addition of two large vectors, with the result written to a third vector. Because there are no dependencies across the different entries of each vector, the operation is embarrassingly parallel.

As Fig. 4.1 shows, the performance gains due to parallelization on different multi- and many-core CPUs quickly saturates. The reason is that only a fraction of the total number of CPU cores is required to saturate the memory channels. For example, a dual-socket system equipped with Haswell 12-core Xeon CPUs achieves more than 80 percent of achievable peak memory bandwidth with only four processes per socket (8 total), cf. Fig. 4.1. Consequently, running with more than 8 MPI ranks on such a system will not increase performance substantially. For the same reason, PETSc-based applications usually do not benefit from hyper-threading.

PETSc provides a simple way to measure memory bandwidth for different numbers of processes via the target make streams executed from $PETSC_DIR. The output provides an overview of the possible speedup one can obtain on the given machine (not necessarily a shared memory system). For example, the following is the most relevant output obtained on a dual-socket system equipped with two six-core-CPPUs with hyperthreading:

<table>
<thead>
<tr>
<th>np</th>
<th>speedup</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.0</td>
<td>1.0</td>
</tr>
<tr>
<td>1.58</td>
<td></td>
</tr>
</tbody>
</table>

(continues on next page)
Fig. 4.1: Memory bandwidth obtained on Intel hardware (dual socket except KNL) over the number of processes used. One can get close to peak memory bandwidth with only a few processes.

<table>
<thead>
<tr>
<th>Processes/Threads</th>
<th>Bandwidth (GB/sec)</th>
</tr>
</thead>
<tbody>
<tr>
<td>3</td>
<td>2.19</td>
</tr>
<tr>
<td>4</td>
<td>2.42</td>
</tr>
<tr>
<td>5</td>
<td>2.63</td>
</tr>
<tr>
<td>6</td>
<td>2.69</td>
</tr>
<tr>
<td>...</td>
<td></td>
</tr>
<tr>
<td>21</td>
<td>3.82</td>
</tr>
<tr>
<td>22</td>
<td>3.49</td>
</tr>
<tr>
<td>23</td>
<td>3.79</td>
</tr>
<tr>
<td>24</td>
<td>3.71</td>
</tr>
</tbody>
</table>

Estimation of possible speedup of MPI programs based on Streams benchmark.

It appears you have 1 node(s)

On this machine, one should expect a speed-up of typical memory bandwidth-bound PETSc applications of at most 4x when running multiple MPI ranks on the node. Most of the gains are already obtained when running with only 4-6 ranks. Because a smaller number of MPI ranks usually implies better preconditioners and better performance for smaller problems, the best performance for PETSc applications may be obtained with fewer ranks than there are physical CPU cores available.

Following the results from the above run of make streams, we recommend to use additional nodes instead of placing additional MPI ranks on the nodes. In particular, weak scaling (i.e. constant load per process, increasing the number of processes) and strong scaling (i.e. constant total work, increasing the number of processes) studies should keep the number of processes per node constant.
Non-Uniform Memory Access (NUMA) and Process Placement

CPUs in nodes with more than one CPU socket are internally connected via a high-speed fabric, cf. Fig. 4.2, to enable data exchange as well as cache coherency. Because main memory on modern systems is connected via the integrated memory controllers on each CPU, memory is accessed in a non-uniform way: A process running on one socket has direct access to the memory channels of the respective CPU, whereas requests for memory attached to a different CPU socket need to go through the high-speed fabric. Consequently, best aggregate memory bandwidth on the node is obtained when the memory controllers on each CPU are fully saturated. However, full saturation of memory channels is only possible if the data is distributed across the different memory channels.

Fig. 4.2: Schematic of a two-socket NUMA system. Processes should be spread across both CPUs to obtain full bandwidth.

Data in memory on modern machines is allocated by the operating system based on a first-touch policy. That is, memory is not allocated at the point of issuing `malloc()`, but at the point when the respective memory segment is actually touched (read or write). Upon first-touch, memory is allocated on the memory channel associated with the respective CPU the process is running on. Only if all memory on the respective CPU is already in use (either allocated or as IO cache), memory available through other sockets is considered.

Maximum memory bandwidth can be achieved by ensuring that processes are spread over all sockets in the respective node. For example, the recommended placement of a 8-way parallel run on a four-socket machine is to assign two processes to each CPU socket. To do so, one needs to know the enumeration of cores and pass the requested information to `mpirun`. Consider the hardware topology information returned by `lstopo` (part of the hwloc package) for the following two-socket machine, in which each CPU consists of six cores and supports hyperthreading:

<table>
<thead>
<tr>
<th>Machine (126GB total)</th>
</tr>
</thead>
<tbody>
<tr>
<td>NUMANode L#0 (P#0 63GB)</td>
</tr>
<tr>
<td>Package L#0 + L3 L#0 (15MB)</td>
</tr>
<tr>
<td>L2 L#0 (256KB) + L1d L#0 (32KB) + L1i L#0 (32KB) + Core L#0</td>
</tr>
<tr>
<td>PU L#0 (P#0)</td>
</tr>
<tr>
<td>PU L#1 (P#12)</td>
</tr>
<tr>
<td>L2 L#1 (256KB) + L1d L#1 (32KB) + L1i L#1 (32KB) + Core L#1</td>
</tr>
<tr>
<td>PU L#2 (P#1)</td>
</tr>
<tr>
<td>PU L#3 (P#13)</td>
</tr>
<tr>
<td>L2 L#2 (256KB) + L1d L#2 (32KB) + L1i L#2 (32KB) + Core L#2</td>
</tr>
<tr>
<td>PU L#4 (P#2)</td>
</tr>
<tr>
<td>PU L#5 (P#14)</td>
</tr>
<tr>
<td>L2 L#3 (256KB) + L1d L#3 (32KB) + L1i L#3 (32KB) + Core L#3</td>
</tr>
<tr>
<td>PU L#6 (P#3)</td>
</tr>
<tr>
<td>PU L#7 (P#15)</td>
</tr>
<tr>
<td>L2 L#4 (256KB) + L1d L#4 (32KB) + L1i L#4 (32KB) + Core L#4</td>
</tr>
<tr>
<td>PU L#8 (P#4)</td>
</tr>
<tr>
<td>PU L#9 (P#16)</td>
</tr>
<tr>
<td>L2 L#5 (256KB) + L1d L#5 (32KB) + L1i L#5 (32KB) + Core L#5</td>
</tr>
<tr>
<td>PU L#10 (P#5)</td>
</tr>
<tr>
<td>PU L#11 (P#17)</td>
</tr>
<tr>
<td>NUMANode L#1 (P#1 63GB)</td>
</tr>
<tr>
<td>Package L#1 + L3 L#1 (15MB)</td>
</tr>
</tbody>
</table>

(continues on next page)
The relevant physical processor IDs are shown in parentheses prefixed by \texttt{P#}. Here, IDs 0 and 12 share the same physical core and have a common L2 cache. IDs 0, 12, 1, 13, 2, 14, 3, 15, 4, 16, 5, 17 share the same socket and have a common L3 cache.

A good placement for a run with six processes is to locate three processes on the first socket and three processes on the second socket. Unfortunately, mechanisms for process placement vary across MPI implementations, so make sure to consult the manual of your MPI implementation. The following discussion is based on how processor placement is done with MPICH and Open MPI, where one needs to pass \texttt{--bind-to core --map-by socket} to \texttt{mpirun}:

\begin{verbatim}
$ mpirun -n 6 --bind-to core --map-by socket ./stream
process 0 binding: 100000000000100000000000
process 1 binding: 010000000000010000000000
process 2 binding: 001000000000001000000000
process 3 binding: 000100000000000100000000
process 4 binding: 000010000000000010000000
process 5 binding: 000001000000000001000000
Triad: 45403.1949 Rate (MB/s)
\end{verbatim}

In this configuration, process 0 is bound to the first physical core on the first socket (with IDs 0 and 12), process 1 is bound to the first core on the second socket (IDs 6 and 18), and similarly for the remaining processes. The achieved bandwidth of 45 GB/sec is close to the practical peak of about 50 GB/sec available on the machine. If, however, all MPI processes are located on the same socket, memory bandwidth drops significantly:

\begin{verbatim}
$ mpirun -n 6 --bind-to core --map-by core ./stream
process 0 binding: 100000000000100000000000
process 1 binding: 010000000000010000000000
process 2 binding: 001000000000001000000000
process 3 binding: 000100000000000100000000
process 4 binding: 000010000000000010000000
process 5 binding: 000001000000000001000000
Triad: 25510.7507 Rate (MB/s)
\end{verbatim}

All processes are now mapped to cores on the same socket. As a result, only the first memory channel is fully saturated at 25.5 GB/sec.

\subsection{4.4. Hints for Performance Tuning}
One must not assume that `mpirun` uses good defaults. To demonstrate, compare the full output of `make streams` from *Memory Bandwidth vs. Processes* on the left with the results on the right obtained by passing `-bind-to core --map-by socket`:

```
$ make streams
np  speedup
  1  1.0
  2  1.58
  3  2.19
  4  2.42
  5  2.63
  6  2.69
  7  2.31
  8  2.42
  9  2.37
 10  2.65
 11  2.3
 12  2.53
 13  2.43
 14  2.63
 15  2.74
 16  2.7
 17  3.28
 18  3.66
 19  3.95
 20  3.07
 21  3.82
 22  3.49
 23  3.79
 24  3.71
```

```
$ make streams
np  speedup
  1  1.0
  2  1.59
  3  2.66
  4  3.5
  5  3.56
  6  4.23
  7  3.95
  8  4.39
  9  4.09
 10  4.46
 11  4.15
 12  4.42
 13  3.71
 14  3.83
 15  4.08
 16  4.22
 17  4.18
 18  4.31
 19  4.22
 20  4.28
 21  4.25
 22  4.23
 23  4.28
```

(continues on next page)
For the non-optimized version on the left, the speedup obtained when using any number of processes between 3 and 13 is essentially constant up to fluctuations, indicating that all processes were by default executed on the same socket. Only with 14 or more processes, the speedup number increases again. In contrast, the results of `make streams` with proper processor placement shown on the right resulted in slightly higher overall parallel speedup (identical baselines), in smaller performance fluctuations, and more than 90 percent of peak bandwidth with only six processes.

Machines with job submission systems such as SLURM usually provide similar mechanisms for processor placements through options specified in job submission scripts. Please consult the respective manuals.

**Additional Process Placement Considerations and Details**

For a typical, memory bandwidth-limited PETSc application, the primary consideration in placing MPI processes is ensuring that processes are evenly distributed among sockets, and hence using all available memory channels. Increasingly complex processor designs and cache hierarchies, however, mean that performance may also be sensitive to how processes are bound to the resources within each socket. Performance on the two processor machine in the preceding example may be relatively insensitive to such placement decisions, because one L3 cache is shared by all cores within a NUMA domain, and each core has its own L2 and L1 caches. However, processors that are less “flat”, with more complex hierarchies, may be more sensitive. In many AMD Opterons or the second-generation “Knights Landing” Intel Xeon Phi, for instance, L2 caches are shared between two cores. On these processors, placing consecutive MPI ranks on cores that share the same L2 cache may benefit performance if the two ranks communicate frequently with each other, because the latency between cores sharing an L2 cache may be roughly half that of two cores not sharing one. There may be benefit, however, in placing consecutive ranks on cores that do not share an L2 cache, because (if there are fewer MPI ranks than cores) this increases the total L2 cache capacity and bandwidth available to the application. There is a trade-off to be considered between placing processes close together (in terms of shared resources) to optimize for efficient communication and synchronization vs. farther apart to maximize available resources (memory channels, caches, I/O channels, etc.), and the best strategy will depend on the application and the software and hardware stack.

Different process placement strategies can affect performance at least as much as some commonly explored settings, such as compiler optimization levels. Unfortunately, exploration of this space is complicated by two factors: First, processor and core numberings may be completely arbitrary, changing with BIOS version, etc., and second—as already noted—there is no standard mechanism used by MPI implementations (or job schedulers) to specify process affinity. To overcome the first issue, we recommend using the `lstopo` utility of the Portable Hardware Locality (`hwloc`) software package (which can be installed by configuring PETSc with `--download-hwloc`) to understand the processor topology of your machine. We cannot fully address the second issue—consult the documentation for your MPI implementation and/or job scheduler—but we offer some general observations on understanding placement options:

- An MPI implementation may support a notion of domains in which a process may be pinned. A domain may simply correspond to a single core; however, the MPI implementation may allow a deal of flexibility in specifying domains that encompass multiple cores, span sockets, etc. Some implementations, such as Intel MPI, provide means to specify whether domains should be “compact”—composed of cores sharing resources such as caches—or “scatter”-ed, with little resource sharing (possibly even spanning sockets).
Separate from the specification of domains, MPI implementations often support different orderings in which MPI ranks should be bound to these domains. Intel MPI, for instance, supports “compact” ordering to place consecutive ranks close in terms of shared resources, “scatter” to place them far apart, and “bunch” to map proportionally to sockets while placing ranks as close together as possible within the sockets.

An MPI implementation that supports process pinning should offer some way to view the rank assignments. Use this output in conjunction with the topology obtained via `lstopo` or a similar tool to determine if the placements correspond to something you believe is reasonable for your application. Do not assume that the MPI implementation is doing something sensible by default!

### 4.4.2 Performance Pitfalls and Advice

This section looks into a potpourri of performance pitfalls encountered by users in the past. Many of these pitfalls require a deeper understanding of the system and experience to detect. The purpose of this section is to summarize and share our experience so that these pitfalls can be avoided in the future.

#### Debug vs. Optimized Builds

PETSc’s `configure` defaults to building PETSc with debug mode enabled. Any code development should be done in this mode, because it provides handy debugging facilities such as accurate stack traces, memory leak checks, and memory corruption checks. Note that PETSc has no reliable way of knowing whether a particular run is a production or debug run. In the case that a user requests profiling information via `-log_view`, a debug build of PETSc issues the following warning:

```
##########################################################
# WARNING!!!                                         #
# This code was compiled with a debugging option,     #
# To get timing results run configure                 #
# using --with-debugging=no, the performance will     #
# be generally two or three times faster.             #
#                                                   
##########################################################
```

Conversely, one way of checking whether a particular build of PETSc has debugging enabled is to inspect the output of `-log_view`.

Debug mode will generally be most useful for code development if appropriate compiler options are set to facilitate debugging. The compiler should be instructed to generate binaries with debug symbols (command line option `-g` for most compilers), and the optimization level chosen should either completely disable optimizations (`-O0` for most compilers) or enable only optimizations that do not interfere with debugging (GCC, for instance, supports a `-Og` optimization level that does this).

Only once the new code is thoroughly tested and ready for production, one should disable debugging facilities by passing `--with-debugging=no` to `configure`. One should also ensure that an appropriate compiler optimization level is set. Note that some compilers (e.g., Intel) default to fairly comprehensive optimization levels, while others (e.g., GCC) default to no optimization at all. The best optimization flags will depend on your code, the compiler, and the target architecture, but we offer a few guidelines for finding those that will offer the best performance:

- Most compilers have a number of optimization levels (with level n usually specified via `-On`) that provide a quick way to enable sets of several optimization flags. We suggest trying the higher optimization levels (the highest level is not guaranteed to produce the fastest executable, so some experimentation
may be merited). With most recent processors now supporting some form of SIMD or vector instructions, it is important to choose a level that enables the compiler’s auto-vectorizer; many compilers do not enable auto-vectorization at lower optimization levels (e.g., GCC does not enable it below `-O3` and the Intel compiler does not enable it below `-O2`).

- For processors supporting newer vector instruction sets, such as Intel AVX2 and AVX-512, it is also important to direct the compiler to generate code that targets these processors (e.g., `-march=native`); otherwise, the executables built will not utilize the newer instructions sets and will not take advantage of the vector processing units.

- Beyond choosing the optimization levels, some value-unsafe optimizations (such as using reciprocals of values instead of dividing by those values, or allowing re-association of operands in a series of calculations) for floating point calculations may yield significant performance gains. Compilers often provide flags (e.g., `-ffast-math` in GCC) to enable a set of these optimizations, and they may be turned on when using options for very aggressive optimization (`-fast` or `-Ofast` in many compilers). These are worth exploring to maximize performance, but, if employed, it important to verify that these do not cause erroneous results with your code, since calculations may violate the IEEE standard for floating-point arithmetic.

**Profiling**

Users should not spend time optimizing a code until after having determined where it spends the bulk of its time on realistically sized problems. As discussed in detail in *Profiling*, the PETSc routines automatically log performance data if certain runtime options are specified.

To obtain a summary of where and how much time is spent in different sections of the code, use one of the following options:

- Run the code with the option `-log_view` to print a performance summary for various phases of the code.
- Run the code with the option `-log_mpe [logfilename]`, which creates a logfile of events suitable for viewing with Jumpshot (part of MPICH).

Then, focus on the sections where most of the time is spent. If you provided your own callback routines, e.g. for residual evaluations, search the profiling output for routines such as `SNESFunctionEval` or `SNESJacobianEval`. If their relative time is significant (say, more than 30 percent), consider optimizing these routines first. Generic instructions on how to optimize your callback functions are difficult; you may start by reading performance optimization guides for your system’s hardware.

**Aggregation**

Performing operations on chunks of data rather than a single element at a time can significantly enhance performance because of cache reuse or lower data motion. Typical examples are:

- Insert several (many) elements of a matrix or vector at once, rather than looping and inserting a single value at a time. In order to access elements in of vector repeatedly, employ `VecGetArray()` to allow direct manipulation of the vector elements.
- When possible, use `VecMDot()` rather than a series of calls to `VecDot()`.
- If you require a sequence of matrix-vector products with the same matrix, consider packing your vectors into a single matrix and use matrix-matrix multiplications.
- Users should employ a reasonable number of `PetscMalloc()` calls in their codes. Hundreds or thousands of memory allocations may be appropriate; however, if tens of thousands are being used, then reducing the number of `PetscMalloc()` calls may be warranted. For example, reusing space
or allocating large chunks and dividing it into pieces can produce a significant savings in allocation overhead. *Data Structure Reuse* gives details.

Aggressive aggregation of data may result in inflexible data structures and code that is hard to maintain. We advise users to keep these competing goals in mind and not blindly optimize for performance only.

**Memory Allocation for Sparse Matrix Factorization**

When symbolically factoring an AIJ matrix, PETSc has to guess how much fill there will be. Careful use of the fill parameter in the *MatFactorInfo* structure when calling *MatLUFactorSymbolic()* or *MatILUFactorSymbolic()* can reduce greatly the number of mallocs and copies required, and thus greatly improve the performance of the factorization. One way to determine a good value for the fill parameter is to run a program with the option `-info`. The symbolic factorization phase will then print information such as:

```
Info:MatILUFactorSymbolic_SeqAIJ:Reallocs 12 Fill ratio:given 1 needed 2.16423
```

This indicates that the user should have used a fill estimate factor of about 2.17 (instead of 1) to prevent the 12 required mallocs and copies. The command line option `-pc_factor_fill 2.17`

will cause PETSc to preallocate the correct amount of space for the factorization.

**Detecting Memory Allocation Problems and Memory Usage**

PETSc provides tools to aid in understanding PETSc memory usage and detecting problems with memory allocation, including leaks and use of uninitialized space. Internally, PETSc uses the routines *PetscMalloc()* and *PetscFree()* for memory allocation; instead of directly calling *malloc()* and *free()*.

This allows PETSc to track its memory usage and perform error checking. Users are urged to use these routines as well when appropriate.

- The option `-malloc_debug` turns on PETSc’s extensive runtime error checking of memory for corruption. This checking can be expensive, so should not be used for production runs. The option `-malloc_test` is equivalent to `-malloc_debug` but only works when PETSc is configured with `--with-debugging` (the default configuration). We suggest setting the environmental variable `PETSC_OPTIONS=-malloc_test` in your shell startup file to automatically enable runtime check memory for developing code but not running optimized code. Using `-malloc_debug` or `-malloc_test` for large runs can slow them significantly, thus we recommend turning them off if you code is painfully slow and you don’t need the testing. In addition, you can use `-check_pointer_intensity 0` for long run debug runs that do not need extensive memory corruption testing. This option is occasionally added to the `PETSC_OPTIONS` environmental variable by some users.

- The option `-malloc_dump` will print a list of memory locations that have not been freed at the conclusion of a program. If all memory has been freed no message is printed. Note that the option `-malloc_dump` activates a call to *PetscMallocDump()* during *PetscFinalize()*. The user can also call *PetscMallocDump()* elsewhere in a program.

- Another useful option is `-malloc_view`, which reports memory usage in all routines at the conclusion of the program. Note that this option activates logging by calling *PetscMallocViewSet()* in *PetscInitialize()* and then prints the log by calling *PetscMallocView()* in *PetscFinalize()*.

- When finer granularity is desired, the user can call *PetscMallocGetCurrentUsage()* and *PetscMallocGetMaximumUsage()* for memory allocated by PETSc, or *PetscMemoryGetCurrentUsage()* and *PetscMemoryGetMaximumUsage()* for the total memory used by the program.
Note that *PetscMemorySetGetMaximumUsage()* must be called before *PetscMemoryGetMaximumUsage()* (typically at the beginning of the program).

- The option `-memory_view` provides a high-level view of all memory usage, not just the memory used by *PetscMalloc()* , at the conclusion of the program.
- When running with `-log_view`, the additional option `-log_view_memory` causes the display of additional columns of information about how much memory was allocated and freed during each logged event. This is useful to understand what phases of a computation require the most memory.

One can also use Valgrind to track memory usage and find bugs, see FAQ: Valgrind usage.

### Data Structure Reuse

Data structures should be reused whenever possible. For example, if a code often creates new matrices or vectors, there often may be a way to reuse some of them. Very significant performance improvements can be achieved by reusing matrix data structures with the same nonzero pattern. If a code creates thousands of matrix or vector objects, performance will be degraded. For example, when solving a nonlinear problem or timestepping, reusing the matrices and their nonzero structure for many steps when appropriate can make the code run significantly faster.

A simple technique for saving work vectors, matrices, etc. is employing a user-defined context. In C and C++ such a context is merely a structure in which various objects can be stashed; in Fortran a user context can be an integer array that contains both parameters and pointers to PETSc objects. See SNES Tutorial ex5 and SNES Tutorial ex5f90 for examples of user-defined application contexts in C and Fortran, respectively.

### Numerical Experiments

PETSc users should run a variety of tests. For example, there are a large number of options for the linear and nonlinear equation solvers in PETSc, and different choices can make a very big difference in convergence rates and execution times. PETSc employs defaults that are generally reasonable for a wide range of problems, but clearly these defaults cannot be best for all cases. Users should experiment with many combinations to determine what is best for a given problem and customize the solvers accordingly.

- Use the options `-snes_view`, `-ksp_view`, etc. (or the routines KSPView(), SNESView(), etc.) to view the options that have been used for a particular solver.
- Run the code with the option `-help` for a list of the available runtime commands.
- Use the option `-info` to print details about the solvers’ operation.
- Use the PETSc monitoring discussed in Profiling to evaluate the performance of various numerical methods.

### Tips for Efficient Use of Linear Solvers

As discussed in KSP: Linear System Solvers, the default linear solvers are

- uniprocess: GMRES(30) with ILU(0) preconditioning
- multiprocess: GMRES(30) with block Jacobi preconditioning, where there is 1 block per process, and each block is solved with ILU(0)

One should experiment to determine alternatives that may be better for various applications. Recall that one can specify the KSP methods and preconditioners at runtime via the options:

```
-ksp_type <ksp_name> -pc_type <pc_name>
```
One can also specify a variety of runtime customizations for the solvers, as discussed throughout the manual. In particular, note that the default restart parameter for GMRES is 30, which may be too small for some large-scale problems. One can alter this parameter with the option `-ksp_gmres_restart <restart>` or by calling `KSPGMRESSetRestart()`. *Krylov Methods* gives information on setting alternative GMRES orthogonalization routines, which may provide much better parallel performance.

For elliptic problems one often obtains good performance and scalability with multigrid solvers. Consult *Algebraic Multigrid (AMG) Preconditioners* for available options. Our experience is that GAMG works particularly well for elasticity problems, whereas hypre does well for scalar problems.

**System-Related Problems**

The performance of a code can be affected by a variety of factors, including the cache behavior, other users on the machine, etc. Below we briefly describe some common problems and possibilities for overcoming them.

- **Problem too large for physical memory size**: When timing a program, one should always leave at least a ten percent margin between the total memory a process is using and the physical size of the machine’s memory. One way to estimate the amount of memory used by given process is with the Unix `getrusage` system routine. The PETSc option `-malloc_view` reports all memory usage, including any Fortran arrays in an application code.

- **Effects of other users**: If other users are running jobs on the same physical processor nodes on which a program is being profiled, the timing results are essentially meaningless.

- **Overhead of timing routines on certain machines**: On certain machines, even calling the system clock in order to time routines is slow; this skews all of the flop rates and timing results. The file `$PETSC_DIR/src/benchmarks/PetscTime.c` contains a simple test problem that will approximate the amount of time required to get the current time in a running program. On good systems it will on the order of $10^{-6}$ seconds or less.

- **Problem too large for good cache performance**: Certain machines with lower memory bandwidths (slow memory access) attempt to compensate by having a very large cache. Thus, if a significant portion of an application fits within the cache, the program will achieve very good performance; if the code is too large, the performance can degrade markedly. To analyze whether this situation affects a particular code, one can try plotting the total flop rate as a function of problem size. If the flop rate decreases rapidly at some point, then the problem may likely be too large for the cache size.

- **Inconsistent timings**: Inconsistent timings are likely due to other users on the machine, thrashing (using more virtual memory than available physical memory), or paging in of the initial executable. *Accurate Profiling and Paging Overheads* provides information on overcoming paging overhead when profiling a code. We have found on all systems that if you follow all the advise above your timings will be consistent within a variation of less than five percent.

### 4.5 The Use of BLAS and LAPACK in PETSc and external libraries

1. BLAS 1 operations (and GPU equivalents) - vector operations such as `VecNorm()`, `VecAXPY()`, and `VecScale()` are used extensively in PETSc. Depending on the simulation the size of the vectors may be from hundreds of entries to many millions.

2. BLAS 2 operations - dense matrix with vector operations, generally the dense matrices are very small.

3. Eigenvalue and SVD computations, generally for very small matrices
4. External packages such as MUMPS and SuperLU_DIST use BLAS 3 operations (and possibly BLAS 1 and 2). The dense matrices may be of modest size, going up to thousands of rows and columns.

For most PETSc simulations (that is not using certain external packages) using an optimized set of BLAS/LAPACK routines only provides a modest improvement in performance. For some external packages using optimized BLAS/LAPACK can make a dramatic improvement in performance.

4.5.1 32 or 64-bit BLAS/LAPACK integers

BLAS/LAPACK libraries may use 32 or 64-bit integers. PETSc configure and compile handles this automatically so long as the arguments to the BLAS/LAPACK routines are set to the type PetscBLASInt. The routine PetscBLASIntCast(PetscInt, PetscBLASInt *) casts a PetscInt to the BLAS/LAPACK size. If the BLAS/LAPACK size is not large enough it generates an error. For the vast majority of simulations, even very large ones, 64-bit BLAS/LAPACK integers are not needed, even when 64-bit PETSc integers are used.

The configure option --with-64-bit-blas-indices attempts to locate and use a 64-bit integer version of BLAS/LAPACK library. Except for MKL Cluster PARDISO, most external packages do not support using 64-bit BLAS/LAPACK integers so if you are using such packages you cannot use 64-bit BLAS/LAPACK integers.

The configure options --with-64-bit-indices and --with-64-bit-blas-indices are independent. --with-64-bit-indices does not imply that the BLAS/LAPACK libraries use 64 bit indices.

4.5.2 Shared memory BLAS/LAPACK parallelism

Some BLAS/LAPACK libraries can make use of shared memory parallelism within the function calls, generally using OpenMP, or possibly PThreads. If this feature is turned on, it is in addition to the MPI based parallelism that PETSc is using. Thus it can result in over-subscription of hardware resources. For example, if a system has 16 cores and PETSc is run with an MPI size of 16 then each core is assigned an MPI process. But if the BLAS/LAPACK is running with OpenMP and 4 threads per process this results in 64 threads competing to use 16 cores which will perform poorly.

If one elects to use both MPI parallelism and shared memory BLAS/LAPACK parallelism one should ensure they do not over subscribe the hardware resources. Since PETSc does not natively use OpenMP this means that phases of the computation that do not use BLAS/LAPACK will be under-subscribed, thus under-utilizing the system. For PETSc simulations which do not use external packages there is generally no benefit to using parallel BLAS/LAPACK. The environmental variable OMP_NUM_THREADS can be used to set the number of threads used by each MPI process for its shared memory parallel BLAS/LAPACK. The additional environmental variables OMP_PROC_BIND and OMP_PLACES may also need to be set appropriately for the system to obtain good parallel performance with BLAS/LAPACK. The configure option --with-openmp will trigger PETSc to try to locate and use a parallel BLAS/LAPACK library.

Certain external packages such as MUMPS may benefit from using parallel BLAS/LAPACK operations. See the manual page MATSOLVERMUMPS for details on how one can restrict the number of MPI processes while running MUMPS to utilize parallel BLAS/LAPACK.
4.5.3 Available BLAS/LAPACK libraries

Most systems (besides Microsoft Windows) come with pre-installed BLAS/LAPACK which are satisfactory for many PETSc simulations.

The freely available Intel MKL mathematics libraries provide BLAS/LAPACK that are generally better performing than the system provided libraries and are generally fine for most users.

For systems that do not provide BLAS/LAPACK, such as Microsoft Windows, PETSc provides the Fortran reference version --download-fblaslapack and a f2c generated C version --download-f2cblaslapack (which also supports 128 bit real number computations). These libraries are less optimized but useful to get started with PETSc easily.

PETSc also provides access to OpenBLAS via the --download-openblas configure option. OpenBLAS uses some highly optimized operations but falls back on reference routines for many other operations. See the OpenBLAS manual for more information. The configure option --download-openblas provides a full BLAS/LAPACK implementation.

BLIS does not bundle LAPACK with it so PETSc’s configure attempts to locate a compatible system LAPACK library to use if --download-blis is selected. One can use --download-f2cblaslapack --download-blis. This is recommended as a portable high-performance option.

4.6 Other PETSc Features

4.6.1 PETSc on a process subset

Users who wish to employ PETSc on only a subset of MPI processes within a larger parallel job, or who wish to use a “manager” process to coordinate the work of “worker” PETSc processes, should specify an alternative communicator for PETSC_COMM_WORLD by directly setting its value, for example to use an existing MPI communicator comm,

```
PETSC_COMM_WORLD = comm; /* To use a previously-defined MPI_Comm */
```

before calling PetscInitialize(), but, obviously, after calling MPI_Init().

4.6.2 Runtime Options

Allowing the user to modify parameters and options easily at runtime is very desirable for many applications. PETSc provides a simple mechanism to enable such customization. To print a list of available options for a given program, simply specify the option -help at runtime, e.g.,

```
$ mpiexec -n 1 ./ex1 -help
```

Note that all runtime options correspond to particular PETSc routines that can be explicitly called from within a program to set compile-time defaults. For many applications it is natural to use a combination of compile-time and runtime choices. For example, when solving a linear system, one could explicitly specify use of the Krylov subspace solver BiCGStab by calling

```
KSPSetType(ksp, KSPBCGS);
```

One could then override this choice at runtime with the option

```
-ksp_type tfqmr
```
to select the Transpose-Free QMR algorithm. (See *KSP: Linear System Solvers* for details.)

The remainder of this section discusses details of runtime options.

### The Options Database

Each PETSc process maintains a database of option names and values (stored as text strings). This database is generated with the command `PetscInitialize()`, which is listed below in its C/C++ and Fortran variants, respectively:

```c
PetscInitialize(int *argc, char ***args, const char *file, const char *help); // C
```

```fortran
call PetscInitialize(integer ierr) ! Fortran
```

The arguments `argc` and `args` (in the C/C++ version only) are the addresses of the usual command line arguments, while the `file` is a name of an optional file that can contain additional options. By default this file is called `.petscrc` in the user’s home directory. The user can also specify options via the environmental variable `PETSC_OPTIONS`. The options are processed in the following order:

1. file
2. environmental variable
3. command line

Thus, the command line options supersede the environmental variable options, which in turn supersede the options file.

The file format for specifying options is

```
-<optionname> possible_value
-<anotheroptionname> possible_value
...
```

All of the option names must begin with a dash (-) and have no intervening spaces. Note that the option values cannot have intervening spaces either, and tab characters cannot be used between the option names and values. For uniformity throughout PETSc, we employ the format `-[prefix_]package_option` (for instance, `-ksp_type`, `-mat_view ::info`, or `-mg_levels_ksp_type`).

Users can specify an alias for any option name (to avoid typing the sometimes lengthy default name) by adding an alias to the `.petsrcrc` file in the format

```
alias -newname -oldname
```

For example,

```
alias -kspt -ksp_type
alias -sd -start_in_debugger
```

Comments can be placed in the `.petsrcrc` file by using `#` in the first column of a line.
Options Prefixes

Options prefixes allow specific objects to be controlled from the options database. For instance, `PCMG` gives prefixes to its nested `KSP` objects; one may control the coarse grid solver by adding the `mg_coarse` prefix, for example `-mg_coarse_ksp_type preonly`. One may also use `KSPSetOptionsPrefix()`, `DMSetOptionsPrefix()`, `SNESSetOptionsPrefix()`, `TSSetOptionsPrefix()`, and similar functions to assign custom prefixes, useful for applications with multiple or nested solvers.

Adding options from a file

PETSc can load additional options from a file using `PetscOptionsInsertFile()`, which can also be used from the command line, e.g. `-options_file my_options.opts`.

One can also use YAML files with `PetscOptionsInsertFileYAML()`. For example, the following file:

```yaml
$$
$&ans: &ans 42
$&eu: &eu 2.72
$&pi: &pi 3.14

opt:
  bool: true
  int: *
  real: *
  imag: 2.72i
  cmplx: -3.14+2.72i
  str: petsc

$&seq-bool [true, false]
$&seq-int [123, 456, 789]
$&seq-real [*pi, *eu]
$&seq-str [abc, ijk, fgh]

seq1: {
  bool: *
  int: *
  real: *
  str: *
}

seq2: {
  bool:
    - true
    - false
  int:
    - 123
    - 456
    - 789
  real:
    - *
    - *
  str:
    - *rst
    - *uvw
$$
```

(continues on next page)
map:
  - key0: 0
  - key1: 1
  - key2: 2
  - $$: ignored
  - $$tail: ignored

corresponds to the following PETSc options:

```plaintext
-map key0,key1,key2 # (source: file)
-map_key0 0 # (source: file)
-map_key1 1 # (source: file)
-map_key2 2 # (source: file)
-opt_bool true # (source: file)
-opt_cmplx -3.14+2.72i # (source: file)
-opt_imag 2.72i # (source: file)
-opt_int 42 # (source: file)
-opt_real 3.14 # (source: file)
-opt_str petsc # (source: file)
-seq1_bool true,false # (source: file)
-seq1_int 123,456,789 # (source: file)
-seq1_real 3.14,2.72 # (source: file)
-seq1_str abc,ijk,fgh # (source: file)
-seq2_bool true,false # (source: file)
-seq2_int 123,456,789 # (source: file)
-seq2_real 3.14,2.72 # (source: file)
-seq2_str rst,uvw,xyz # (source: file)
```

With `-options_file`, PETSc will parse the file as YAML if it ends in a standard YAML or JSON extension or if one uses a `.yaml` postfix, e.g. `-options_file my_options.yaml` or `-options_file my_options.txt:yaml`

PETSc will also check the first line of the options file itself and parse the file as YAML if it matches certain criteria, for example.

```yaml
---
name: value
```
User-Defined PetscOptions

Any subroutine in a PETSc program can add entries to the database with the command

```c
PetscOptionsSetValue(PetscOptions options, char *name, char *value);
```

though this is rarely done. To locate options in the database, one should use the commands

```c
PetscOptionsHasName(PetscOptions options, char *pre, char *name, PetscBool *flg);
PetscOptionsGetInt(PetscOptions options, char *pre, char *name, PetscInt *value, PetscBool *flg);
PetscOptionsGetReal(PetscOptions options, char *pre, char *name, PetscReal *value, PetscBool *flg);
PetscOptionsGetString(PetscOptions options, char *pre, char *name, char *value, size_t maxlen, PetscBool *flg);
PetscOptionsGetStringArray(PetscOptions options, char *pre, char *name, char **values, PetscInt *nmax, PetscBool *flg);
PetscOptionsGetIntArray(PetscOptions options, char *pre, char *name, PetscInt *value, PetscInt *nmax, PetscBool *flg);
PetscOptionsGetRealArray(PetscOptions options, char *pre, char *name, PetscReal *value, PetscInt *nmax, PetscBool *flg);
```

All of these routines set flg=PETSC_TRUE if the corresponding option was found, flg=PETSC_FALSE if it was not found. The optional argument pre indicates that the true name of the option is the given name (with the dash “-” removed) prepended by the prefix pre. Usually pre should be set to NULL (or PETSC_NULL_CHARACTER for Fortran); its purpose is to allow someone to rename all the options in a package without knowing the names of the individual options. For example, when using block Jacobi preconditioning, the KSP and PC methods used on the individual blocks can be controlled via the options -sub_ksp_type and -sub_pc_type.

Keeping Track of Options

One useful means of keeping track of user-specified runtime options is use of -options_view, which prints to stdout during PetscFinalize() a table of all runtime options that the user has specified. A related option is -options_left, which prints the options table and indicates any options that have not been requested upon a call to PetscFinalize(). This feature is useful to check whether an option has been activated for a particular PETSc object (such as a solver or matrix format), or whether an option name may have been accidentally misspelled.

4.6.3 Viewers: Looking at PETSc Objects

PETSc employs a consistent scheme for examining, printing, and saving objects through commands of the form

```c
XXXView(XXX obj, PetscViewer viewer);
```

Here obj is a PETSc object of type XXX, where XXX is Mat, Vec, SNES, etc. There are several predefined viewers.

- Passing in a zero (0) for the viewer causes the object to be printed to the screen; this is useful when viewing an object in a debugger but should be avoided in source code.
- PETSC_VIEWER_STDOUT_SELF and PETSC_VIEWER_STDOUT_WORLD causes the object to be printed to the screen.
• **PETSC_VIEWER_DRAW_SELF** **PETSC_VIEWER_DRAW_WORLD** causes the object to be drawn in a default X window.

• Passing in a viewer obtained by **PetscViewerDrawOpen()** causes the object to be displayed graphically. See **Graphics** for more on PETSc’s graphics support.

• To save an object to a file in ASCII format, the user creates the viewer object with the command **PetscViewerASCIIOpen(MPI_Comm comm, char* file, PetscViewer *viewer)**. This object is analogous to **PETSC_VIEWER_STDOUT_SELF** (for a communicator of **MPI_COMM_SELF**) and **PETSC_VIEWER_STDOUT_WORLD** (for a parallel communicator).

• To save an object to a file in binary format, the user creates the viewer object with the command **PetscViewerBinaryOpen(MPI_Comm comm, char* file, PetscViewerBinaryType type, PetscViewer *viewer)**. Details of binary I/O are discussed below.

• To save an object to a file in binary format, the user creates the viewer object with the command **PetscViewerSocketOpen(MPI_Comm comm, char *machine, int port, PetscViewer *viewer)**. See **Sending Data to an Interactive MATLAB Session**.

The user can control the format of ASCII printed objects with viewers created by **PetscViewerASCIIOpen()** by calling

```c
PetscViewerPushFormat(PetscViewer viewer, PetscViewerFormat format);
```

Formats include **PETSC_VIEWER_DEFAULT**, **PETSC_VIEWER_ASCII_MATLAB**, and **PETSC_VIEWER_ASCII_IMPL**. The implementation-specific format, **PETSC_VIEWER_ASCII_IMPL**, displays the object in the most natural way for a particular implementation.

The routines

```c
PetscViewerPushFormat(PetscViewer viewer, PetscViewerFormat format);
PetcViewerPopFormat(PetscViewer viewer);
```

allow one to temporarily change the format of a viewer.

As discussed above, one can output PETSc objects in binary format by first opening a binary viewer with **PetscViewerBinaryOpen()** and then using **MatView()**, **VecView()**, etc. The corresponding routines for input of a binary object have the form **XXXLoad()**. In particular, matrix and vector binary input is handled by the following routines:

```c
MatLoad(Mat newmat, PetscViewer viewer);
VecLoad(Vec newvec, PetscViewer viewer);
```

These routines generate parallel matrices and vectors if the viewer’s communicator has more than one process. The particular matrix and vector formats are determined from the options database; see the manual pages for details.

One can provide additional information about matrix data for matrices stored on disk by providing an optional file **matrixfilename.info**, where **matrixfilename** is the name of the file containing the matrix. The format of the optional file is the same as the **.petscrc** file and can (currently) contain the following:

```-matload_block_size <bs>
```

The block size indicates the size of blocks to use if the matrix is read into a block oriented data structure (for example, **MATMPIBAIJ**). The diagonal information **s1,s2,s3,...** indicates which (block) diagonals in the matrix have nonzero values. The info file is automatically created when **VecView()** or **MatView()** is used with a binary viewer; hence if you save a matrix with a given block size with **MatView()**, then a **MatLoad()** on that file will automatically use the saved block size.

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Viewing From Options

Command-line options provide a particularly convenient way to view PETSc objects. All options of the form -xxx_view accept colon(:)-separated compound arguments which specify a viewer type, format, and/or destination (e.g. file name or socket) if appropriate. For example, to quickly export a binary file containing a matrix, one may use -mat_view binary:matrix.out, or to output to a MATLAB-compatible ASCII file, one may use -mat_view ascii:matrix.m:ascii_matlab. See the PetscOptionsGetViewer() man page for full details, as well as the XXXViewFromOptions() man pages (for instance, PetscDrawSetFromOptions()) for many other convenient command-line options.

Using Viewers to Check Load Imbalance

The PetscViewer format PETSC_VIEWER_LOAD_BALANCE will cause certain objects to display simple measures of their imbalance. For example

```
-n 4 ./ex32 -ksp_view_mat ::load_balance
```

will display

```
Nonzeros: Min 162 avg 168 max 174
```

indicating that one process has 162 nonzero entries in the matrix, the average number of nonzeros per process is 168 and the maximum number of nonzeros is 174. Similar for vectors one can see the load balancing with, for example,

```
-n 4 ./ex32 -ksp_view_rhs ::load_balance
```

The measurements of load balancing can also be done within the program with calls to the appropriate object viewer with the viewer format PETSC_VIEWER_LOAD_BALANCE.

4.6.4 Using SAWs with PETSc

The Scientific Application Web server, SAWs\(^1\), allows one to monitor running PETSc applications from a browser. To use SAWs you must configure PETSc with the option --download-saws. Options to use SAWs include

- **-saws_options** - allows setting values in the PETSc options database via the browser (works only on one process).
- **-stack_view saws** - allows monitoring the current stack frame that PETSc is in; refresh to see the new location.
- **-snes_monitor_saws, -ksp_monitor_saws** - monitor the solvers’ iterations from the web browser.

For each of these you need to point your browser to http://hostname:8080, for example http://localhost:8080. Options that control behavior of SAWs include

- **-saws_log filename** - log all SAWs actions in a file.
- **-saws_https certfile** - use HTTPS instead of HTTP with a certificate.
- **-saws_port_auto_select** - have SAWs pick a port number instead of using 8080.
- **-saws_port port** - use port instead of 8080.

\(^1\) Saws wiki on Bitbucket
4.6.5 Debugging

PETSc programs may be debugged using one of the two options below.

- **-start_in_debugger [noxterm, dbx, xgdb, xdb, xlddb, lldb] [-display name]** - start all processes in debugger
- **-on_error_attach_debugger [noxterm, dbx, xgdb, xdb, xlddb, lldb] [-display name]** - start debugger only on encountering an error

Note that, in general, debugging MPI programs cannot be done in the usual manner of starting the programming in the debugger (because then it cannot set up the MPI communication and remote processes).

By default on Linux systems the GNU debugger *gdb* is used, on macOS systems *lldb* is used.

By default, the debugger will be started in a new xterm (Apple Terminal on macOS), to enable running separate debuggers on each process, unless the option *noxterm* is used. In order to handle the MPI startup phase, the debugger command *cont* should be used to continue execution of the program within the debugger. Rerunning the program through the debugger requires terminating the first job and restarting the processor(s); the usual *run* option in the debugger will not correctly handle the MPI startup and should not be used. Not all debuggers work on all machines, the user may have to experiment to find one that works correctly.

You can select a subset of the processes to be debugged (the rest just run without the debugger) with the option

```
-dbugger_ranks rank1,rank2,...
```

where you simply list the ranks you want the debugger to run with.

4.6.6 Error Handling

Errors are handled through the routine *PetscError()* . This routine checks a stack of error handlers and calls the one on the top. If the stack is empty, it selects *PetscTraceBackErrorHandler()* , which tries to print a traceback. A new error handler can be put on the stack with

```
PetscPushErrorHandler(PetscErrorCode (*HandlerFunction)(int line, char *dir, char *file, char *message, int number, void *HandlerContext), void *HandlerContext)
```

The arguments to *HandlerFunction()* are the line number where the error occurred, the file in which the error was detected, the corresponding directory, the error message, the error integer, and the *HandlerContext*. The routine

```
PetscPopErrorHandler()
```

removes the last error handler and discards it.

PETSc provides two additional error handlers besides *PetscTraceBackErrorHandler()* :
The function `PetscAbortErrorHandler()` calls `abort` on encountering an error, while `PetscAttachDebuggerErrorHandler()` attaches a debugger to the running process if an error is detected. At runtime, these error handlers can be set with the options `-on_error_abort` or `-on_error_attach_debugger [noxterm, dbx, xxgdb, xldb] [-display DISPLAY]`.

All PETSc calls can be traced (useful for determining where a program is hanging without running in the debugger) with the option `-log_trace [filename]` where `filename` is optional. By default the traces are printed to the screen. This can also be set with the command `PetscLogTraceBegin(FILE*)`.

It is also possible to trap signals by using the command `PetscPushSignalHandler(PetscErrorCode (*Handler)(int, void *), void *ctx);`

The default handler `PetscSignalHandlerDefault()` calls `PetscError()` and then terminates. In general, a signal in PETSc indicates a catastrophic failure. Any error handler that the user provides should try to clean up only before exiting. By default all PETSc programs turn on the default PETSc signal handler in `PetscInitialize()`, this can be prevented with the option `-no_signal_handler` that can be provided on the command line, in the ~/.petsrc file, or with the call `PetscCall(PetscOptionsSetValue(PETSC_NULL_OPTIONS, "-no_signal_handler", "true"));`.

Once the first PETSc signal handler has been pushed it is impossible to go back to to a signal handler that was set directly by the user with the UNIX signal handler API or by the loader.

Some Fortran compilers/loaders cause, by default, a traceback of the Fortran call stack when a segmentation violation occurs to be printed. This is handled by them setting a special signal handler when the program is started up. This feature is useful for debugging without needing to start up a debugger. If `PetscPushSignalHandler()` has been called this traceback will not occur, hence if the Fortran traceback is desired one should put `PetscCallA(PetscOptionsSetValue(PETSC_NULL_OPTIONS, "-no_signal_handler","true",ierr))` before the call to `PetscInitialize()`. This prevents PETSc from defaulting to using a signal handler.

There is a separate signal handler for floating-point exceptions. The option `-fp_trap` turns on the floating-point trap at runtime, and the routine `PetscFPTrapPush(PetscFPTrap flag);` can be used within a program. A flag of `PETSC_FP_TRAP_ON` indicates that floating-point exceptions should be trapped, while a value of `PETSC_FP_TRAP_OFF` (the default) indicates that they should be ignored.

`PetscFPTrapPop(void);` should be used to revert to the previous handling of floating point exceptions before the call to `PetscFPTrapPush()`.

A small set of macros is used to make the error handling lightweight. These macros are used throughout the PETSc libraries and can be employed by the application programmer as well. When an error is first detected, one should set it by calling...
The user should check the return codes for all PETSc routines (and possibly user-defined routines as well) with

```c
PetscCall(PetscRoutine(...));
```

Likewise, all memory allocations should be checked with

```c
PetscCall(PetscMalloc1(n, &ptr));
```

If this procedure is followed throughout all of the user’s libraries and codes, any error will by default generate a clean traceback of the location of the error.

Note that the macro `PETSC_FUNCTION_NAME` is used to keep track of routine names during error tracebacks. Users need not worry about this macro in their application codes; however, users can take advantage of this feature if desired by setting this macro before each user-defined routine that may call `SETERRQ()`, `PetscCall()`. A simple example of usage is given below.

```c
PetscErrorCode MyRoutine1()
{
    /* Declarations Here */
    PetscFunctionBeginUser;
    /* code here */
    PetscFunctionReturn(PETSC_SUCCESS);
}
```

### 4.6.7 Numbers

PETSc supports the use of complex numbers in application programs written in C, C++, and Fortran. To do so, we employ either the C99 `complex` type or the C++ versions of the PETSc libraries in which the basic “scalar” datatype, given in PETSc codes by `PetscScalar`, is defined as `complex` (or `complex<double>` for machines using templated complex class libraries). To work with complex numbers, the user should run `configure` with the additional option `--with-scalar-type=complex`. The installation instructions provide detailed instructions for installing PETSc. You can use `--with-clanguage=c` (the default) to use the C99 complex numbers or `--with-clanguage=c++` to use the C++ complex type\(^2\).

Recall that each configuration of the PETSc libraries is stored in a different directory, given by `PETSC_DIR/PETSC_ARCH` according to the architecture. Thus, the libraries for complex numbers are maintained separately from those for real numbers. When using any of the complex numbers versions of PETSc, all vector and matrix elements are treated as complex, even if their imaginary components are zero. Of course, one can elect to use only the real parts of the complex numbers when using the complex versions of the PETSc libraries; however, when working only with real numbers in a code, one should use a version of PETSc for real numbers for best efficiency.

The program KSP Tutorial ex11 solves a linear system with a complex coefficient matrix. Its Fortran counterpart is KSP Tutorial ex11f.

\(^2\) Note that this option is not required to use PETSc with C++
4.6.8 Parallel Communication

When used in a message-passing environment, all communication within PETSc is done through MPI, the message-passing interface standard [For94]. Any file that includes `petscsys.h` (or any other PETSc include file) can freely use any MPI routine.

4.6.9 Graphics

The PETSc graphics library is not intended to compete with high-quality graphics packages. Instead, it is intended to be easy to use interactively with PETSc programs. We urge users to generate their publication-quality graphics using a professional graphics package. If a user wants to hook certain packages into PETSc, he or she should send a message to `petsc-maint@mcs.anl.gov`; we will see whether it is reasonable to try to provide direct interfaces.

Windows as PetscViewers

For drawing predefined PETSc objects such as matrices and vectors, one may first create a viewer using the command

```c
PetscViewerDrawOpen(MPI_Comm comm, char *display, char *title, int x, int y, int w, int h, PetscViewer *viewer);
```

This viewer may be passed to any of the `XXXView()` routines. Alternately, one may use command-line options to quickly specify viewer formats, including `PetscDraw`-based ones; see Viewing From Options.

To draw directly into the viewer, one must obtain the `PetscDraw` object with the command

```c
PetscViewerDrawGetDraw(PetscViewer viewer, PetscDraw *draw);
```

Then one can call any of the `PetscDrawXXX()` commands on the `draw` object. If one obtains the `draw` object in this manner, one does not call the `PetscDrawOpenX()` command discussed below.

Predefined viewers, `PETSC_VIEWER_DRAW_WORLD` and `PETSC_VIEWER_DRAW_SELF`, may be used at any time. Their initial use will cause the appropriate window to be created.

Implementations using OpenGL, TikZ, and other formats may be selected with `PetscDrawSetType()`. PETSc can also produce movies; see `PetscDrawSetSaveMovie()`, and note that command-line options can also be convenient; see the `PetscDrawSetFromOptions()` man page.

By default, PETSc drawing tools employ a private colormap, which remedies the problem of poor color choices for contour plots due to an external program’s mangling of the colormap. Unfortunately, this may cause flashing of colors as the mouse is moved between the PETSc windows and other windows. Alternatively, a shared colormap can be used via the option `-draw_x_shared_colormap`.

Simple PetscDrawing

With the default format, one can open a window that is not associated with a viewer directly under the X11 Window System with the command

```c
PetscDrawCreate(MPI_Comm comm, char *display, char *title, int x, int y, int w, int h, PetscDraw *win);
PetscDrawSetFromOptions(win);
```
All drawing routines are performed relative to the window’s coordinate system and viewport. By default, the
drawing coordinates are from \((0,0)\) to \((1,1)\), where \((0,0)\) indicates the lower left corner of the window.
The application program can change the window coordinates with the command

\[
\text{PetscDrawSetCoordinates(PetscDraw win, PetscReal xl, PetscReal yl, PetscReal xr, yr);}\]

By default, graphics will be drawn in the entire window. To restrict the drawing to a portion of the window,
one may use the command

\[
\text{PetscDrawSetViewPort(PetscDraw win, PetscReal xl, PetscReal yl, PetscReal xr, yr);}\]

These arguments, which indicate the fraction of the window in which the drawing should be done, must
satisfy \(0 \leq xl \leq xr \leq 1\) and \(0 \leq yl \leq yr \leq 1\).

To draw a line, one uses the command

\[
\text{PetscDrawLine(PetscDraw win, PetscReal xl, PetscReal yl, PetscReal xr, yr, int cl);}\]

The argument \(cl\) indicates the color (which is an integer between 0 and 255) of the line. A list of predefined
colors may be found in include/petscdraw.h and includes PETSC_DRAW_BLACK, PETSC_DRAW_RED,
PETSC_DRAW_BLUE etc.

To ensure that all graphics actually have been displayed, one should use the command

\[
\text{PetscDrawFlush(PetscDraw win);}\]

When displaying by using double buffering, which is set with the command

\[
\text{PetscDrawSetDoubleBuffer(PetscDraw win);}\]

all processes must call

\[
\text{PetscDrawFlush(PetscDraw win);}\]

in order to swap the buffers. From the options database one may use \(-\text{draw\_pause}\ n\), which causes the
PETSc application to pause \(n\) seconds at each PetscDrawPause(). A time of \(-1\) indicates that the
application should pause until receiving mouse input from the user.

Text can be drawn with commands

\[
\text{PetscDrawString(PetscDraw win, PetscReal x, PetscReal y, int color, char *text);}\]
\[
\text{PetscDrawStringVertical(PetscDraw win, PetscReal x, PetscReal y, int color, const char *text);}\]
\[
\text{PetscDrawStringCentered(PetscDraw win, PetscReal x, PetscReal y, int color, const char *text);}\]
\[
\text{PetscDrawStringBoxed(PetscDraw draw, PetscReal sxl, PetscReal syl, int sc, int bc, const char text[], PetscReal *w, PetscReal *h);}\]

The user can set the text font size or determine it with the commands

\[
\text{PetscDrawStringSetSize(PetscDraw win, PetscReal width, PetscReal height);}\]
\[
\text{PetscDrawStringGetSize(PetscDraw win, PetscReal *width, PetscReal *height);}\]
Line Graphs

PETSc includes a set of routines for manipulating simple two-dimensional graphs. These routines, which begin with `PetscDrawAxisDraw()`, are usually not used directly by the application programmer. Instead, the programmer employs the line graph routines to draw simple line graphs. As shown in the listing below, line graphs are created with the command

```c
PetscDrawLGCreate(PetscDraw win, PetscInt ncurves, PetscDrawLG *ctx);
```

The argument `ncurves` indicates how many curves are to be drawn. Points can be added to each of the curves with the command

```c
PetscDrawLGAddPoint(PetscDrawLG ctx, PetscReal *x, PetscReal *y);
```

The arguments `x` and `y` are arrays containing the next point value for each curve. Several points for each curve may be added with

```c
PetscDrawLGAddPoints(PetscDrawLG ctx, PetscInt n, PetscReal **x, PetscReal **y);
```

The line graph is drawn (or redrawn) with the command

```c
PetscDrawLGDraw(PetscDrawLG ctx);
```

A line graph that is no longer needed can be destroyed with the command

```c
PetscDrawLGDestroy(PetscDrawLG *ctx);
```

To plot new curves, one can reset a linegraph with the command

```c
PetscDrawLGReset(PetscDrawLG ctx);
```

The line graph automatically determines the range of values to display on the two axes. The user can change these defaults with the command

```c
PetscDrawLGSetLimits(PetscDrawLG ctx, PetscReal xmin, PetscReal xmax, PetscReal ymin, PetscReal ymax);
```

It is also possible to change the display of the axes and to label them. This procedure is done by first obtaining the axes context with the command

```c
PetscDrawLGGetAxis(PetscDrawLG ctx, PetscDrawAxis *axis);
```

One can set the axes’ colors and labels, respectively, by using the commands

```c
PetscDrawAxisSetColors(PetscDrawAxis axis, int axis_lines, int ticks, int text);
PetscDrawAxisSetLabels(PetscDrawAxis axis, char *top, char *x, char *y);
```

It is possible to turn off all graphics with the option `-nox`. This will prevent any windows from being opened or any drawing actions to be done. This is useful for running large jobs when the graphics overhead is too large, or for timing.

The full example, Draw Test ex3, follows.

Listing: `src/classes/draw/tests/ex3.c`
static char help[] = "Plots a simple line graph.\n";

#if defined(PETSC_APPLE_FRAMEWORK)
    #import <PETSc/petscsys.h>
    #import <PETSc/petscdraw.h>
#else
    #include <petscsys.h>
    #include <petscdraw.h>
#endif

int main(int argc, char **argv)
{
    PetscDraw draw;
    PetscDrawLG lg;
    PetscDrawAxis axis;
    PetscInt n = 15, i, x = 0, y = 0, width = 400, height = 300, nports = 1;
    PetscBool useports, flg;
    const char *xlabel, *ylabel, *toplabel, *legend;
    PetscReal xd, yd;
    PetscDrawViewPorts *ports = NULL;

toplabel = "Top Label";
xlabel = "X-axis Label";
ylabel = "Y-axis Label";
legend = "Legend";

PetscFunctionBeginUser;
PetscCall(PetscInitialize(&argc, &argv, NULL, help));
PetscCall(PetscOptionsGetInt(NULL, NULL, "-x", &x, NULL));
PetscCall(PetscOptionsGetInt(NULL, NULL, "-y", &y, NULL));
PetscCall(PetscOptionsGetInt(NULL, NULL, "-width", &width, NULL));
PetscCall(PetscOptionsGetInt(NULL, NULL, "-height", &height, NULL));
PetscCall(PetscOptionsGetInt(NULL, NULL, "-n", &n, NULL));
PetscCall(PetscOptionsGetInt(NULL, NULL, "-nports", &nports, &useports));
PetscCall(PetscOptionsHasName(NULL, NULL, "-nolegend", &flg));
    if (flg) legend = NULL;
PetscCall(PetscOptionsHasName(NULL, NULL, "-notoplabel", &flg));
    if (flg) toplabel = NULL;
PetscCall(PetscOptionsHasName(NULL, NULL, "-noxlabel", &flg));
    if (flg) xlabel = NULL;
PetscCall(PetscOptionsHasName(NULL, NULL, "-noylabel", &flg));
    if (flg) ylabel = NULL;
PetscCall(PetscOptionsHasName(NULL, NULL, "-nolabels", &flg));
    if (flg) {
        toplabel = NULL;
        xlabel = NULL;
        ylabel = NULL;
    }

PetscCall(PetscDrawCreate(PETSC_COMM_WORLD, 0, "Title", x, y, width, height, &draw));
PetscCall(PetscDrawSetFromOptions(draw));
if (useports) {
    PetscCall(PetscDrawViewPortsCreate(draw, nports, &ports));
PetscCall(PetscDrawViewPortsSet(ports, 0));
}

(continues on next page)
Graphical Convergence Monitor

For both the linear and nonlinear solvers default routines allow one to graphically monitor convergence of the iterative method. These are accessed via the command line with `-ksp_monitor draw::draw_lg` and `-snes_monitor draw::draw_lg`. See also `Convergence Monitoring` and `Convergence Monitoring`.

Disabling Graphics at Compile Time

To disable all X-window-based graphics, run `configure` with the additional option `-with-x=0`

4.7 Developer Environments

4.7.1 Emacs Users

Many PETSc developers use Emacs, which can be used as a “simple” text editor or a comprehensive development environment. For a more integrated development environment, we recommend using `lsp-mode` (or `eglot`) with `clangd`. The most convenient way to teach `clangd` what compilation flags to use is to install `Bear` (“build ear”) and run:

```
bear make -B
```

which will do a complete rebuild (-B) of PETSc and capture the compilation commands in a file named `compile_commands.json`, which will be automatically picked up by `clangd`. You can use the same procedure when building examples or your own project. It can also be used with any other editor that
supports clangd, including VS Code and Vim. When lsp-mode is accompanied by flycheck, Emacs will provide real-time feedback and syntax checking, along with refactoring tools provided by clangd.

The easiest way to install packages in recent Emacs is to use the “Options” menu to select “Manage Emacs Packages”.

### Tags

It is sometimes useful to cross-reference tags across projects. Regardless of whether you use lsp-mode, it can be useful to use GNU Global (install `gtags`) to provide reverse lookups (e.g. find all call sites for a given function) across all projects you might work on/browse. Tags for PETSc can be generated by running `make allgtags` from `$PETSC_DIR`, or one can generate tags for all projects by running a command such as

```bash
find $PETSC_DIR/{include,src,tutorials,$PETSC_ARCH/include} any/other/paths \
  -regex '.*\.(cc|hh|cpp|cxx|C|hpp|c|h|h\)cu\)$' \n  | grep -v ftn-auto | gtags -f -
```

from your home directory or wherever you keep source code. If you are making large changes, it is useful to either set this up to run as a cron job or to make a convenient alias so that refreshing is easy. Then add the following to `~/.emacs` to enable gtags and specify key bindings.

```lisp
(setq require 'gtags)
(global-set-key (kbd "C-c f") 'gtags-find-file)
(global-set-key (kbd "C-c .") 'gtags-find-tag)
(global-set-key (kbd "C-c r") 'gtags-find-rtag)
(global-set-key (kbd "C-c ,") 'gtags-pop-stack)
(add-hook 'c-mode-common-hook
  '(lambda () (gtags-mode t)))) ; Or add to existing hook
```

A more basic alternative to the GNU Global (`gtags`) approach that does not require adding packages is to use the builtin `etags` feature. First, run `make alletags` from the PETSc home directory to generate the file `$PETSC_DIR/TAGS`, and then from within Emacs, run

```
M-x visit-tags-table
```

where M denotes the Emacs Meta key, and enter the name of the TAGS file. Then the command M-. will cause Emacs to find the file and line number where a desired PETSc function is defined. Any string in any of the PETSc files can be found with the command M-x `tags-search`. To find repeated occurrences, one can simply use M-, to find the next occurrence.

### 4.7.2 VS Code Users

VS Code (unlike Visual Studio Users, described below) is an open-source editor with a rich extension ecosystem. It has excellent integration with clangd and will automatically pick up `compile_commands.json` as produced by a command such as `bear make -B` (see Developer Environments). If you have no prior attachment to a specific code editor, we recommend trying VS Code.
4.7.3 Vi and Vim Users

This section lists helpful Vim commands for PETSc. Ones that configure Vim can be placed in a .vimrc file in the top of the PETSc directory and will be loaded automatically.

Vim has configurable keymaps: all of the “command mode” commands given that start with a colon (such as :help) can be assigned to short sequences in “normal mode,” which is how most Vim users use their most frequently used commands.

See the Developer Environments discussion above for configuration of clangd, which provides integrated development environment.

Tags

The tags feature can be used to search PETSc files quickly and efficiently. To use this feature, one should first check if the file, $PETSC_DIR/CTAGS exists. If this file is not present, it should be generated by running make alltags from the PETSc home directory. Once the file exists, from Vi/Vim the user should issue the command

```
:set tags=CTAGS
```

from the $PETSC_DIR directory and enter the name of the CTAGS file. The command :tag functionname will cause Vi/Vim to open the file and line number where a desired PETSc function is defined in the current window. <Ctrl-o> will return the screen to your previous location.

The command :stag functionname will split the current window and then open the file and line number for that function in one half. Some prefer this because it is easier to compare the file you are editing to the function definition this way.

Cscope and gtags

Vim can also use the cscope utility to navigate source code. One useful thing it can do that the basic tags feature can’t is search for references to a symbol, rather than its definition, which can be useful for refactoring. The command

```
:cs find s functionname
```

opens a list of all of the places the function is called in PETSc, and opens the file and line that you choose. The variant :scs find s functionname does the same but splits the window like stag.

The PETSc makefile does not have a command for building a cscope database, but GNU Global is cross-compatible with cscope: call make allgtags to make the gtags database, and run the commands

```
:set csprg=gtags-cscope
:cs add GTAGS
```
Quickfix

Rather than exiting editing a file to build the library and check for errors or warnings, calling `:make` runs the make command without leaving Vim and collects the errors and warnings in a “quickfix” window. Move the cursor to one of the errors or warnings in the quickfix window and press `<Enter>` and the main window will jump to the file and line with the error. The following commands filter lines of out PETSc’s make output that can clutter the quickfix window:

```vim
:set efm^=%-GStarting\ make\ run\ on\ %.%#
:set efm^=%-GMachine\ characteristics:\ %.%#
:set efm^=%-G#define\ PETSC%.%#
```

Autocompletion and snippets

Autocompletion of long function names can be helpful when working with PETSc. If you have a tags file, you can press `<Ctrl-N>` when you have partially typed a word to bring up a list of potential completions that you can choose from with `<Tab>`.

More powerful autocompletion, such as completing the fieldname of a struct, is available from external plugins that can be added to Vim, such as SuperTab, VimCompletesMe, or YouCompleteMe.

Along the same lines, plugins can be added that fill in the boilerplate associated with PETSc programming with code snippets. One such tool is UltiSnips.

LSP for Vim

Several plugins provide the equivalent of emacs’ lsp-mode: YouCompleteMe, mentioned above, is one; another popular one is ale. These can check for syntax errors, check for compilation errors in the background, and provide sophisticated tools for refactoring. Like lsp-mode, they also rely on a compilation database, so `bear -- make -B` should be used as well to generate the file `compile_commands.json`.

See online tutorials for additional Vi/Vim options.

4.7.4 Eclipse Users

If you are interested in developing code that uses PETSc from Eclipse or developing PETSc in Eclipse and have knowledge of how to do indexing and build libraries in Eclipse, please contact us at petsc-dev@mcs.anl.gov.

One way to index and build PETSc in Eclipse is as follows.

1. Open “File→Import→Git→Projects from Git”. In the next two panels, you can either add your existing local repository or download PETSc from Bitbucket by providing the URL. Most Eclipse distributions come with Git support. If not, install the EGit plugin. When importing the project, select the wizard “Import as general project”.

2. Right-click on the project (or the “File” menu on top) and select “New → Convert to a C/C++ Project (Adds C/C++ Nature)”. In the setting window, choose “C Project” and specify the project type as “Shared Library”.

3. Right-click on the C project and open the “Properties” panel. Under “C/C++ Build → Builder Settings”, set the Build directory to `$PETSC_DIR` and make sure “Generate Makefiles automatically” is unselected. Under the section “C/C++ General→Paths and Symbols”, add the PETSc paths to “Includes”.

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1. Configure PETSc normally outside Eclipse to generate a makefile and then build the project in Eclipse.

   The source code will be parsed by Eclipse.

   If you launch Eclipse from the Dock on Mac OS X, .bashrc will not be loaded (a known OS X behavior, for security reasons). This will be a problem if you set the environment variables $PETSC_DIR and $PETSC_ARCH in .bashrc. A solution which involves replacing the executable can be found at `/questions/829749/launch-mac-eclipse-with-environment-variables-set` . Alternatively, you can add $PETSC_DIR and $PETSC_ARCH manually under “Properties ! C/C++ Build ! Environment”.

   To allow an Eclipse code to compile with the PETSc include files and link with the PETSc libraries, a PETSc user has suggested the following.

   1. Right-click on your C project and select “Properties → C/C++ Build → Settings”
   2. A new window on the righthand side appears with various settings options. Select “Includes” and add the required PETSc paths,

```
$PETSC_DIR/include
$PETSC_DIR/$PETSC_ARCH/include
```

   1. Select “Libraries” under the header Linker and set the library search path:

```
$PETSC_DIR/$PETSC_ARCH/lib
```

   and the libraries, for example

```
m, petsc, stdc++, mpichxx, mpich, lapack, blas, gfortran, dl, rt, gcc_s, pthread, X11
```

   Another PETSc user has provided the following steps to build an Eclipse index for PETSc that can be used with their own code, without compiling PETSc source into their project.

   1. In the user project source directory, create a symlink to the PETSC src/ directory.
   2. Refresh the project explorer in Eclipse, so the new symlink is followed.
   3. Right-click on the project in the project explorer, and choose “Index → Rebuild”. The index should now be built.
   4. Right-click on the PETSc symlink in the project explorer, and choose “Exclude from build...” to make sure Eclipse does not try to compile PETSc with the project.

   For further examples of using Eclipse with a PETSc-based application, see the documentation for LaMEM³.

³ See the doc/ directory at https://bitbucket.org/bkaus/lamem
4.7.5 Qt Creator Users

This information was provided by Mohammad Mirzadeh. The Qt Creator IDE is part of the Qt SDK, developed for cross-platform GUI programming using C++. It is available under GPL v3, LGPL v2 and a commercial license and may be obtained, either as part of the Qt SDK or as stand-alone software. It supports automatic makefile generation using cross-platform qmake and CMake build systems as well as allowing one to import projects based on existing, possibly hand-written, makefiles. Qt Creator has a visual debugger using GDB and LLDB (on Linux and OS X) or Microsoft’s CDB (on Microsoft Windows) as backends. It also has an interface to Valgrind’s “memcheck” and “callgrind” tools to detect memory leaks and profile code. It has built-in support for a variety of version control systems including git, mercurial, and subversion. Finally, Qt Creator comes fully equipped with auto-completion, function look-up, and code refactoring tools. This enables one to easily browse source files, find relevant functions, and refactor them across an entire project.

Creating a Project

When using Qt Creator with qmake, one needs a .pro file. This configuration file tells Qt Creator about all build/compile options and locations of source files. One may start with a blank .pro file and fill in configuration options as needed. For example:

```make
# The name of the application executable
TARGET = ex1

# There are two ways to add PETSc functionality
# 1-Manual: Set all include path and libs required by PETSc
PETSC_INCLUDE = path/to/petsc_includes # e.g. obtained via running `make
    --getincludedirs`
PETSC_LIBS = path/to/petsc_libs # e.g. obtained via running `make getlinklibs`

# 2-Automatic: Use the PKGCONFIG functionalty
# NOTE: petsc.pc must be in the pkgconfig path. You might need to adjust PKG_CONFIG_   
   --PATH
CONFIG += link pkgconfig
PKGCONFIG += PETSc

# Set appropriate compiler and its flags
QMAKE_CC = path/to/mpicc
QMAKE_CXX = path/to/mpicxx # if this is a cpp project
QMAKE_LINK = path/to/mpicxx # if this is a cpp project
QMAKE_CFLAGS += -O3 # add extra flags here
QMAKE_CXXFLAGS += -O3
QMAKE_LFLAGS += -O3

# Add all files that must be compiled
SOURCES += ex1.c source1.c source2.cpp
HEADERS += source1.h source2.h

# OTHER_FILES are ignored during compilation but will be shown in file panel in QtCreator
OTHER_FILES += 
```

(continues on next page)
path/to/resource_file \\ path/to/another_file

In this example, keywords include:

- **TARGET**: The name of the application executable.
- **INCLUDEPATH**: Used at compile time to point to required include files. Essentially, it is used as an 
  `-I \$\$INCLUDEPATH` flag for the compiler. This should include all application-specific header files 
  and those related to PETSc (which may be found via `make getincludedirs`).
- **LIBS**: Defines all required external libraries to link with the application. To get PETSc’s linking 
  libraries, use `make getlinklibs`.
- **CONFIG**: Configuration options to be used by `qmake`. Here, the option `link_pkgconfig` instructs 
  `qmake` to internally use `pkgconfig` to resolve `INCLUDEPATH` and `LIBS` variables.
- **PKGCONFIG**: Name of the configuration file (the `.pc` file – here `petsc.pc`) to be passed to `pkg-
  config`. Note that for this functionality to work, `petsc.pc` must be in path which might require 
  adjusting the `PKG_CONFIG_PATH` environment variable. For more information see the Qt Creator 
  documentation.
- **QMAKE_CC** and **QMAKE_CXX**: Define which C/C++ compilers use.
- **QMAKE_LINK**: Defines the proper linker to be used. Relevant if compiling C++ projects.
- **QMAKE_CFLAGS**, **QMAKE_CXXFLAGS** and **QMAKE_LFLAGS**: Set the corresponding compile and linking 
  flags.
- **SOURCES**: Source files to be compiled.
- **HEADERS**: Header files required by the application.
- **OTHER_FILES**: Other files to include (source, header, or any other extension). Note that none of the 
  source files placed here are compiled.

More options can be included in a `.pro` file; see [https://doc.qt.io/qt-5/qmake-project-files.html](https://doc.qt.io/qt-5/qmake-project-files.html). Once the 
`.pro` file is generated, the user can simply open it via Qt Creator. Upon opening, one has the option to 
create two different build options, debug and release, and switch between the two. For more information 
on using the Qt Creator interface and other more advanced aspects of the IDE, refer to [https://www.qt.io/ 

### 4.7.6 Visual Studio Users

To use PETSc from Microsoft Visual Studio, one would have to compile a PETSc example with its corre-
spanding makefile and then transcribe all compiler and linker options used in this build into a Visual Studio 
project file, in the appropriate format in Visual Studio project settings.
4.7.7 Xcode IDE Users

See doc_macos_install for the standard Unix command line tools approach to development on macOS. The information below is only if you plan to write code within the Xcode IDE.

Apple Xcode IDE for macOS Applications

Follow the instructions in $PETSC_DIR/systems/Apple/OSX/bin/makeall to build the PETSc framework and documentation suitable for use in Xcode.

You can then use the PETSc framework in $PETSC_DIR/arch-osx/PETSc.framework in the usual manner for Apple frameworks. See the examples in $PETSC_DIR/systems/Apple/OSX/examples. When working in Xcode, things like function name completion should work for all PETSc functions as well as MPI functions. You must also link against the Apple Accelerate.framework.

Apple Xcode IDE for iPhone/iPad iOS Applications

Follow the instructions in $PETSC_DIR/systems/Apple/iOS/bin/iosbuilder.py to build the PETSc library for use on the iPhone/iPad.

You can then use the PETSc static library in $PETSC_DIR/arch-osx/libPETSc.a in the usual manner for Apple libraries inside your iOS XCode projects; see the examples in $PETSC_DIR/systems/Apple/iOS/examples. You must also link against the Apple Accelerate.framework.

A thorough discussion of the procedure is given in Comparison of Migration Techniques for High-Performance Code to Android and iOS.

For Android, you must have your standalone bin folder in the path, so that the compilers are visible.

The installation process has not been tested for iOS or Android since 2017.

4.8 Advanced Features of Matrices and Solvers

This chapter introduces additional features of the PETSc matrices and solvers.

4.8.1 Extracting Submatrices

One can extract a (parallel) submatrix from a given (parallel) using

\[
\text{MatCreateSubMatrix(Mat } A, \text{IS } \text{rows}, \text{IS } \text{cols}, \text{MatReuse } \text{call}, \text{Mat } *B);\\
\]

This extracts the rows and cols of the matrix A into B. If call is MAT_INITIAL_MATRIX it will create the matrix B. If call is MAT_REUSE_MATRIX it will reuse the B created with a previous call. This function is used internally by PCFIELDSPLIT.

One can also extract one or more submatrices per MPI process with

\[
\text{MatCreateSubMatrices(Mat } A, \text{PetscInt } n, \text{IS } \text{rows[]}, \text{IS } \text{cols[]}, \text{MatReuse } \text{call}, \text{Mat } *B[]);\\
\]

This extracts n (zero or more) matrices with the rows[k] and cols[k] of the matrix A into an array of sequential matrices B[k] on this process. If call is MAT_INITIAL_MATRIX it will create the array of matrices B. If call is MAT_REUSE_MATRIX it will reuse the B created with a previous call. The IS arguments are sequential. The array of matrices should be destroyed with MatDestroySubMatrices(). This function is used by PCBJACOBI and PCASM.
Each submatrix may be parallel, existing on a \texttt{MPI	extunderscore Comm} associated with each pair of \texttt{IS} \texttt{rows[k]} and \texttt{cols[k]}, using

\begin{verbatim}
MatCreateSubMatricesMPI(Mat A, PetscInt n, IS rows[], IS cols[], MatReuse call, Mat *B[]);
\end{verbatim}

Finally this version has a specialization

\begin{verbatim}
MatGetMultiProcBlock(Mat A, MPI	extunderscore Comm subComm, MatReuse scall, Mat *subMat);
\end{verbatim}

where collections of non-overlapping MPI processes share a single parallel matrix on their sub-communicator. This function is used by \texttt{PCBJACOBI} and \texttt{PCASM}.

The routine

\begin{verbatim}
MatCreateRedundantMatrix(Mat A, PetscInt nsubcomm, MPI	extunderscore Comm subcomm, MatReuse reuse, Mat *matredundant);
\end{verbatim}

where \texttt{nsubcomm} copies of the entire matrix are stored, one on each \texttt{subcomm}. The routine \texttt{PetscSubcommCreate()} and its \texttt{PetscSubcomm} object may, but need not be, used to construct the \texttt{subcomm}.

The routine

\begin{verbatim}
MatMPIAdjToSeq(Mat A, Mat *B);
\end{verbatim}

is a specialization that duplicates an entire \texttt{MATMPIADJ} matrix on each MPI process.

### 4.8.2 Matrix Factorization

Normally, PETSc users will access the matrix solvers through the \texttt{KSP} interface, as discussed in \textit{KSP: Linear System Solvers}, but the underlying factorization and triangular solve routines are also directly accessible to the user.

The ILU, LU, ICC, Cholesky, and QR matrix factorizations are split into two or three stages depending on the user’s needs. The first stage is to calculate an ordering for the matrix. The ordering generally is done to reduce fill in a sparse factorization; it does not make much sense for a dense matrix.

\begin{verbatim}
MatGetOrdering(Mat matrix, MatOrderingType type, IS *rowperm, IS *colperm);
\end{verbatim}

The currently available alternatives for the ordering \texttt{type} are

- \texttt{MATORDERINGNATURAL} - Natural
- \texttt{MATORDERINGND} - Nested Dissection
- \texttt{MATORDERING1WD} - One-way Dissection
- \texttt{MATORDERINGRCM} - Reverse Cuthill-McKee
- \texttt{MATORDERINGQMD} - Quotient Minimum Degree

These orderings can also be set through the options database.

Certain matrix formats may support only a subset of these. All of these orderings are symmetric at the moment; ordering routines that are not symmetric may be added. Currently we support orderings only for sequential matrices.

Users can add their own ordering routines by providing a function with the calling sequence

\begin{verbatim}
int reorder(Mat A, MatOrderingType type, IS *rowperm, IS *colperm);
\end{verbatim}
Here $A$ is the matrix for which we wish to generate a new ordering, type may be ignored and rowperm and colperm are the row and column permutations generated by the ordering routine. The user registers the ordering routine with the command

```c
MatOrderingRegister(MatOrderingType ordname, char *path, char *sname, PetscErrorCode (*reorder)(Mat, MatOrderingType, IS*, IS*));
```

The input argument ordname is a string of the user's choice, either an ordering defined in petscmat.h or the name of a new ordering introduced by the user. See the code in src/mat/impls/order/sorder.c and other files in that directory for examples on how the reordering routines may be written.

Once the reordering routine has been registered, it can be selected for use at runtime with the command line option `-pc_factor_mat_ordering_type ordname`. If reordering from the API, the user should provide the ordname as the second input argument of MatGetOrdering().

PETSc matrices interface to a variety of external factorization/solver packages via the MatSolverType which can be MATSOLVERSUPERLU_DIST, MATSOLVERMUMPS, MATSOLVERPASTIX, MATSOLVERMKL_PARDISO, MATSOLVERMKL_CPARDISO, MATSOLVERUMFPACK, MATSOLVERCHOLMOD, MATSOLVERKLU, MATSOLVECUSPARSE, and MATSOLVERCUDA. The last three of which can run on GPUs, while MATSOLVERSUPERLU_DIST can partially run on GPUs. See doc_linsolve for a table of the factorization based solvers in PETSc.

Most of these packages compute their own orderings and cannot use ones provided so calls to the following routines with those packages can pass NULL as the IS permutations.

The following routines perform incomplete and complete, in-place, symbolic, and numerical factorizations for symmetric and nonsymmetric matrices, respectively:

```c
MatICCFactor(Mat matrix, IS permutation, const MatFactorInfo *info);
MatLUFactor(Mat matrix, IS rowpermutation, IS columnpermutation, const MatFactorInfo *info);
MatCholeskyFactor(Mat matrix, IS permutation, const MatFactorInfo *info);
MatLUFactor(Mat matrix, IS rowpermutation, IS columnpermutation, const MatFactorInfo *info);
MatQRFactor(Mat matrix, IS columnpermutation, const MatFactorInfo *info);
```

The argument info->fill > 1 is the predicted fill expected in the factored matrix, as a ratio of the original fill. For example, info->fill=2.0 would indicate that one expects the factored matrix to have twice as many nonzeros as the original.

For sparse matrices it is very unlikely that the factorization is actually done in-place. More likely, new space is allocated for the factored matrix and the old space deallocated, but to the user it appears in-place because the factored matrix replaces the unfactored matrix.

The two factorization stages can also be performed separately, by using the preferred out-of-place mode, first one obtains that matrix object that will hold the factor using

```c
MatGetFactor(Mat matrix, MatSolverType package, MatFactorType ftype, Mat *factor);
```

and then performs the factorization

```c
MatICCFactorSymbolic(Mat factor, Mat matrix, IS perm, const MatFactorInfo *info);
MatCholeskyFactorSymbolic(Mat factor, Mat matrix, IS perm, const MatFactorInfo *info);
MatILUFactorSymbolic(Mat factor, Mat matrix, IS rowperm, IS colperm, const MatFactorInfo *info);
MatLUFactorSymbolic(Mat factor, Mat matrix, IS rowperm, IS colperm, const MatFactorInfo *info);
MatCholeskyFactorNumeric(Mat factor, Mat matrix, const MatFactorInfo);
MatLUFactorNumeric(Mat factor, Mat matrix, const MatFactorInfo *info);
```

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MatQRFactorSymbolic(Mat factor, Mat matrix, IS perm, const MatFactorInfo *info);
MatQRFactorNumeric(Mat factor, Mat matrix, const MatFactorInfo *info);

In this case, the contents of the matrix result is undefined between the symbolic and numeric factorization stages. It is possible to reuse the symbolic factorization. For the second and succeeding factorizations, one simply calls the numerical factorization with a new input matrix and the same factored result matrix. It is essential that the new input matrix have exactly the same nonzero structure as the original factored matrix. (The numerical factorization merely overwrites the numerical values in the factored matrix and does not disturb the symbolic portion, thus enabling reuse of the symbolic phase.) In general, calling XXXFactorSymbolic with a dense matrix will do nothing except allocate the new matrix; the XXXFactorNumeric routines will do all of the work.

Why provide the plain XXXfactor routines when one could simply call the two-stage routines? The answer is that if one desires in-place factorization of a sparse matrix, the intermediate stage between the symbolic and numeric phases cannot be stored in a result matrix, and it does not make sense to store the intermediate values inside the original matrix that is being transformed. We originally made the combined factor routines do either in-place or out-of-place factorization, but then decided that this approach was not needed and could easily lead to confusion.

We do not provide our own sparse matrix factorization with pivoting for numerical stability. This is because trying to both reduce fill and do pivoting can become quite complicated. Instead, we provide a poor stepchild substitute. After one has obtained a reordering, with MatGetOrdering(Mat A, MatOrdering type, IS *row, IS *col) one may call

MatReorderForNonzeroDiagonal(Mat A, PetscReal tol, IS row, IS col);

which will try to reorder the columns to ensure that no values along the diagonal are smaller than tol in an absolute value. If small values are detected and corrected for, a nonsymmetric permutation of the rows and columns will result. This is not guaranteed to work, but may help if one was simply unlucky in the original ordering. When using the KSP solver interface the option -pc_factor_nonzeros_along_diagonal <tol> may be used. Here, tol is an optional tolerance to decide if a value is nonzero; by default it is 1.e-10.

The external MatSolverType’s MATSOLVERSUPERLU_DIST and MATSOLVERMUMPS do manage numerical pivoting internal to their API.

The external factorization packages each provide a wide number of options to chose from, details on these may be found by consulting the manual page for the solver package, such as, MATSOLVERSUPERLU_DIST. Most of the options can be easily set via the options database even when the factorization solvers are accessed via KSP.

Once a matrix has been factored, it is natural to solve linear systems. The following four routines enable this process:

MatSolve(Mat A, Vec x, Vec y);
MatSolveTranspose(Mat A, Vec x, Vec y);
MatSolveAdd(Mat A, Vec x, Vec y, Vec w);
MatSolveTransposeAdd(Mat A, Vec x, Vec y, Vec w);

matrix A of these routines must have been obtained from a factorization routine; otherwise, an error will be generated. In general, the user should use the KSP solvers introduced in the next chapter rather than using these factorization and solve routines directly.

Some of the factorizations also support solves with multiple right hand sides stored in a Mat using
and

\texttt{MatMatSolveTranspose(Mat A, Mat B, Mat X);}

Finally, \texttt{MATSOLVERMUMPS}, provides access to Schur complements obtained after partial factorizations as well as the inertia of a matrix via \texttt{MatGetInertia()}.

## 4.8.3 Matrix-Matrix Products

PETSc matrices provide code for computing various matrix-matrix products. This section will introduce the two sets of routines available. For now consult \texttt{MatCreateProduct()} and \texttt{MatMatMult()}.

## 4.8.4 Creating PC’s Directly

Users obtain their preconditioner contexts from the \texttt{KSP} context with the command \texttt{KSPGetPC()}. It is possible to create, manipulate, and destroy \texttt{PC} contexts directly, although this capability should rarely be needed. To create a \texttt{PC} context, one uses the command

\texttt{PCCreate(MPI_Comm comm, PC \*pc);} 

The routine

\texttt{PCSetType(PC pc, PCTYPE method);} 

sets the preconditioner method to be used. The routine

\texttt{PCSetOperators(PC pc, Mat Amat, Mat Pmat);} 

set the matrices that are to be used with the preconditioner. The routine

\texttt{PCGetOperators(PC pc, Mat \*Amat, Mat \*Pmat);} 

returns the values set with \texttt{PCSetOperators()}.

The preconditioners in PETSc can be used in several ways. The two most basic routines simply apply the preconditioner or its transpose and are given, respectively, by

\texttt{PCApply(PC pc, Vec x, Vec y);} 
\texttt{PCApplyTranspose(PC pc, Vec x, Vec y);} 

In particular, for a preconditioner matrix, \(B\), that has been set via \texttt{PCSetOperators(pc, Amat, Pmat)}, the routine \texttt{PCApply(pc, x, y)} computes \(y = B^{-1}x\) by solving the linear system \(By = x\) with the specified preconditioner method.

Additional preconditioner routines are

\texttt{PCApplyBAorAB(PC pc, PCSide right, Vec x, Vec y, Vec work);} 
\texttt{PCApplyBAorABTranspose(PC pc, PCSide right, Vec x, Vec y, Vec work);} 
\texttt{PCApplyRichardson(PC pc, Vec x, Vec y, Vec work, PetscReal rtol, PetscReal atol, PetscReal \_dtol, PetscInt maxits, PetscBool zeroguess, PetscInt \_*outits, \_PCRichardsonConvergedReason);}
The first two routines apply the action of the matrix followed by the preconditioner or the preconditioner followed by the matrix depending on whether the right is PC_LEFT or PC_RIGHT. The final routine applies its iterations of Richardson’s method. The last three routines are provided to improve efficiency for certain Krylov subspace methods.

A PC context that is no longer needed can be destroyed with the command

```c
PCDestroy(PC *pc);
```

### 4.9 Running PETSc Tests

#### 4.9.1 Quick start with the tests

Users should set `$PETSC_DIR` and `$PETSC_ARCH` before running the tests, or can provide them on the command line as below.

To check if the libraries are working do:

```bash
$ make PETSC_DIR=<PETSC_DIR> PETSC_ARCH=<PETSC_ARCH> check
```

For comprehensive testing of builds, the general invocation from the `$PETSC_DIR` is:

```bash
$ make PETSC_DIR=<PETSC_DIR> $PETSC_ARCH=<PETSC_ARCH> alltests
```

or

```bash
$ make [-j <n>] test PETSC_ARCH=<PETSC_ARCH>
```

For testing `configure` that used the `--prefix` option, the general invocation from the installation (prefix) directory is:

```bash
$ make [-j <n>] -f share/petsc/examples/gmakefile.test test
```

which will create/use the directories `tests/*` in the current directory for generated test files. You may pass an additional argument `TESTDIR=mytests` to place these generated files elsewhere.

For a full list of options, use

```bash
$ make help-test
```

#### 4.9.2 Understanding test output and more information

Depending on your machine’s configuration running the full test suite (above) can take from a few minutes to a couple hours. Note that currently we do not have a mechanism for automatically running the test suite on batch computer systems except to obtain an interactive compute node (via the batch system) and run the tests on that node (this assumes that the compilers are available on the interactive compute nodes.

The test reporting system classifies them according to the Test Anywhere Protocol (TAP).¹¹ In brief, the categories are

- **ok** The test passed.
- **not ok** The test failed.

¹¹ See https://testanything.org/tap-specification.html
The tests are a series of shell scripts, generated by information contained within the test source file, that are invoked by the makefile system. The tests are run in \$PETSC_DIR/$PETSC_ARCH/tests with the same directory as the source tree underneath. For testing installs, the default location is \${PREFIX_DIR}/tests but this can be changed with the TESTDIR location. (See Directory Structure). A label is used to denote where it can be found within the source tree. For example, test vec_vec_tutorials-ex6, which can be run e.g. with

```
$ make test search='vec_vec_tutorials-ex6'
```

(see the discussion of search below), denotes the shell script:

```
$ \$PETSC_DIR/$PETSC_ARCH/tests/vec/vec/tutorials/runex6.sh
```

These shell scripts can be run independently in those directories, and take arguments to show the commands run, change arguments, etc. Use the -h option to the shell script to see these options.

Often, you want to run only a subset of tests. Our makefiles use \gmake's wildcard syntax. In this syntax, % is a wild card character and is passed in using the search argument. Two wildcard characters cannot be used in a search, so the searchin argument is used to provide the equivalent of %pattern% search. The default examples have default arguments, and we often wish to test examples with various arguments; we use the argsearch argument for these searches. Like searchin, it does not use wildcards, but rather whether the string is within the arguments.

Some examples are:

```
$ make test search='ts%' # Run all TS examples
$ make test searchin='tutorials' # Run all tutorials
$ make test search='ts%' searchin='tutorials' # Run all TS tutorials
$ make test argsearch='cuda' # Run examples with cuda in arguments
$ make test test-fail='1'
$ make test query='requires' queryval='*MPI_PROCESS_SHARED_MEMORY*'
```

It is useful before invoking the tests to see what targets will be run. The print-test target helps with this:

```
$ make print-test argsearch='cuda'
```

To see all of the test targets which would be run, this command can be used:

```
$ make print-test
```

To learn more about the test system details, one can look at the the PETSc developers documentation.
4.9.3 Using the test harness for your own code

Select a package name, for example, `mypkg` and create a sub-directory with that name, say `/home/mine/mypackage/src/mypkg`. In any sub-directory of that directory named `tests` or `tutorials` put a PETSc makefile, for example, `src/ts/tutorials/makefile` and standalone test applications that the makefile can compile with, for example

```
$ make mytest
```

Include at the bottom of the test code a formatted comment indicating what tests should be run, see `test_harness`. Also select a directory where you wish the tests to be compiled and run, say `/home/mine/mytests`.

You can build and run the tests with

```
$ make -f ${PETSC_DIR}/gmakefile.test TESTSRCDIR=/home/mine/mypackage/src TESTDIR=/home/mine/mytests pkgs=mypkg
```

There is not yet a mechanism to have your test code also link against your library, contact us for ideas.


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