

## ML 5.0 Smoothed Aggregation User's Guide

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### Abstract

**ML** is a multigrid preconditioning package intended to solve linear systems of equations  $Ax = b$  where  $A$  is a user supplied  $n \times n$  sparse matrix,  $b$  is a user supplied vector of length  $n$  and  $x$  is a vector of length  $n$  to be computed. **ML** should be used on large sparse linear systems arising from partial differential equation (PDE) discretizations. While technically any linear system can be considered, **ML** should be used on linear systems that correspond to things that work well with multigrid methods (e.g. elliptic PDEs). **ML** can be used as a stand-alone package or to generate preconditioners for a traditional iterative solver package (e.g. Krylov methods). We have supplied support for working with the AZTEC 2.1 and AZTECOO iterative packages [19]. However, other solvers can be used by supplying a few functions.

This document describes one specific algebraic multigrid approach: smoothed aggregation. This approach is used within several specialized multigrid methods: one for the eddy current formulation for Maxwell's equations, and a multilevel and domain decomposition method for symmetric and non-symmetric systems of equations (like elliptic equations, or compressible and incompressible fluid dynamics problems). Other methods exist within **ML** but are not described in this document. Examples are given illustrating the problem definition and exercising multigrid options.

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# 1 Notational Conventions

In this guide, we show typed commands in this font:

```
% a_really_long_command
```

The character % indicates any shell prompt<sup>1</sup>. Function names are shown as `ML_Gen_Solver`. Names of packages or libraries as reported in small caps, as EPETRA. Mathematical entities are shown in italics.

## 2 Overview

This guide describes the use of an algebraic multigrid method within the **ML** package. The algebraic multigrid method can be used to solve linear system systems of type

$$Ax = b \tag{1}$$

where  $A$  is a user supplied  $n \times n$  sparse matrix,  $b$  is a user supplied vector of length  $n$  and  $x$  is a vector of length  $n$  to be computed. **ML** is intended to be used on (distributed) large sparse linear systems arising from partial differential equation (PDE) discretizations. While technically any linear system can be considered, **ML** should be used on linear systems that correspond to things that work well with multigrid methods (e.g. elliptic PDEs).

The **ML** package is used by creating a **ML** object and then associating a matrix,  $A$ , and a set of multigrid parameters which describe the specifics of the solver. Once created and initialized, the **ML** object can be used to solve linear systems.

This manual is structured as follows. Multigrid and multilevel methods are briefly recalled in Section 3. A quick start is reported in Section 4. The process of configuring and building **ML** is outlined in Section 5. Section 6 shows the basic usage of **ML** as a black-box preconditioner for EPETRA matrices. The definition of (parallel) preconditioners using `ML_Epetra::MultiLevelPreconditioner` is detailed. This class only requires the linear system matrix, and a list of options. Available parameters for `ML_Epetra::MultiLevelPreconditioner` are reported in Section 6.4. Section 7 reports how to use the Maxwell solvers of **ML**. More advanced uses of **ML** are presented in Section 8. Here, we present how to define and fine-tune smoothers, coarse grid solver, and the multilevel hierarchy. Multigrid options are reported in Section 9. Smoothing options are reported in Section 10, where we also present how to construct a user's defined smoother. Advanced usage of **ML** with EPETRA objects is reported in Section 11. Section 12 reports how to define matrices in **ML** format without depending on EPETRA.

## 3 Multigrid Background

A brief multigrid description is given (see [3], [9], or [10] for more information). A multigrid solver tries to approximate the original PDE problem of interest on a hierarchy of grids and use 'solutions' from coarse grids to accelerate the convergence on the finest grid. A simple multilevel iteration is illustrated in Figure 1. In the above method, the  $S_k^1()$ 's and  $S_k^2()$ 's

---

<sup>1</sup>For simplicity, commands are shown as they would be issued in a Linux or Unix environment. Note, however, that **ML** has and can be built successfully in a Windows environment.

```

/* Solve  $A_k u = b$  ( $k$  is current grid level)          */
proc multilevel( $A_k, b, u, k$ )
   $u = S_k^1(A_k, b, u)$ ;
  if (  $k \neq \mathbf{Nlevel} - 1$  )
     $P_k = \text{determine\_interpolant}( A_k )$ ;
     $\hat{r} = P_k^T(b - A_k u)$  ;
     $\hat{A}_{k+1} = P_k^T A_k P_k$ ;   $v = 0$ ;
    multilevel( $\hat{A}_{k+1}, \hat{r}, v, k + 1$ );
     $u = u + P_k v$ ;
     $u = S_k^2(A_k, b, u)$ ;

```

Figure 1: High level multigrid V cycle consisting of ‘Nlevel’ grids to solve (1), with  $A_0 = A$ .

are approximate solvers corresponding to  $k$  steps of pre and post smoothing, respectively. These smoothers are discussed in Section 9. For now, it suffices to view them as basic iterative methods (e.g. Gauss-Seidel) which effectively smooth out the error associated with the current approximate solution. The  $P_k$ ’s (interpolation operators that transfer solutions from coarse grids to finer grids) are the key ingredient that are determined automatically by the algebraic multigrid method<sup>2</sup>. For the purposes of this guide, it is important to understand that when the multigrid method is used, a hierarchy of grids, grid transfer operators ( $P_k$ ), and coarse grid discretizations ( $A_k$ ) are created. To complete the specification of the multigrid method, smoothers must be supplied on each level. There are several smoothers within **ML** or an iterative solver package can be used, or users can write their own smoother (see Section 9).

## 4 Quick Start

This section is intended for the impatient user. It’s assumed that you’ve already have a local copy of TRILINOS<sup>3</sup>. Using the instructions here, your build of TRILINOS will have the following libraries: AZTECOO, EPETRA, EPETRAEXT, IFPACK, LOCA, **ML**, NEW\_PACKAGE, NOX, AMESOS and TEUCHOS.

1. `cd` into the TRILINOS directory.
2. Make a build directory, e.g., `mkdir sandbox`.
3. `cd sandbox`.
4. Configure TRILINOS:
  - (a) `../configure --enable-teuchos --enable-amesos --enable-aztecoo --enable-epetra`  
if you do not want to use MPI.
  - (b) `../configure --enable-teuchos`  
`--enable-amesos --enable-aztecoo --enable-epetra --with-mpi-compilers=/usr/loc`  
(or wherever your MPI compilers are located) to use MPI.

<sup>2</sup>The  $P_k$ ’s are usually determined as a preprocessing step and not computed within the iteration.

<sup>3</sup>Please refer to the web page <http://software.sandia.gov/trilinos> to know how to obtain a copy of TRILINOS.

5. Build TRILINOS: `make`.<sup>4</sup>
6. If your build finished without errors, you should see the directory `Trilinos/sandbox/packages/`, with subdirectories below that for each individual library. **ML**'s subdirectory, `ml`, should contain files `config.log`, `config.status`, `Makefile`, and `Makefile.export`, and directories `src` and `examples`. Directory `src` contains object files and `libml.a`. Directory `examples` contains executables with extension `.exe`, symbolic links to the corresponding source code, and object files. Directory `test` is intended primarily for developers and can be ignored.
7. Look in `Trilinos/sandbox/packages/ml/examples` for examples of how to use **ML**. File `Trilinos/packages/ml/examples/README` suggests how to use the examples.

## 5 Configuring and Building ML

**ML** is configured and built using the GNU `autoconf` [7] and `automake` [8] tools. It can be configured and build as a standalone package without or with **AZTEC** 2.1 support (as detailed in Section 5.1 and 5.2), or as a part of the **TRILINOS** framework [11] (as described in Section 5.3). Even though **ML** can be compiled and used as a standalone package, the recommended approach is to build **ML** as part of the **TRILINOS** framework, as a richer set of features are then available.

**ML** has been configured and built successfully on a wide variety of operating systems, and with a variety of compilers (as reported in Table 1).

Operating System	Compilers(s)
Linux	GNU, Intel, Portland Group
MAC OS X	GNU
IRIX N32, IRIX 64, HPUX, Solaris, DEC	native
ASC Red Storm	native and Portland Group
Windows	Microsoft and Cygwin

Table 1: Main operating systems and relative compilers supported by **ML**.

Although it is possible to configure directly in the **ML** home directory, we strongly advise against this. Instead, we suggest working in an independent directory and configuring and building there.

### 5.1 Building in Standalone Mode

To configure and build **ML** as a standalone package without any **AZTEC** support, do the following. It's assumed that the shell variable `$ML_HOME` identifies the **ML** directory.

```
% cd $ML_HOME
% mkdir standalone
% cd standalone
```

---

<sup>4</sup>If you are using GNU's `make` on a machine with more than one processor, then you can speed up the build with `make -j XX`, where `XX` is the number of processors. You can also reduce the size of the link lines with the option `--with-gnumake`.

```
% $ML_HOME/configure --disable-epetra --disable-aztecoo \
    --prefix=$ML_HOME/standalone
% make
% make install
```

The **ML** library file `libml.a` and the header files will be installed in the directory specified in `--prefix`.

## 5.2 Building with AZTEC 2.1 Support

To enable the supports for AZTEC 2.1, **ML** must be configured with the options reported in the previous section, plus `--with-ml_aztec2.1` (defaulted to `no`).

All of the AZTEC 2.1 functionality that **ML** accesses is contained in the file `ml_aztec_utils.c`. In principal by creating a similar file, other solver packages could work with **ML** in the same way. For the AZTEC users there are essentially three functions that are important. The first is `AZ_ML_Set_Amat` which converts AZTEC matrices into **ML** matrices by making appropriate **ML** calls (see Section 12.1 and Section 12.2). It is important to note that when creating **ML** matrices from AZTEC matrices information is not copied. Instead, wrapper functions are made so that **ML** can access the same information as AZTEC. The second is `ML_Gen_SmoothingAztec` that is used for defining AZTEC iterative methods as smoothers (discussed in Section 9). The third function, `AZ_set_ML_preconditioner`, can be invoked to set the AZTEC preconditioner to use the multilevel ‘V’ cycle constructed in **ML**. Thus, it is possible to invoke several instances of AZTEC within one solve: smoother on different multigrid levels and/or outer iterative solve.

## 5.3 Building with TRILINOS Support (RECOMMENDED)

We recommend to configure and build **ML** as part of the standard TRILINOS build and configure process. In fact, **ML** is built by default if you follow the standard TRILINOS configure and build directions. Please refer to the TRILINOS documentation for information about the configuration and building of other TRILINOS packages.

To configure and build **ML** through TRILINOS, you may need do the following (actual configuration options may vary depending on the specific architecture, installation, and user’s need). It’s assumed that shell variable `$TRILINOS_HOME` (here introduced for the sake of simplicity only) identifies the TRILINOS directory, and, for example, that we are compiling under LINUX and MPI.

```
% cd $TRILINOS_HOME
% mkdir LINUX_MPI
% cd LINUX_MPI
% $TRILINOS_HOME/configure \
    --enable-teuchos \
    --enable-amesos \
    --with-mpi-compilers \
    --prefix=$TRILINOS_HOME/LINUX_MPI
% make
% make install
```



If required, other TRILINOS and **ML** options can be specified in the configure line. A complete list of **ML** options is given in Section 5.3.1 and 5.3.2. You can also find a complete list and explanations by typing `./configure --help` in the **ML** home directory.

### 5.3.1 Enabling Third Party Library Support

**ML** can be configured with the following third party libraries (TPLs): SUPERLU, SUPERLU\_DIST, PARASAILS, ZOLTAN, METIS, and PARMETIS. It can take advantage of the following TRILINOS packages: IFPACK, TEUCHOS, TRIUTILS, AMESOS, EPETRAEXT. Through AMESOS, **ML** can interface with the direct solvers KLU, UMFPACK, SUPERLU, SUPERLU\_DIST<sup>5</sup>, MUMPS. It is assumed that you have already built the appropriate libraries (e.g., `libsuperlu.a`) and have the header files. To configure **ML** with one of the above TPLs, you must enable the particular TPL interface in **ML**.

The same configure options that enable certain other Trilinos packages also enable the interfaces to those packages within **ML**.

<code>--enable-epetra</code>	Enable support for the EPETRA package. (Recommended)
<code>--enable-epetraext</code>	Enable support for the EPETRAEXT package. (Recommended)
<code>--enable-aztecoo</code>	Enable support for the AZTECOO package. (Recommended)
<code>--enable-amesos</code>	Enables support for the AMESOS package. AMESOS is an interface with several direct solvers. <b>ML</b> supports UMFPACK [5], KLU, SUPERLU_DIST (1.0 and 2.0), MUMPS [1]. This package is necessary to use the AMESOS interface to direct solvers. (Recommended)
<code>--enable-teuchos</code>	Enables support for the TEUCHOS package. This package is necessary to use the <code>MLEpetra::MultiLevelPreconditioner</code> class. (Recommended)
<code>--enable-triutils</code>	Enables support for the TRIUTILS package. <b>ML</b> uses TRIUTILS in some examples and tests to create the linear system matrix.
<code>--enable-galeri</code>	Enables support for the GALERI package. <b>ML</b> uses GALERI in some examples and tests to create the linear system matrix.
<code>--enable-ifpack</code>	Enable support for the IFPACK package [12], to use IFPACK factorizations as smoothers for <b>ML</b> .

---

<sup>5</sup>Currently, **ML** can support SUPERLU\_DIST directly (without AMESOS support), or through AMESOS.

To know whether the `--enable-package` options below reported are enabled or disabled by default<sup>6</sup>, please consult the configure at the Trilinos level by typing

```
$TRILINOS_HOME/configure --help
```

The following configure line options enable interfaces in **ML** to certain TPLs. By default, all the `--with-ml_TPL` options are disabled. For the most up-to-date list, please type `configure --help` at the **ML** package level.

<code>--with-ml_parasails</code>	Enables interface for PARASAILS [4].
<code>--with-ml_metis</code>	Enables interface for METIS [16].
<code>--with-ml_parmetis2x</code>	Enables interface for PARMETIS, version 2.x.
<code>--with-ml_parmetis3x</code>	Enables interface for PARMETIS [15], version 3.x.
<code>--with-ml_superlu</code>	Enables <b>ML</b> 's direct interface for serial SUPERLU [6] 1.0. This interface is deprecated in favor of the AMESOS interface. Note: you cannot configure <b>ML</b> with both this option and <code>--enable-amesos</code> .
<code>--with-ml_superlu2</code>	Enables <b>ML</b> 's direct interface for serial SUPERLU [6] 2.0. This interface is deprecated in favor of the AMESOS interface. Note: you cannot configure <b>ML</b> with both this option and <code>--enable-amesos</code> .
<code>--with-ml_superlu_dist</code>	Enables <b>ML</b> interface for SUPERLU_DIST [6]. This interface is deprecated in favor of the AMESOS interface. Note: you cannot configure <b>ML</b> with both this option and <code>--enable-amesos</code> .

If one of the above options is enabled, then the user must specify the location of the header files, with the option

```
--with-incdirs=<include-locations>
```

(Header files for TRILINOS libraries are automatically located if **ML** is built through the TRILINOS configure.) In order to link the **ML** examples, the user must indicate the location of all the enabled packages' libraries , with the option

```
--with-ldflags="<lib-locations>"
--with-libs="<lib-list>"
```

The user might find useful the option

```
--disable-examples
```

---

<sup>6</sup>This refers to configurations from the Trilinos top level (that is, using `$TRILINOS_HOME/configure`). If **ML** is configured from the **ML** level (that is, the user directly calls `$ML_HOME/configure`), then all the `--enable-package` are off by default.

which turns off compilation and linking of all Trilinos examples, or

`--disable-ml-examples`

which turns off compilation and linking of **ML** examples.

Here is an example configure line, where METIS, PARMETIS, ZOLTAN, and SUPERLU are all used:

```
./configure --with-mpi-compilers="/usr/local/mpich/bin" CXXFLAGS="-O3" CFLAGS="-O3"
FFLAGS="-O3" --cache-file=config.cache --with-gnumake --with-mlmetis --with-mlzoltan
--with-mlparmetis3x --with-ldflags="-L/usr/local/superlu-3.0/lib -L/usr/local/parmetis
-L/usr/local/zoltan/lib" --with-libs="-lsuperlu -lparmetis -lmetis -lzoltan"
--with-incdirs="-I/usr/local/superlu-3.0/SRC -I/usr/local/parmetis-3.1 -I/usr/local/par
-I/usr/local/parmetis-3.1/ParMETISLib -I/usr/local/zoltan/include"
```

More details about the installation of TRILINOS can be found at the TRILINOS web site,

<http://software.sandia.gov/Trilinos>

and [18, Chapter 1].

### 5.3.2 Enabling Profiling

All of the options below are disabled by default.

`--enable-ml-timing` This prints out timing of key **ML** routines.

`--enable-ml-flops` This enables printing of flop counts.

Timing and flop counts are printed when the associated object is destroyed.

## 6 ML and Epetra: Getting Started with the MultiLevelPreconditioner Class

In this Section we show how to use **ML** as a preconditioner to EPETRA and AZTECOO through the MultiLevelPreconditioner class. This class is derived from the Epetra\_RowMatrix class, and is defined in the ML\_Epetra namespace.

The MultiLevelPreconditioner class automatically constructs all the components of the preconditioner, using the parameters specified in a TEUCHOS parameter list. The constructor of this class takes as input an Epetra\_RowMatrix pointer<sup>7</sup> and a TEUCHOS parameter list<sup>8</sup>.

In order to compile, it may also be necessary to include the following files: `ml_config.h` (as first **ML** include), `Epetra_ConfigDefs.h` (as first EPETRA include), `Epetra_RowMatrix.h`, `Epetra_MultiVector.h`, `Epetra_LinearProblem.h`, and `AztecOO.h`. Check the EPETRA and AZTECOO documentation for more details. Additionally, the user must include the header file "`ml_MultiLevelPreconditioner.h`". Also note that the macro `HAVE_CONFIG_H` must be defined either in the user's code or as a compiler flag.

### 6.1 Example 1: `ml_preconditioner.cpp`

We now give a very simple fragment of code that uses the MultiLevelPreconditioner. For the complete code, see `$ML_HOME/examples/BasicExamples/ml_preconditioner.cpp`. The linear operator **A** is derived from an Epetra\_RowMatrix, **Solver** is an AztecOO object, and **Problem** is an Epetra\_LinearProblem object.

```
#include "ml_include.h"
#include "ml_MultiLevelPreconditioner.h"
#include "Teuchos_ParameterList.hpp"

...

Teuchos::ParameterList MList;

// set default values for smoothed aggregation in MList
ML_Epetra::SetDefaults("SA",MList);

// overwrite with user's defined parameters
MList.set("max levels",6);
MList.set("increasing or decreasing","decreasing");
MList.set("aggregation: type", "MIS");
MList.set("coarse: type","Amesos-KLU");

// create the preconditioner
```

---

<sup>7</sup>Note that not all Epetra matrices can be used with ML. Clearly, the input matrix must be a square matrix. Besides, it is supposed that the `OperatorDomainMap()`, the `OperatorRangeMap()` and the `RowMatrixRowMap()` of the matrix all coincide, and that each row is assigned to exactly one process.

<sup>8</sup>In order to use the MultiLevelPreconditioner class, **ML** must be configured with options `-enable-epetra` `--enable-teuchos`.

```

ML_Epetra::MultiLevelPreconditioner* MLPrec =
    new ML_Epetra::MultiLevelPreconditioner(A, MLList, true);

// create an Aztec00 solver
Aztec00 Solver(Problem)

// set preconditioner and solve
Solver.SetPrecOperator(MLPrec);
Solver.SetAztecOption(AZ_solver, AZ_gmres);
Solver.Iterate(Niters, 1e-12);

...

delete MLPrec;

```

A MultiLevelPreconditioner may be defined as follows. First, the user defines a TEUCHOS parameter list<sup>9</sup>. Table 2 briefly reports the most important methods of this class.

<code>set(Name,Value)</code>	Add entry <code>Name</code> with value and type specified by <code>Value</code> . Any C++ type (like <code>int</code> , <code>double</code> , a pointer, etc.) is valid.
<code>get(Name,DefValue)</code>	Get value (whose type is automatically specified by <code>DefValue</code> ). If not present, return <code>DefValue</code> .
<code>subList(Name)</code>	Get a reference to sublist <code>List</code> . If not present, create the sublist.

Table 2: Some methods of Teuchos::ParameterList class.

Input parameters are set via method `set(Name,Value)`, where `Name` is a string containing the parameter name, and `Value` is the specified parameter value, whose type can be any C++ object or pointer. A complete list of parameters available for class MultiLevelPreconditioner is reported in Section 6.4.

The parameter list is passed to the constructor, together with a pointer to the matrix, and a boolean flag. If this flag is set to `false`, the constructor will not create the multilevel hierarchy until when `MLPrec->ComputePreconditioner()` is called. The hierarchy can be destroyed using `MLPrec->DestroyPreconditioner()`<sup>10</sup>. For instance, the user may define a code like:

```

// A is still not filled with numerical values
ML_Epetra::MultiLevelPreconditioner* MLPrec =
    new ML_Epetra::MultiLevelPreconditioner(A, MLList, false);

// compute the elements of A
...
// now compute the preconditioner
MLPrec->ComputePreconditioner();

```

<sup>9</sup>See the TEUCHOS documentation for a detailed overview of this class.

<sup>10</sup>We suggest to always create the preconditioning object with `new` and to delete it using `delete`. Some MPI calls occur in `DestroyPreconditioner()`, so the user should not call `MPI_Finalize()` or delete the communicator used by `ML` before the preconditioning object is destroyed.

```

// solve the linear system
...
// destroy the previously define preconditioner, and build a new one
MLPrec->DestroyPreconditioner();

// re-compute the elements of A
// now re-compute the preconditioner, using either
MLPrec->ComputePreconditioner();
// or
MLPrec->ReComputePreconditioner();

// re-solve the linear system

// .. finally destroy the object
delete MLPrec;

```

In this fragment of code, the user defines the **ML** preconditioner, but the preconditioner is created only with the call `ComputePreconditioner()`. This may be useful, for example, when **ML** is used in conjunction with nonlinear solvers (like NOX [17]). Method `ReComputePreconditioner()` can be used to recompute the preconditioner using already available information about the aggregates. `ReComputePreconditioner()` reuses the already computed tentative prolongator, then recomputes the smoothed prolongators and the other components of the hierarchy, like smoothers and coarse solver<sup>11</sup>.

## 6.2 Example 2: ml\_2level\_DD.cpp

In the second example, a two level domain decomposition preconditioner is constructed. The coarse space is defined using aggregation. It is worthwhile to compare the parameters selected here to the default parameters stated in Table 6.

File `$ML_HOME/examples/TwoLevelDD/ml_2level_DD.cpp` reports the entire code. In the example, the linear system matrix **A**, is an `Epetra.CrsMatrix` corresponding to the discretization of a Laplacian on a 2D Cartesian grid. The solution vector and right-hand side are **x** and **b** respectively.

The AztecOO linear problem is defined as

```

Epetra_LinearProblem problem(&A, &x, &b);
AztecOO solver(problem);

```

We create the TEUCHOS parameter list as follows:

```

ParameterList MLList;
ML_Epetra::SetDefaults("DD", MLList);
MLList.set("max levels",2);
MLList.set("increasing or decreasing","increasing");

MLList.set("aggregation: type", "METIS");

```

<sup>11</sup>Note that the hierarchy produced by `ReComputePreconditioner()` can differ from that produced by `ComputePreconditioner()` for non-zero threshold values.

```

MLList.set("aggregation: nodes per aggregate", 16);
MLList.set("smoother: pre or post", "both");
MLList.set("coarse: type", "Amesos-KLU");
MLList.set("smoother: type", "Aztec");

```

The last option tells **ML** to use the AZTEC preconditioning function as a smoother. All AZTEC preconditioning options can be used as **ML** smoothers. AZTEC requires an integer vector **options** and a double vector **params**. Those can be defined as follows:

```

int options[AZ_OPTIONS_SIZE];
double params[AZ_PARAMS_SIZE];
AZ_defaults(options, params);
options[AZ_precond] = AZ_dom_decomp;
options[AZ_subdomain_solve] = AZ_icc;
MLList.set("smoother: Aztec options", options);
MLList.set("smoother: Aztec params", params);

```

The last two commands set the pointer to **options** and **params** in the parameter list<sup>12</sup>.

The **ML** preconditioner is created as in the previous example,

```

ML_Epetra::MultiLevelPreconditioner* MLPrec =
    new ML_Epetra::MultiLevelPreconditioner(A, MLList, true);

```

and we can check that no options have been misspelled, using

```

MLPrec->PrintUnused();

```

The AztecOO solver is called using, for instance,

```

solver.SetPrecOperator(MLPrec);
solver.SetAztecOption(AZ_solver, AZ_cg_condnum);
solver.SetAztecOption(AZ_kspace, 160);
solver.Iterate(1550, 1e-12);

```

Finally, some (limited) information about the preconditioning phase are obtained using

```

cout << MLPrec->GetOutputList();

```

Note that the input parameter list is *copied* in the construction phase, hence later changes to **MLList** will not affect the preconditioner. Should the user need to modify parameters in the **MLPrec**'s internally stored parameter list, he or she can get a reference to the internally stored list:

```

ParameterList& List = MLPrec->GetList();

```

and then directly modify **List**. Method **GetList()** should be used carefully, as a change to **List** may modify the behavior of **MLPrec**.

### 6.3 Tips for Achieving Good Parallel Performance

This section gives a few tips on tuning **ML**'s parallel performance.

<sup>12</sup>Only the *pointer* is copied in the parameter list, not the array itself. Therefore, **options** and **params** should not go out of scope before the destruction of the preconditioner.

Uncoupled	Attempts to construct aggregates of optimal size ( $3^d$ nodes in $d$ dimensions). Each process works independently, and aggregates cannot span processes.
Coupled MIS	As <b>Uncoupled</b> , but aggregates can span processes (deprecated). Uses a maximal independent set technique to define the aggregates. Aggregates can span processes. May provide better quality aggregates than either <b>Coupled</b> or <b>uncoupled</b> . Computationally more expensive than either because it requires matrix-matrix product.
Uncoupled-MIS	Uses <b>Uncoupled</b> for all levels until there is 1 aggregate per processor, then switches over to <b>MIS</b> . The coarsening scheme on a given level cannot be specified with this option.
METIS	Use a graph partitioning algorithm to create the aggregates, working process-wise. The number of nodes in each aggregate is specified with the option <b>aggregation: nodes per aggregate</b> . Requires <b>ML</b> to be configured with <b>--with-mlmetis</b> .
ParMETIS	As <b>METIS</b> , but partition the global graph. Requires <b>--with-mlparmetis2x</b> or <b>--with-mlparmetis3x</b> . Aggregates can span arbitrary number of processes. Global number of aggregates can be specified with the option <b>aggregation: global number</b> .

Table 3: ML\_Epetra::MultiLevelPreconditioner: Available coarsening schemes.

### 6.3.1 Hints for the Impatient User

1. Use the matrix repartitioning option. This is discussed in detail in the next section, §6.3.2.
2. In order to reduce message-passing latency, try limiting the number of multigrid levels to 3 or 4. The syntax is described in §6.4.2.
3. Instead of doing a direct solve on the coarsest level, try a few smoothing sweeps instead. See §6.4.5.
4. For a symmetric positive definite linear system, choose a smoother whose computational kernel is a matvec, such as the Chebyshev polynomial smoother. See §6.4.4.

### 6.3.2 Dynamic Load-balancing

**ML** supports dynamic load-balancing of coarse-level matrices in the multigrid preconditioner. This is extremely important for two reasons. First, it is not unusual for the message-passing latency on a coarse level to be roughly equivalent to the latency of the fine level, even though the number of matrix unknowns on the coarse level is much smaller than on the fine level. Second, as the number of unknowns per processor becomes small, **ML**'s aggregation methods become less effective. The coarsening rate may drop, thus leading to more multigrid levels than one might observe in serial.

We now give two code fragments to demonstrate how to use the repartitioning options. The complete description of options that control repartitioning is given in §6.4.6. The following code demonstrates repartitioning for a matrix with unknowns that are node-based.



Jacobi	Point-Jacobi. Damping factor is specified using <code>smoother: damping factor</code> , and the number of sweeps with <code>smoother: sweeps</code>
Gauss-Seidel	Point Gauss-Seidel. Damping factor is specified using <code>smoother: damping factor</code> , and the number of sweeps with <code>smoother: sweeps</code>
symmetric Gauss-Seidel	Point symmetric Gauss-Seidel. Damping factor is specified using <code>smoother: damping factor</code> , and the number of sweeps with <code>smoother: sweeps</code>
Aztec	Use AZTECOO's built-in preconditioning functions as smoothers. Or, if <code>smoother: Aztec as solver</code> is <code>true</code> , use approximate solutions with AZTECOO (with <code>smoothers: sweeps</code> iterations as smoothers. The AZTECOO vectors <code>options</code> and <code>params</code> can be set using <code>smoother: Aztec options</code> and <code>smoother: Aztec params</code> .
MLS	Use MLS smoother. The polynomial order is specified by <code>smoother: MLS polynomial order</code> , and the alpha value by <code>smoother: MLS alpha</code> .
user-defined	User-defined smoothing function. The function must conform to the prototype <code>int (*)(ML_Smoother *, int inLength, double *invec, int outLength, double *outvec)</code> . The number of sweeps is controlled via <code>smoother: sweeps</code> . See <code>ml/examples/BasicExamples/ml_user_smoothing.cpp</code> .

Table 4: ML\_Epetra::MultiLevelPreconditioner: Commonly used smoothers.

```

Teuchos::ParameterList MList;
MList.set("repartition: enable",1);
MList.set("repartition: max min ratio",1.3);
MList.set("repartition: min per proc",500);
MList.set("repartition: partitioner","Zoltan");
MList.set("repartition: Zoltan dimensions",2);

// Zoltan requires coordinates
double *xcoord, *ycoord, *zcoord;

// user must load the coordinates, of coarse ...

MList.set("x-coordinates", xcoord);
MList.set("y-coordinates", xcoord);

```

The following code demonstrates repartitioning for a system arising from the eddy current approximations to Maxwell's equations.

```

Teuchos::ParameterList MList;
// these correspond to the main edge matrix
MList.set("repartition: enable",1);
MList.set("repartition: max min ratio",1.3);
MList.set("repartition: min per proc",1000);

```

Jacobi	Use <code>coarse: sweeps</code> steps of Jacobi (with damping parameter <code>coarse: damping parameter</code> ) as a solver.
Gauss-Seidel	Use <code>coarse: sweeps</code> steps of Gauss-Seidel (with damping parameter <code>coarse: damping parameter</code> ) as a solver.
MLS	Use degree <code>coarse: MLS polynomial order</code> Chebyshev polynomial as a solver.
Hiptmair	Use <code>coarse: sweeps</code> steps of two-stage Hiptmair smoothing, with subsmoothen <code>coarse: subsmoothen type</code> , as a solver.
Amesos-KLU	Use KLU through AMESOS. Coarse grid problem is shipped to proc 0, solved, and solution is broadcast
Amesos-Superlu	Use SUPERLU, version 3.0, through AMESOS.
Amesos-UMFPACK	Use UMFPACK through AMESOS. Coarse grid problem is shipped to proc 0, solved, and solution is broadcasted.
Amesos-Superludist	Use SUPERLU_DIST through AMESOS.
Amesos-MUMPS	Use double precision version of MUMPS through AMESOS.
user-defined	User-defined smoothing function. The function must conform to the prototype <code>int (*)(ML_Smoothen *, int inLength, double *invec, int outLength, double *outvec)</code> . The number of sweeps is controlled via <code>coarse: sweeps</code> . See <code>ml/examples/BasicExamples/ml_user_smoothing.cpp</code> .
SuperLU	Use <b>ML</b> interface to SUPERLU(deprecated).

Table 5: ML\_Epetra::MultiLevelPreconditioner: Some of the available coarse matrix solvers. Note: Amesos solvers requires **ML** to be configured with `with-ml_amesos`, and Amesos to be properly configured to support the specified solver.

```

MLList.set("repartition: partitioner","Zoltan");

// Zoltan requires coordinates
double *xcoord, *ycoord, *zcoord;

// user must load the coordinates, of coarse ...

MLList.set("x-coordinates", xcoord);
MLList.set("y-coordinates", ycoord);
MLList.set("z-coordinates", zcoord);

// these correspond to the auxiliary node matrix
MLList.set("repartition: node max min ratio",1.5);
MLList.set("repartition: node min per proc",300);
MLList.set("repartition: node partitioner","Zoltan");
MLList.set("repartition: Zoltan dimensions",3);

double *node_xcoord, *node_ycoord, *node_zcoord;

// user must load the coordinates, of coarse ...

MLList.set("x-coordinates", node_xcoord);

```

```
MLList.set("y-coordinates", node_ycoord);
MLList.set("z-coordinates", node_zcoord);
```

## 6.4 List of All Parameters for MultiLevelPreconditioner Class

In this section we give general guidelines for using the MultiLevelPreconditioner class effectively. The complete list of input parameters is also reported. It is important to point out that some options can be effectively used only if **ML** has been properly configured. In particular:

- METIS aggregation scheme requires `--with-ml_metis`, or otherwise the code will include all nodes in the calling processor in a unique aggregate;
- PARMETIS aggregation scheme requires `--with-ml_metis --enable-epetra` and `--with-ml_parmetis2x` or `--with-ml_parmetis3x`.
- AMESOS coarse solvers require `--enable-amesos --enable-epetra --enable-teuchos`. Moreover, AMESOS must have been configured to support the requested coarse solver. Please refer to the AMESOS documentation for more details.
- IFPACK smoothers require `--enable-ifpack --enable-epetra --enable-teuchos`.
- PARASAILS smoother requires `--with-ml_parasails`.

**Note that, in the parameters name, spaces are important: Do not include non-required leading or trailing spaces, and do separate words by just one space! Misspelled parameters will not be detected.** One may find useful to print unused parameters by calling `PrintUnused()` *after* the construction of the multilevel hierarchy.

Some of the parameters that affect the MultiLevelPreconditioner class can in principle be different from level to level. By default, the set method for the MultiLevelPreconditioner class affects all levels in the multigrid hierarchy. In order to change a setting on a particular level (say, `d`), the string `"(level d)"` is appended to the option string (note that a space must separate the option and the level specification). For instance, assuming decreasing levels starting from 4, one could set the aggregation schemes as follows:

```
MLList.set("aggregation: type","Uncoupled");
MLList.set("aggregation: type (level 1)","METIS");
MLList.set("aggregation: type (level 3)","MIS");
```

If the finest level is 0, and one has 5 levels, the code will use `Uncoupled` for level 0, `METIS` for levels 1 and 2, then `MIS` for levels 3 and 4. Parameters that can be set differently on individual levels are denoted with the symbol `*` (this symbol is not part of the parameter name).

### 6.4.1 General Options

output	[int] Controls the amount of information printed to the screen. Ranges from 0 to 10 (0 is no output, and 10 is incredibly detailed output). Default: 10.
print unused	[int] If non-negative, will print all the unused parameters on the specified processor. If -1, the unused parameters will be printed on all processes. If -2, nothing will be printed. Default: -2.
PDE equations	[int] Number of PDE equations for each grid node. This value is not considered for <code>Epetra_VbrMatrix</code> objects, as in this case is obtained from the block map used to construct the object. Note that only block maps with constant element size can be considered. Default: 1.

### 6.4.2 Multigrid Cycle Options

cycle applications	[int] Number of applications of the multilevel V-cycle. Default: 1.
max levels	[int] Maximum number of levels. Default: 10.
increasing or decreasing	[string] If set to <code>increasing</code> , level 0 will correspond to the finest level. If set to <code>decreasing</code> , <code>max levels - 1</code> will correspond to the finest level. Default: <code>increasing</code> .
prec type	[string] Multigrid cycle type. Possible values: MGW, MGW, full-MGW, one-level-postsmoothing, two-level-additive, two-level-hybrid, two-level-hybrid2, projected MGW. Default: MGW.
projected modes	[double**] Array of double vectors containing modes that should be projected out before and after the AMG V-cycle is applied. For use with the <code>projected</code> MGW option. Default: NULL.

number of projected modes	[int] Number of modes that are to be projected out before and after the AMG V-cycle is applied. For use with the <code>projected</code> MGv option. Possible values: 1,2, or 3. Default: 0.
---------------------------	---

#### 6.4.3 Aggregation Parameters

aggregation: type *	[string] Define the aggregation scheme. See Table 3. Default: <code>Uncoupled</code> .
aggregation: threshold	[double] Dropping threshold in aggregation. Default: 0.0.
aggregation: damping factor	[double] Damping factor for smoothed aggregation. Default: 4/3.
aggregation: smoothing sweeps	[int] Number of smoothing sweeps to use in the prolongator smoother. This should be used in conjunction with larger aggregates, from using either <code>METIS</code> or <code>ParMETIS</code> aggregation schemes. If sweeps is equal to 1, then the value of <code>aggregation: damping factor</code> is used. If sweeps is greater than 1, the damping factors are calculated automatically. Default: 1.
aggregation: global aggregates *	[int] Defines the global number of aggregates (only for <code>METIS</code> and <code>ParMETIS</code> aggregation schemes).
aggregation: local aggregates *	[int] Defines the number of aggregates of the calling processor (only for <code>METIS</code> and <code>ParMETIS</code> aggregation schemes). Note: this value overwrites <code>aggregation: global aggregates</code> .
aggregation: nodes per aggregate *	[int] Defines the number of nodes to be assigned to each aggregate (only for <code>METIS</code> and <code>ParMETIS</code> aggregation schemes). Note: this value overwrites <code>aggregation: local aggregates</code> . If none among <code>aggregation: global aggregates</code> , <code>aggregation: local aggregates</code> and <code>aggregation: nodes per aggregate</code> is specified, the default value is 1 aggregate per process.

eigen-analysis: type	[string] Defines the numerical scheme to be used to compute an estimation of the maximum eigenvalue of $D^{-1}A$ , where $D = \text{diag}(A)$ (for smoothed aggregation only). Choices are: <b>cg</b> (10 steps of conjugate gradient method), <b>Anomr</b> (the A-norm of the matrix), <b>Anasazi</b> (use the Arnoldi method), <b>power-method</b> . Default: <b>Anorm</b> .
aggregation: next-level aggregates per process *	[int] Defines the maximum number of next-level matrix rows per process (only for ParMETIS aggregation scheme). Default: 128.
aggregation: use tentative restriction	[bool] If <b>true</b> , then the final restriction operator at level $\ell$ is computed as $R_\ell = P_{\ell,0}^T$ , where $P_{\ell,0}$ is the tentative prolongator. Note that the final prolongator is still computed as specified by parameter <b>aggregation: damping factor</b> . Default: <b>false</b> .
aggregation: symmetrize	[bool] If <b>true</b> , then the final prolongator for level $\ell$ is computed as $P_\ell = (I - \omega/\lambda_{\max}(A + A^T))P_{0,\ell}$ , where $\omega$ is the damping parameter specified using <b>aggregation: damping factor</b> , $\lambda_{\max}$ is the maximum eigenvalue of $D^{-1}A$ , and $P_{0,\ell}$ is the tentative prolongator at level $\ell$ . The restriction is defined as $R_\ell = P_\ell^T$ . Default: <b>false</b> .

#### 6.4.4 Smoothing Parameters

smoother: sweeps *	[int] Number of sweeps of smoother. Default: 1.
smoother: damping factor *	[double] Smoother damping factor. Default: 0.67.
smoother: pre or post *	[string] It can assume on the following values: <b>pre</b> , <b>post</b> , <b>both</b> . Default: <b>both</b> .
smoother: type *	[string] Type of the smoother. See Table 4. Default: <b>symmetric Gauss-Seidel</b> .
smoother: Aztec options *	[int*] Pointer to AZTEC's options vector (only for Aztec smoother) .

smoother: Aztec params *	[double*] Pointer to AZTEC's params vector (only for Aztec smoother) .
smoother: Aztec as solver *	[bool] If true, smoother: sweeps iterations of AZTEC solvers will be used as smoothers. If false, only the AZTEC's preconditioner function will be used as smoother (only for Aztec smoother). Default: false.
smoother: MLS polynomial order *	[int] Polynomial order for MLS smoothers. Default: 3.
smoother: MLS alpha *	[double] Alpha value for MLS smoothers. Default: 30.0.
smoother: user-defined function	[int (*)(ML_Smoother *, int inLength, double *invec, int outLength, double *outvec)] Pointer to user-defined smoothing function. Default: NULL.
smoother: user-defined name	[string] Label for user-defined smoother. Default: "User-defined".
smoother: Hiptmair sweeps *	[int] Number of Hiptmair iterations to perform. Default: 1.
smoother: Hiptmair efficient symmetric	[bool] Reduce the preconditioner computational work while maintaining symmetry by doing edge-node relaxation on the first leg of the V-cycle, and node-edge relaxation on the second leg. Default: true.
subsmoother: type *	[string] Smoother to be used as a subsmoother at each step of Hiptmair smoothing. It can assume on the following values: MLS, symmetric Gauss-Seidel. Default: MLS.
subsmoother: MLS polynomial order *	[int] Polynomial order for Hiptmair MLS sub-smoother. Default: 3.
subsmoother: MLS alpha *	[int] Polynomial coefficient for Hiptmair MLS sub-smoother. Default: 3.
subsmoother: SGS damping factor *	[double] Damping factor for Hiptmair sub-smoother symmetric Gauss-Seidel. Automatically calculated in the range (0, 2).

subsmoother: SGS edge sweeps *	[int] Number of iterations of Hiptmair sub-smoother SGS to perform on edge problem. Default: 1.
subsmoother: SGS node sweeps *	[int] Number of iterations of Hiptmair sub-smoother SGS to perform on nodal projection problem. Default: 1.

#### 6.4.5 Coarsest Grid Parameters

coarse: max size	[int] Maximum dimension of the coarse grid. <b>ML</b> will not coarsen further if current level's size is less than this value. Default: 128.
coarse: type	[string] Coarse solver. See Table 5. Default: <b>Amesos_KLU</b> .
coarse: sweeps	[int] (Only for Jacobi, Gauss-Seidel, symmetric Gauss-Seidel, and Hiptmair.) Number of sweeps in the coarse solver. Default: 1.
coarse: user-defined function	[int (*)(ML.Smoother *, int inLength, double *invec, int outLength, double *outvec)] Pointer to user-defined smoothing function. Default: <b>NULL</b> .
coarse: user-defined name	[string] Label for user-defined smoother. Default: "User-defined".
coarse: damping factor	[double] (Only for Jacobi and Gauss-Seidel.) Damping factor in the coarse solver. Default: 0.67.
coarse: max processes	[int] Maximum number of processes to be used in the coarse grid solution (only for <b>Amesos-Superludist</b> , <b>Amesos-MUMPS</b> ). If -1, <b>AMESOS</b> will decide the optimal number of processors to be used. Default: -1.

#### 6.4.6 Load-balancing Options

repartition: enable	[int] Enable/disable repartitioning. Default: 0 (off).
---------------------	--



repartition: partitioner	[string] Partitioning package to use. Can assume the following values: Zoltan, ParMETIS. Default: Zoltan.
repartition: max min ratio	[double] Specifies desired ratio of $g_{max}/g_{min}$ , where $g_{max}$ is the maximum number rows over all processors, and $g_{min}$ is the minimum number of rows over all processors. If the actual ratio is larger, then repartitioning occurs. Default: 1.1.
repartition: min per proc	[int] Specifies desired minimum number of rows per processor. If the actual number is greater, then repartitioning occurs. Default: 20.
repartition: node max min ratio	[double] Maxwell-specific option for auxiliary nodal hierarchy. Specifies desired ratio of $g_{max}/g_{min}$ , where $g_{max}$ is the maximum number rows over all processors, and $g_{min}$ is the minimum number of rows over all processors. If the actual ratio is larger, then repartitioning occurs. Default: 1.1.
repartition: node min per proc	[int] Maxwell-specific option for auxiliary nodal hierarchy. Specifies desired minimum number of rows per processor. If the actual number is greater, then repartitioning occurs. Default: 20.
repartition: Zoltan dimensions	[int] Dimension of the problem. Can assume the following values: {2, 3}.

#### 6.4.7 Visualization

viz: enable	[bool] Enable/disable visualization. Default: false.
viz: output format	[string] The format of the visualization files. Can assume the following values: vtk, xyz, openx. Default: vtk.
viz: print starting solution	[bool] Enable/disable printing of initial starting solution.
viz: equation to plot	[bool] In the case of systems of PDEs, the equation number to print. Default: all equations.

#### 6.4.8 Miscellaneous

<code>x-coordinates</code>	<code>[double*]</code> Pointer to array of $x$ -coordinates of mesh nodes (edges, in the case of Maxwell).
<code>y-coordinates</code>	<code>[double*]</code> Pointer to array of $y$ -coordinates of mesh nodes (edges, in the case of Maxwell).
<code>z-coordinates</code>	<code>[double*]</code> Pointer to array of $z$ -coordinates of mesh nodes (edges, in the case of Maxwell).
<code>node: x-coordinates</code>	<code>[double*]</code> Pointer to array of $x$ -coordinates of mesh nodes. (Maxwell only)
<code>node: y-coordinates</code>	<code>[double*]</code> Pointer to array of $y$ -coordinates of mesh nodes. (Maxwell only)
<code>node: z-coordinates</code>	<code>[double*]</code> Pointer to array of $z$ -coordinates of mesh nodes. (Maxwell only)

#### 6.4.9 Null Space Detection

If used in conjunction with ANASAZI, **ML** can compute an approximate null space of the linear operator to precondition. This can be done using the following options:

<code>null space: type</code>	<code>[string]</code> If <code>default vectors</code> , the default null space are used (no computation required). If <code>pre-computed</code> , a pointer to the already-computed null is obtained from option <code>null space: vectors</code> . If <code>enriched</code> , <b>ML</b> will compute, using ANASAZI, an approximation of the null space of the operator. Default: <code>default vectors</code> .
<code>null space: vectors</code>	<code>[double*]</code> . Pointer to user-supplied null space vectors.
<code>null space: dimension</code>	<code>[int]</code> . Number of user-supplied null space vectors.
<code>null space: vectors to compute</code>	<code>[int]</code> . If <code>null space: type</code> is set to <code>enriched</code> , this option indicates the number of eigenvectors to compute. Default: 1.
<code>null space: add default vectors</code>	<code>[bool]</code> . If <code>true</code> , the default null space (one constant vector for each unknown) will be added to the computed null space. Default: <code>true</code> .

<code>eigen-analysis: tolerance</code>	[double] Tolerance for ANASAZI. Default: 0.1.
<code>eigen-analysis: restart</code>	[int] Number of restarts for ANASAZI. Default: 2.

#### 6.4.10 Aggregation Strategies

In some case, it is preferable to generate the aggregates using a matrix defined as follows:

$$L_{i,j} = (\mathbf{x}_i - \mathbf{x}_j)^{-2}, \quad i \neq j, \quad L_{i,i} = -\sum_{i \neq j} L_{i,j}, \quad (2)$$

(where  $\mathbf{x}_i$  represents the coordinates of node  $i$ ) in conjunction with a given nonzero dropping threshold. This may happen, for instance, when working with bilinear finite elements on stretched grids. **ML** can be instructed to generate the aggregates using matrix (2), then build the preconditioner using the actual linear system matrix, as done in the following code fragment:

```
double* x_coord;
double* y_coord; // set to 0 for 1D problems
double* z_coord; // set to 0 for 2D problems
// here we define the nodal coordinates...

MLList.set("x-coordinates", x_coord );
MLList.set("y-coordinates", y_coord );
MLList.set("z-coordinates", z_coord );
```

The double vectors `y_coord` and/or `z_coord` can be set to 0. `x_coord` (and `y_coord` if 2D, and `z_coord` if 3D) must be allocated to contain the coordinates for all nodes assigned to the calling processor, plus that of the ghost nodes. For systems of PDE equations, the coordinates must refer to the block nodes (that is, x-coordinate of local row `i` will be reported in `x_coord[i/NumPDEEqns]`).

The following option can be used:

<code>aggregation: aux: enable</code>	[bool]. Enable/disable the use of matrix (2). Default: <code>false</code> .
---------------------------------------	--

## 6.5 Default Parameter Settings for Common Problem Types

The `MultiLevelPreconditioner` class provides default values for four different preconditioner types:

1. Linear elasticity
2. Classical 2-level domain decomposition for the advection diffusion operator
3. 3-level algebraic domain decomposition for the advection diffusion operator

#### 4. Eddy current formulation of Maxwell's equations

Default values are listed in Table 6. In the table, **SA** refers to “classical” smoothed aggregation (with small aggregates and relative large number of levels), **DD** and **DD-ML** to domain decomposition methods (whose coarse matrix is defined using aggressive coarsening and limited number of levels). **Maxwell** refers to the solution of Maxwell's equations.

Default values for the parameter list can be set by `ML_Epetra::SetDefaults()`. The user can easily put the desired default values in a given parameter list as follows:

```
Teuchos::ParameterList MList;  
ML_Epetra::SetDefaults(ProblemType, MList);
```

or as

```
Teuchos::ParameterList MList;  
ML_Epetra::SetDefaults(ProblemType, MList);
```

For **DD** and **DD-ML**, the default smoother is **Aztec**, with an incomplete factorization **ILUT**, and minimal overlap. Memory for the two **AZTEC** vectors is allocated using **new**, and the user is responsible to free this memory, for instance as follows:

```
int* options;  
options = MList.get("smoother: Aztec options", options);  
double* params;  
params = MList.get("smoother: Aztec params", params);  
.  
.  
.  
// Make sure solve is completed before deleting options & params!!  
delete [] options;  
delete [] params;
```

The rational behind this is that the parameter list stores a *pointer* to those vectors, not the content itself. (As a general rule, the vectors stored in the parameter list should not be prematurely destroyed or permitted to go out of scope.)

## 6.6 Analyzing the ML preconditioner

A successful multilevel preconditioner require the careful choice of a large variety of parameters, for instance each level's smoother, the aggregation schemes, or the coarse solver. Often, for non-standard problems, there is no theory to support these choices. Also, sometimes it is difficult to understand which component of the multilevel cycle is not properly working. To help to set the multilevel components, **ML** offers a set of tools, to (empirically) analyze several components of the multilevel cycles, and the finest-level matrix.

Two examples are included in the **ML** distribution:

- File `examples/Visualization/ml_viz.cpp` shows how to visualize the effect of the **ML** cycle and each level's smoother on a random vector;
- File `examples/Advanced/ml_analyze.cpp` shows how to get some quantitative information about each level's matrix, and multilevel preconditioner.

Option Name	Type	SA	DD	DD-ML	maxwell
max levels	int	10	2	3	10
output	int	8	8	8	10
increasing or decreasing	string	increasing	increasing	increasing	decreasing
PDE equations	int	1	1	1	1
print unused	int	0	0	0	0
aggregation: type	string	Uncoupled-MIS	METIS	METIS	Uncoupled-MIS
aggregation: local aggregates	int	–	1	–	–
aggregation: nodes per aggregate	int	–	–	512	–
aggregation: damping factor	double	4/3	4/3	4/3	4/3
eigen-analysis: type	string	Anorm	Anorm	Anorm	Anorm
aggregation: threshold	double	0.0	0.0	0.0	0.0
aggregation: next-level aggregates	int	–	–	128	–
per process					
smoother: sweeps	int	2	2	2	1
smoother: damping factor	double	0.67	–	–	1.0
smoother: pre or post	string	both	both	both	both
smoother: type	string	symmetric Gauss-Seidel	Aztec	Aztec	Hiptmair/MLS
smoother: Aztec as solver	bool	–	false	false	false
smoother: MLS polynomial order	int	–	–	–	3
smoother: MLS alpha	double	–	–	–	27.0
coarse: type	string	Amesos_KLU	Amesos_KLU	Amesos_KLU	Amesos_KLU
coarse: max size	int	16	128	128	75
coarse: sweeps	int	1	1	1	1
coarse: damping factor	double	1.0	1.0	1.0	1.0
coarse: max processes	int	16	16	16	–

Table 6: Default values for ML\_Epetra::MultiLevelPreconditioner for the 4 currently supported problem types SA, DD, DD-ML, Maxwell. “\_” means not set.

In the following subsections, we suppose that a `MultiLevelPreconditioner` object has already be defined, and the preconditioner computed.

### 6.6.1 Cheap Analysis of All Level Matrices

Method `AnalyzeMatrixCheap()` will report on standard output general information about each level's matrix. An example of output is as reported below. (Here, we report only the part of output related to the finest level.)

```
*** Analysis of ML_Operator 'A matrix level 0' ***

Number of global rows           = 256
Number of equations             = 1
Number of stored elements       = 1216
Number of nonzero elements      = 1216
Minimum number of nonzero elements/row = 3
Maximum number of nonzero elements/row = 5
Average number of nonzero elements/rows = 4.750000
Nonzero elements in strict lower part = 480
Nonzero elements in strict upper part = 480
Max |i-j|, a(i,j) != 0         = 16
Number of diagonally dominant rows = 86 (= 33.59%)
Number of weakly diagonally dominant rows = 67 (= 26.17%)
Number of Dirichlet rows       = 0 (= 0.00%)
||A||_F                        = 244.066240
Min_{i,j} ( a(i,j) )          = -14.950987
Max_{i,j} ( a(i,j) )          = 15.208792
Min_{i,j} ( abs(a(i,j)) )      = 0.002890
Max_{i,j} ( abs(a(i,j)) )      = 15.208792
Min_i ( abs(a(i,i)) )          = 2.004640
Max_i ( abs(a(i,i)) )          = 15.208792
Min_i ( \sum_{j!=i} abs(a(i,j)) ) = 2.004640
Max_i ( \sum_{j!=i} abs(a(i,j)) ) = 15.205902
max eig(A) (using power method) = 27.645954
max eig(D^{-1}A) (using power method) = 1.878674
```

Total time for analysis = 3.147979e-03 (s)

This analysis is “cheap” in the sense that it involves only element-by-element comparison, plus the computation of the largest-magnitude eigenvalue (which requires some matrix-vector products). `AnalyzeMatrixCheap()` can be used for both serial and parallel runs.

### 6.6.2 Analyze the Effect of Smoothers

For each level, method `AnalyzeSmoothers()` computes the eigenvectors of the matrix (say,  $A$ ). Then, for each eigenvector (say,  $v$ ) of  $A$ , the smoother is applied to the solution of the homogeneous system

$$Ae = 0$$

with starting solution  $e_0 = v$ . The code reports on file the real and imaginary values of the eigenvalue corresponding to eigenvector  $v$ , and the  $\|e\|/\|e_0\|$ .

The syntax is `AnalyzeSmoother(NumPre, NumPost)`. `NumPre` is the number of pre-smoother applications, and `NumPost` the number of post-smoother applications. This method reports on the standard output the following information:

```
Solving Ae = 0, with a random initial guess
- number of pre-smoother cycle(s) = 5
- number of post-smoother cycle(s) = 5
- all reported data are scaled with their values
  before the application of the solver
  (0 == perfect solution, 1 == no effect)
- SF is the smoothness factor
```

Solver	Linf	L2	SF
Presmoother (level 0, eq 0)	0.827193	0.804528	0.313705
Postsmoother (level 0, eq 0)	0.822015	0.810521	0.342827
Presmoother (level 1, eq 0)	0.972593	0.908874	2.51318
Postsmoother (level 1, eq 0)	0.982529	0.922668	2.53639

### 6.6.3 Analyze the effect of the ML cycle on a random vector

Method `AnalyzeCycle(NumCycles)`, where `NumCycles` is the number of multilevel cycles to apply, applies the already computed **ML** preconditioner to a random vector, and reports on standard output the following information:

```
Solving Ae = 0, with a random initial guess
using 5 ML cycle(s).
- (eq 0) scaled Linf norm after application(s) = 0.0224609
- (eq 0) scaled L2 norm after application(s)   = 0.000249379
- (eq 0) scaled smoothness factor              = 10.6517
```

### 6.6.4 Test different smoothers

The `MultiLevelPreconditioner` class offers a very easy way to test the effect of a variety of smoothers on the problem at hand. Once the preconditioning object has been created, a call to `TestSmoother()` performs the following operations:

1. Creates a new linear system, whose matrix is the one used to construct the `MultiLevelPreconditioner` object;
2. Defines a random solution, and the corresponding right-hand side;
3. Defines a zero starting vector for the Krylov solver;
4. Creates a new preconditioning object, with the same options as in the current preconditioner, except for the choice of the smoothers;
5. Solve the linear system with the newly created preconditioner;

6. Reports in a table the iterations to converge and the corresponding CPU time.

The following options, to be set *before* calling `ComputePreconditioner()`, can be used to tune the test session:

<code>test: max iters</code>	[int] Maximum number of iterations for the Krylov solver. Default: 500.
<code>test: tolerance</code>	[double] Tolerance for the Krylov solver. Default: 1e-5.
<code>test: Jacobi</code>	[bool] Enable/disable test with Jacobi smoother. Default: true.
<code>test: Gauss-Seidel</code>	[bool] Enable/disable test with Gauss-Seidel smoother. Default: true.
<code>test: symmetric Gauss-Seidel</code>	[bool] Enable/disable test with symmetric Gauss-Seidel smoother. Default: true.
<code>test: block Gauss-Seidel</code>	[bool] Enable/disable test with block Gauss-Seidel smoother. Default: true.
<code>test: Aztec</code>	[bool] Enable/disable test with AZTECOO smoother. Default: true.
<code>test: Aztec as solver</code>	[bool] Enable/disable test with AZTECOO as a solver smoother. Default: true.
<code>test: ParaSails</code>	[bool] Enable/disable test with PARASAILSSmoothing. Default: true.
<code>test: IFPACK</code>	[bool] Enable/disable test with IFPACKsmoothing. Default: true.

An example of output is reported below. Note that some smoothers (PARASAILS, IFPACK) will be tested only if **ML** has been properly configured. Note also that `TestSmoothers()` requires **ML** to be configured with option `--enable-aztecoo`.

---

```
*** *****
*** Analysis of ML parameters (smoothers) ***
*** *****
```

```
*** maximum iterations = 500
*** tolerance          = 1e-05
```

```
*** All options as in the input parameter list, except that
*** all levels have the same smoother
```



\*\*\* M: maximum iterations exceeded without convergence  
 \*\*\* N: normal exit status (convergence achieved)  
 \*\*\* B: breakdown occurred  
 \*\*\* I: matrix is ill-conditioned  
 \*\*\* L: numerical loss of precision occurred

count smoother type.....its.....||r||/||r\_0||..time (s).....

- Jacobi

#0....n=5, omega=2.50e-01.....12.....2.97314e-06....0.0839319.....N  
 #1....n=5, omega=5.00e-01.....12.....6.21844e-06....0.0820519.....N  
 #2....n=5, omega=7.50e-01.....12.....7.52614e-06....0.082267.....N  
 #3....n=5, omega=1.00e+00.....12.....3.80406e-06....0.082956.....N  
 #4....n=5, omega=1.25e+00.....12.....2.15858e-06....0.0824361.....N

- Gauss-Seidel

#5....n=5, omega=2.50e-01.....7.....2.20736e-06....0.0857691.....N  
 #6....n=5, omega=5.00e-01.....7.....1.91864e-06....0.0789189.....N  
 #7....n=5, omega=7.50e-01.....6.....8.40948e-06....0.076307.....N  
 #8....n=5, omega=1.00e+00.....7.....1.36415e-06....0.0792729.....N  
 #9....n=5, omega=1.25e+00.....7.....1.4833e-06....0.0790809.....N

- Gauss-Seidel (sym)

#10....n=5, omega=2.50e-01.....4.....2.32835e-06....0.149586.....N  
 #11....n=5, omega=5.00e-01.....4.....2.68576e-06....0.092068.....N  
 #12....n=5, omega=7.50e-01.....4.....8.51966e-07....0.0793679.....N  
 #13....n=5, omega=1.00e+00.....4.....1.34439e-06....0.0787291.....N  
 #14....n=5, omega=1.25e+00.....4.....5.09185e-06....0.0790669.....N

- Gauss-Seidel (block)

#15....n=5, omega=2.50e-01.....6.....6.56673e-06....0.0920131.....N  
 #16....n=5, omega=5.00e-01.....7.....1.77309e-06....0.0881529.....N  
 #17....n=5, omega=7.50e-01.....6.....8.53488e-06....0.0846661.....N  
 #18....n=5, omega=1.00e+00.....6.....2.66381e-06....0.0839909.....N  
 #19....n=5, omega=1.25e+00.....6.....4.87356e-06....0.083786.....N

- Aztec preconditioner

#20....ILU(fill=0).....7.....4.93736e-06....0.0712331.....N  
 #21....ILU(fill=1).....6.....3.54992e-06....0.0647091.....N  
 #22....ILU(fill=2).....7.....1.4724e-06....0.0678911.....N

- Aztec as solver

#23....iterations=1.....7.....1.94081e-06....0.140772.....N  
 #24....iterations=3.....4.....8.90029e-08....0.0687031.....N  
 #25....iterations=5.....3.....1.00778e-07....0.069193.....N

```

- ParaSails
#26....default.....20.....6.60045e-06....0.214094.....N

*** The best iteration count was obtain in test 25
*** The best CPU-time was obtain in test 21

*** Total time = 2.43798(s)
-----

```

## 6.7 Visualization Capabilities

### 6.7.1 Visualizing the Aggregates

**ML** offers the possibility to visualize the aggregates for all levels. Aggregates generated by **Uncoupled** and **METIS** aggregation schemes can be visualized for serial and parallel runs, while aggregates generated using **MIS** and **ParMETIS** can be visualized only for serial runs.

Data can be stored in the “xyz” format, readable by packages such as **XD3D** [?]. For each aggregate, the file contains a line of type

```
x-coord y-coord <z-coord> aggregate-number
```

(**z-coord** is present only for 3D computations.) **XD3D** can be used to visualize files in this format, but only for 2D problems. Results are reported in Figure 2.

Data can also be stored in the “vtk legacy” format, readable by packages such as **PARAVIEW** [?]. **PARAVIEW** allows for 2D and 3D point clouds visualization. It is also possible to visualize sections of 3D data via cutting planes. In order to use this format, fine level coordinates must be supplied. Coarse level coordinates are derived from the fine level coordinates via the multigrid restriction operators. Data is stored as follows. The first section begins with the header

```
POINTS XX float
```

where **XX** is the total number of points. For each point, the coordinates are given as

```
x-coord y-coord <z-coord>
```

Next, connectivity information is given in a pair-wise fashion. For example, if the  $(i, j)$  entry of matrix  $A$  is nonzero, then there is a line in the data file of the form

```
2 i j
```

where the first entry specifies the number of points that follow. (This first entry will always be 2 because the connectivity is pair-wise.) The next data section specifies the aggregate to which each point belongs. It begins with the header

```
POINT_DATA XX
```

```
SCALARS aggregate_number int 1
```

```
LOOKUP_TABLE default
```

The section then has a line for each point of the form

```
aggregate_number
```

The final section specifies the solution (residual) at each point. The section begins with the header

```
SCALARS solution float 1  
LOOKUP_TABLE default
```

and then has a line for each point of the form

```
solution_value
```

With the exception of connectivity, all information is given in the same order. Hence, all information for the  $j$ th unknown can be found at the  $j$ th line of each section.

### 6.7.2 Visualizing the effect of the ML Preconditioner and Smoothers

In some cases, it may be useful to visualize the effect of the **ML** preconditioner, or of each level's smoother, on a random vector (whose components are contained in the interval  $[0.5, 1]$ ), to understand if there are zones or directions that are not affected by the current multilevel preconditioner. This can be easily done with the following code fragment:

```
double* x_coord;  
double* y_coord;  
// here we define the nodal coordinates...  
  
MLList.set("viz: enable", true);  
//you can also specify "xyz" on the line below  
MLList.set("viz: output format", "vtk");  
MLList.set("viz: x-coordinates", x_coord);  
MLList.set("viz: y-coordinates", y_coord);  
// by default the starting solution is not printed  
MLList.set("viz: print starting solution", true);  
  
// create the preconditioner  
ML_Epetra::MultiLevelPreconditioner * MLPrec =  
    new ML_Epetra::MultiLevelPreconditioner(*A, MLList, true);  
  
// visualize the effect of the ML preconditioner on a  
// random vector. We ask for 10 multilevel cycles  
MLPrec->VisualizeCycle(10);  
  
// visualize the effect of each level's smoother on a  
// random vector. We ask for 5 steps of presmoothers,  
// and 1 step of postsmoother  
MLPrec->VisualizeSmoothers(5,1);
```

(File `ml_viz.cpp` contains the compilable code.) We note that the parameters must be set *before* calling `ComputePreconditioner()`. See also Section 6.4.10 for the requirements on the coordinates vectors. Results will be written in the following files:

- `before-presmoothing-eqX-levelY.vtk` contains the random vector before the application of the presmoothing, for equation X at level Y;
- `after-presmoothing-eqX-levelY.vtk` contains the random vector after the application of the presmoothing, for equation X at level Y;
- `before-postsmoothing-eqX-levelY.vtk` contains the random vector before the application of the postsmoothing, for equation X at level Y;
- `after-postsmoothing-eqX-levelY.vtk` contains the random vector after the application of the postsmoothing, for equation X at level Y;
- `before-cycle-eqX-levelY.vtk` contains the random vector before the application of the MLcycle, for equation X at the finest level;
- `after-cycle-eqX-levelY.vtk` contains the random vector after the application of the ML cycle, for equation X at finest level.

## 6.8 Print the Computational Stencil for a 2D Cartesian Grid

Method `PrintStencil2D()` can be used to print out the computational stencil for problems defined on 2D Cartesian grids, if the nodes numbering follows the x-axis. The following fragment of code shows the use of this method:

```
int Nx = 16; // nodes along the x-axis
int Ny = 32; // nodes along the y-axis
int NodeID = -1; // print the stencil for this node
int EquationID = 0; // equation 0, useful for vector problems

// MLPrec is a pointer to an already created
// ML_Epetra::MultiLevelPreconditioner
MLPrec->PrintStencil2D(Nx,Ny,NodeID, EquationID);

// also valid in this case
MLPrec->PrintStencil2D(Nx,Ny);
```

If `NodeID == -1`, the code considers a node in the center of the computational domain. The result can be something as follows:

2D computational stencil for equation 0 at node 136 (grid is 16 x 16)

0	-1	0
-1	4	-1
0	-1	0

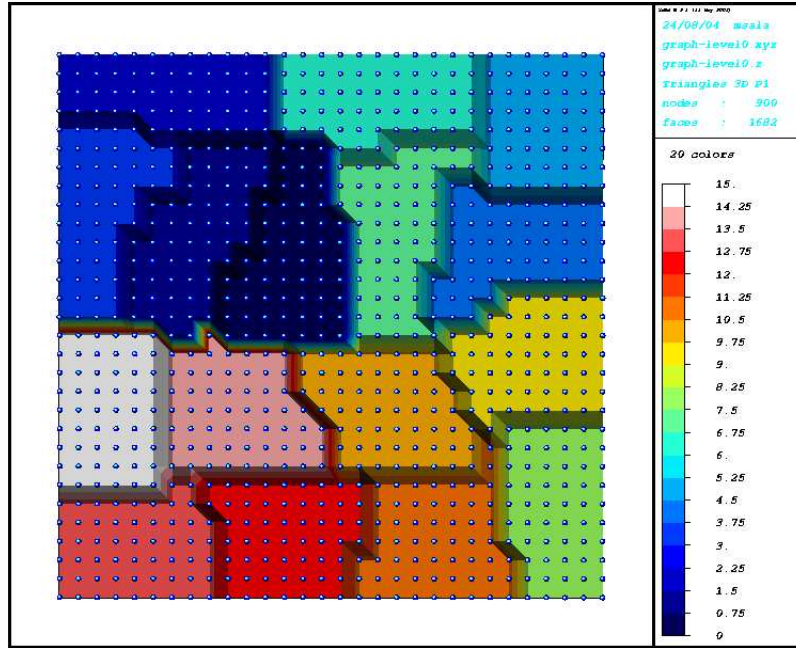


Figure 2: Decomposition into aggregates. Uniform colors between blue spheres represent the interior of aggregates. Visualization was done with XD3D.

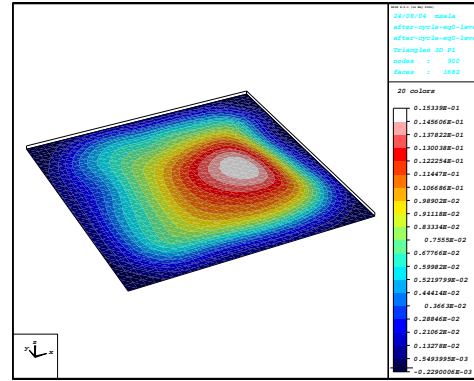
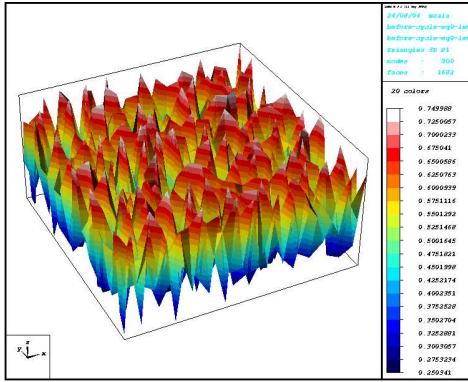


Figure 3: Starting solution (left) and solution after 10 **ML** cycles (right). Visualization was done using XD3D.

## 7 Using the Maxwell Solver in ML

This section gives a brief overview of how to ML’s AMG preconditioner for the eddy current formulation of Maxwell’s equations. The solver is intended only for equations in the time domain (real-valued), not the frequency domain (complex-valued).

### 7.1 Background

The eddy current formulation of Maxwell’s equations can be written as

$$\nabla \times \nabla \times \vec{E} + \sigma \vec{E} = \vec{f}, \quad (3)$$

where  $\vec{E}$  is the unknown electric field to be computed,  $\sigma$  is the spatially-varying electrical conductivity, and  $\vec{f}$  is the known right-hand side. Neumann, Dirichlet, and/or periodic boundary conditions are supported.

Although we assume that (3) is discretized with first-order edge elements, it is possible to use this preconditioner for higher-order discretizations, as long as the user provides the sub-matrix that corresponds to just the first-order discretization. For more theoretical background and algorithm descriptions, please see [14, 2].

### 7.2 Notational conventions

For the remainder of the discussion,  $K^{(e)}$  denotes the matrix corresponding to the first-order element discretization of (3).  $K^{(e)}$  can be written as

$$K^{(e)} = S + M,$$

where  $S$  is the stiffness matrix corresponding to the  $\nabla \times \nabla \times$  term in (3), and  $M$  is the mass matrix corresponding to the  $\sigma \vec{E}$  term in (3). The null-space of  $S$  is given by the discrete gradient matrix,  $T$ .  $T$  corresponds to the null space of the  $\nabla \times \nabla \times$  term in (3).  $K^{(n)}$  is an auxiliary nodal matrix that is described in §7.3.1.

### 7.3 Operators that the user must supply

The user must provide three matrices:

1. the first-order edge element matrix  $K^{(e)}$ .
2. the auxiliary nodal matrix  $K^{(n)}$ .
3. the discrete gradient matrix  $T$ .

#### 7.3.1 Description of the auxiliary nodal matrix $K^{(n)}$

$K^{(n)}$  is a square,  $N_n \times N_n$  matrix, where  $N_n$  is the number of nodes in the mesh. There are two ways to construct  $K^{(n)}$ . The first method is to discretize the PDE

$$\int_{\Omega} \sigma u \cdot v + \int_{\Omega} \nabla u \cdot \nabla v, \quad (4)$$

using nodal linear finite elements on the same mesh as (3). The second method assumes that you already have  $T$  available.  $K^{(n)}$  can be formed via the triple-matrix product

$$T^T K^{(e)} T = T^T M T,$$

where we have used the fact that  $ST = 0$ .

### 7.3.2 Description of the discrete gradient matrix $T$

$T$  is a rectangular,  $N_e \times N_n$  matrix, where  $N_e$  is the number of edges in the mesh (rows in  $K^{(e)}$ ) and  $N_n$  is the number of nodes in the mesh (rows in  $K^{(n)}$ ). Each row of  $T$  has at most two entries, a +1 and/or a -1. There are two ways to construct  $T$ . In the first method, it's assumed that the mesh topology is available.  $T$  can be viewed as a node-edge incidence matrix of the mesh as a directed graph. As such, each row of  $T$  corresponds to an edge, and the +1/-1 entries in the row are the head/tail of the edge, respectively. Hence,  $T$  can be built one row at a time by visiting each edge and inserting +1/-1 in the appropriate columns. The second method assumes that  $K^{(n)}$  is already constructed. Each off-diagonal nonzero entry,  $(i, j)$ , in the upper triangular portion of  $K^{(n)}$  corresponds to a row in  $T$  containing a 1 and a -1 in columns  $i$  and  $j$ .

## 7.4 Smoother options

The default smoother for Maxwell's equations is a two-stage smoother that we call Hiptmair smoothing [13]. This smoother consists of three steps: one step of a smoother  $\mathcal{S}$  on the entire system  $Kx = f$ ; one step of a smoother  $\mathcal{S}$  on the projected system  $T^T M T e = T^T r$ , where  $r = f - Kx$ ; and finally one step of a smoother  $\mathcal{S}$  on the entire system with initial guess  $x + e$ .

In serial, the default sub-smoother  $\mathcal{S}$  is symmetric Gauss-Seidel (SGS). In parallel, the default sub-smoother  $\mathcal{S}$  is a degree 3 Chebyshev polynomial. The coefficients of this polynomial are calculated automatically based on the coarsening rate of the multigrid hierarchy. The Chebyshev degree can be changed with the option

`smoother: Hiptmair MLS polynomial order`



## 8 Advanced Usage of ML

Sections 6 and 6.4 have detailed the use of **ML** as a black box preconditioner. In some cases, instead, the user may need to explicitly construct the **ML** hierarchy. This is reported in the following sections.

A brief sample program is given in Figure 4. The function `ML_Create` creates a multilevel

```
ML_Create          (&ml_object, N_grids);

ML_Init_Amatrix    (ml_object, 0,  nlocal, nlocal, (void *) A_data);
ML_Set_Amatrix_Getrow(ml_object, 0,  user_getrow, NULL, nlocal_allcolumns);
ML_Set_Amatrix_Matvec(ml_object, 0,  user_matvec);

N_levels = ML_Gen_MGHierarchy_UsingAggregation(ml_object, 0,
                                                ML_INCREASING, NULL);
ML_Gen_Smoothen_Jacobi(ml_object, ML_ALL_LEVELS, ML_PRESMOOTHER, 1,
                       ML_DEFAULT);
ML_Gen_Solver      (ml_object, ML_MGV, 0, N_levels-1);
ML_Iterate(ml_object, sol, rhs);
ML_Destroy(&ml_object);
```

Figure 4: High level multigrid sample code.

solver object that is used to define the preconditioner. It requires the maximum number of multigrid levels be specified. In almost all cases, `N_grids= 20` is more than adequate. The three ‘Amatrix’ statements are used to define the discretization matrix,  $A$ , that is solved. This is discussed in greater detail in Section 12.1. The multigrid hierarchy is generated via `ML_Gen_MGHierarchy_UsingAggregation`. Controlling the behavior of this function is discussed in Section 10. For now, it is important to understand that this function takes the matrix  $A$  and sets up relevant multigrid operators corresponding to the smoothed aggregation multigrid method [22] [21]. In particular, it generates a graph associated with  $A$ , coarsens this graph, builds functions to transfer vector data between the original graph and the coarsened graph, and then builds an approximation to  $A$  on the coarser graph. Once this second multigrid level is completed, the same operations are repeated to the second level approximation to  $A$  generating a third level. This process continues until the current graph is sufficiently coarse. The function `ML_Gen_Smoothen_Jacobi` indicates that a Jacobi smoother should be used on all levels. Smoothers are discussed further in Section 9. Finally, `ML_Gen_Solver` is invoked when the multigrid preconditioner is fully specified. This function performs any needed initialization and checks for inconsistent options. After `ML_Gen_Solver` completes `ML_Iterate` can be used to solve the problem with an initial guess of `sol` (which will be overwritten with the solution) and a right hand side of `rhs`. At the present time, the external interface to vectors are just arrays. That is, `rhs` and `sol` are simple one-dimensional arrays of the same length as the number of rows in  $A$ . In addition to `ML_Iterate`, the function `ML_Solve_MGV` can be used to perform one multigrid ‘V’ cycle as a preconditioner.

## 9 Multigrid & Smoothing Options

Several options can be set to tune the multigrid behavior. In this section, smoothing and high level multigrid choices are discussed. In the next section, the more specialized topic of the grid transfer operator is considered.

For most applications, smoothing choices are important to the overall performance of the multigrid method. Unfortunately, there is no simple advice as to what smoother will be best and systematic experimentation is often necessary. **ML** offers a variety of standard smoothers. Additionally, user-defined smoothers can be supplied and it is possible to use AZTEC as a smoother. A list of **ML** functions that can be invoked to use built-in smoothers are given below along with a few general comments.

<code>ML_Gen_Smoothers_Jacobi</code>	Typically, not the fastest smoother. Should be used with damping. For Poisson problems, the recommended damping values are $\frac{2}{3}$ (1D), $\frac{4}{5}$ (2D), and $\frac{5}{7}$ (3D). In general, smaller damping numbers are more conservative.
<code>ML_Gen_Smoothers_GaussSeidel</code>	Probably the most popular smoother. Typically, faster than Jacobi and damping is often not necessary nor advantageous.
<code>ML_Gen_Smoothers_SymGaussSeidel</code>	Symmetric version of Gauss Seidel. When using multigrid preconditioned conjugate gradient, the multigrid operator must be symmetrizable. This can be achieved by using a symmetric smoother with the same number of pre and post sweeps on each level.
<code>ML_Gen_Smoothers_BlockGaussSeidel</code>	Block Gauss-Seidel with a fixed block size. Often used for PDE systems where the block size is the number of degrees of freedom (DOFs) per grid point.
<code>ML_Gen_Smoothers_VBlockJacobi</code>	Variable block Jacobi smoother. This allows users to specify unknowns to be grouped into different blocks when doing block Jacobi.
<code>ML_Gen_Smoothers_VBlockSymGaussSeidel</code>	Symmetric variable block Gauss-Seidel smoothing. This allows users to specify unknowns to be grouped into different blocks when doing symmetric block Gauss-Seidel.

It should be noted that the parallel Gauss-Seidel smoothers are not true Gauss-Seidel. In particular, each processor does a Gauss-Seidel iteration using off-processor information from the previous iteration.

AZTEC user's [19] can invoke `ML_Gen_SmoothersAztec` to use either AZTEC solvers or AZTEC preconditioners as smoothers on any grid level. Thus, for example, it is possible to use preconditioned conjugate-gradient (where the preconditioner might be an incomplete

Cholesky factorization) as a smoother within the multigrid method. Using Krylov smoothers as a preconditioner could potentially be more robust than using the simpler schemes provided directly by **ML**. However, one must be careful when multigrid is a preconditioner to an outer Krylov iteration. Embedding an inner Krylov method within a preconditioner to an outer Krylov method may not converge due to the fact that the preconditioner can no longer be represented by a simple matrix. Finally, it is possible to pass user-defined smoothing functions into **ML** via `ML_Set_Smoother`. The signature of the user defined smoother function is

```
int user_smoothing(ML_Smoother *smoother, int x_length, double x[],
                  int rhs_length, double rhs[])
```

where `smoother` is an internal **ML** object, `x` is a vector (of length `x_length`) that corresponds to the initial guess on input and is the improved solution estimate on output, and `rhs` is the right hand side vector of length `rhs_length`. The function `ML_Get_MySmootherData(smoother)` can be used to get a pointer back to the user's data (i.e. the data pointer given with the `ML_Set_Smoother` invocation). A simple (and suboptimal) damped Jacobi smoother for the finest grid of our example is given below:

```
int user_smoothing(ML_Smoother *smoother, int x_length, double x[], int rhs_length, double rhs[])
{
    int i;
    double ap[5], omega = .5; /* temp vector and damping factor */

    Poisson_matvec(ML_Get_MySmootherData(smoother), x_length, x, rhs_length, ap);
    for (i = 0; i < x_length; i++) x[i] = x[i] + omega*(rhs[i] - ap[i])/2.;

    return 0;
}
```

A more complete smoothing example that operates on all multigrid levels is given in the file `mlguide.c`. This routine uses the functions `ML_Operator_Apply`, `ML_Operator_Get_Diag`, and `ML_Get_Amatrix` to access coarse grid matrices constructed during the algebraic multigrid process. By writing these user-defined smoothers, it is possible to tailor smoothers to a particular application or to use methods provided by other packages. In fact, the AZTEC methods within **ML** have been implemented by writing wrappers to existing AZTEC functions and passing them into **ML** via `ML_Set_Smoother`.

At the present time there are only a few supported general parameters that may be altered by users. However, we expect that this list will grow in the future. When using `ML_Iterate`, the convergence tolerance (`ML_Set_Tolerance`) and the frequency with which residual information is output (`ML_Set_ResidualOutputFrequency`) can both be set. Additionally, the level of diagnostic output from either `ML_Iterate` or `ML_Solve_MGV` can be set via `ML_Set_OutputLevel`. The maximum number of multigrid levels can be set via `ML_Create` or `ML_Set_MaxLevels`. Otherwise, **ML** continues coarsening until the coarsest grid is less than or equal to a specified size (by default 10 degrees of freedom). This size can be set via `ML_Aggregate_Set_MaxCoarseSize`.

## 10 Smoothed Aggregation Options

When performing smooth aggregation, the matrix graph is first coarsened (actually vertices are aggregated together) and then a grid transfer operator is constructed. A number of parameters can be altered to change the behavior of these phases.

### 10.1 Aggregation Options

A graph of the matrix is usually constructed by associating a vertex with each equation and adding an edge between two vertices  $i$  and  $j$  if there is a nonzero in the  $(i, j)^{th}$  or  $(j, i)^{th}$  entry. It is this matrix graph whose vertices are aggregated together that effectively determines the next coarser mesh. The above graph generation procedure can be altered in two ways. First, a block matrix graph can be constructed instead of a point matrix graph. In particular, all the degrees of freedom (DOFs) at a grid point can be collapsed into a single vertex of the matrix graph. This situation arises when a PDE system is being solved where each grid point has the same number of DOFs. The resulting block matrix graph is significantly smaller than the point matrix graph and by aggregating the block matrix graph, all unknowns at a grid point are kept together. This usually results in better convergence rates (and the coarsening is actually less expensive to compute). To indicate the number of DOFs per node, the function `ML_Aggregate_Set_NullSpace` is used. The second way in which the graph matrix can be altered is by ignoring small values. In particular, it is often preferential to ignore weak coupling during coarsening. The error between weakly coupled points is generally hard to smooth and so it is best not to coarsen in this direction. For example, when applying a Gauss-Seidel smoother to a standard discretization of

$$u_{xx} + \epsilon u_{yy} = f$$

(with  $0 \leq \epsilon \leq 10^{-6}$ ), there is almost no coupling in the  $y$  direction. Consequently, simple smoothers like Gauss-Seidel do not effectively smooth the error in this direction. If we apply a standard coarsening algorithm, convergence rates suffer due to this lack of  $y$ -direction smoothing. There are two principal ways to fix this: use a more sophisticated smoother or coarsen the graph only in the  $x$  direction. By ignoring the  $y$ -direction coupling in the matrix graph, the aggregation phase effectively coarsens in only the  $x$ -direction (the direction for which the errors are smooth) yielding significantly better multigrid convergence rates. In general, a drop tolerance,  $tol_d$ , can be set such that an individual matrix entry,  $A(i, j)$  is dropped in the *coarsening phase* if

$$|A(i, j)| \leq tol_d * \sqrt{|A(i, i)A(j, j)|}.$$

This drop tolerance (whose default value is zero) is set by `ML_Aggregate_Set_Threshold`.

There are two different groups of graph coarsening algorithms in **ML**:

- schemes with fixed ratio of coarsening between levels: uncoupled aggregation, coupled aggregation, and MIS aggregation. A description of those three schemes along with some numerical results are given in [20]. As the default, the **Uncoupled-MIS** scheme is used which does uncoupled aggregation on finer grids and switches to the more expensive MIS aggregation on coarser grids;

- schemes with variable ratio of coarsening between levels: METIS and PARMETISaggregation. Those schemes use the graph decomposition algorithms provided by METIS and PARMETIS, to create the aggregates.

Poorly done aggregation can adversely affect the multigrid convergence and the time per iteration. In particular, if the scheme coarsens too rapidly multigrid convergence may suffer. However, if coarsening is too slow, the number of multigrid levels increases and the number of nonzeros per row in the coarse grid discretization matrix may grow rapidly. We refer the reader to the above paper and indicate that users might try experimenting with the different schemes via `ML_Aggregate_Set_CoarsenScheme_Uncoupled`, `ML_Aggregate_Set_CoarsenScheme_Coupled`, `ML_Aggregate_Set_CoarsenScheme_MIS`, `ML_Aggregate_Set_CoarsenScheme_METIS`, and `ML_Aggregate_Set_CoarsenScheme_ParMETIS`.

## 10.2 Interpolation Options

An interpolation operator is built using coarsening information, seed vectors, and a damping factor. We refer the reader to [21] for details on the algorithm and the theory. In this section, we explain a few essential features to help users direct the interpolation process.

Coarsening or aggregation information is first used to create a tentative interpolation operator. This process takes a seed vector or seed vectors and builds a grid transfer operator. The details of this process are not discussed in this document. It is, however, important to understand that only a few seed vectors are needed (often but not always equal to the number of DOFs at each grid point) and that these seed vectors should correspond to components that are difficult to smooth. The tentative interpolation that results from these seed vectors will interpolate the seed vectors perfectly. It does this by ensuring that all seed vectors are in the range of the interpolation operator. This means that each seed vector can be recovered by interpolating the appropriate coarse grid vector. The general idea of smoothed aggregation (actually all multigrid methods) is that errors not eliminated by the smoother must be removed by the coarse grid solution process. If the error after several smoothing iterations was known, it would be possible to pick this error vector as the seed vector. However, since this is not the case, we look at vectors associated with small eigenvalues (or singular values in the nonsymmetric case) of the discretization operator. Errors in the direction of these eigenvectors are typically difficult to smooth as they appear much smaller in the residual ( $r = Ae$  where  $r$  is the residual,  $A$  is discretization matrix, and  $e$  is the error). For most scalar PDEs, a single seed vector is sufficient and so we seek some approximation to the eigenvector associated with the lowest eigenvalue. It is well known that a scalar Poisson operator with Neumann boundary conditions is singular and that the null space is the constant vector. Thus, when applying smoothed aggregation to Poisson operators, it is quite natural to choose the constant vector as the seed vector. In many cases, this constant vector is a good choice as all spatial derivatives within the operator are zero and so it is often associated with small singular values. Within **ML** the default is to choose the number of seed vectors to be equal to the number of DOFs at each node (given via `ML_Aggregate_Set_NullSpace`). Each seed vector corresponds to a constant vector for that DOF component. Specifically, if we have a PDE system with two DOFs per node. Then one seed vector is one at the first DOF and zero at the other DOF throughout the graph. The second seed vector is zero at the first DOF and one at the other DOF throughout

the graph. In some cases, however, information is known as to what components will be difficult for the smoother or what null space is associated with an operator. In elasticity, for example, it is well known that a floating structure has six rigid body modes (three translational vectors and three rotation vectors) that correspond to the null space of the operator. In this case, the logical choice is to take these six vectors as the seed vectors in smoothed aggregation. When this type of information is known, it should be given to **ML** via the command `ML_Aggregate_Set_NullSpace`.

Once the tentative prolongator is created, it is smoothed via a damped Jacobi iteration. The reasons for this smoothing are related to the theory where the interpolation basis functions must have a certain degree of smoothness (see [21]). However, the smoothing stage can be omitted by setting the damping to zero using the function `ML_Aggregate_Set_DampingFactor`. Though theoretically poorer, unsmoothed aggregation can have considerably less set up time and less cost per iteration than smoothed aggregation. When smoothing, **ML** has two ways to determine the Jacobi damping parameter and each require some estimate of the largest eigenvalue of the discretization operator. The current default is to use a few iterations of a conjugate-gradient method to estimate this value. However, if the matrix is nonsymmetric, the infinity norm of the matrix should be used instead via `ML_Aggregate_Set_SpectralNormScheme_Anorm`. There are several other internal parameters that have not been discussed in this document. In the future, it is anticipated that some of these will be made available to users.

## 11 Advanced Usage of ML and Epetra

Class `ML_Epetra::MultiLevelOperator` is defined in a header file, that must be included as

```
#include "ml_MultiLevelOperator.h"
```

Users may also need to include `ml_config.h`, `Epetra_Operator.h`, `Epetra_MultiVector.h`, `Epetra_LinearProblem.h`, `AztecOO.h`. Check the EPETRA and AztecOO documentation for more details.

Let **A** be an `Epetra_RowMatrix` for which we aim to construct a preconditioner, and let `ml_handle` be the structure **ML** requires to store internal data (see Section 8), created with the instruction

```
ML_Create(&ml_handle, N_levels);
```

where `N_levels` is the specified (maximum) number of levels. As already pointed out, **ML** can accept in input very general matrices. Basically, the user has to specify the number of local rows, and provide a function to update the ghost nodes (that is, nodes requires in the matrix-vector product, but assigned to another process). For Epetra matrices, this is done by the following function

```
EpetraMatrix2MLMatrix(ml_handle, 0, &A);
```

and it is important to note that **A** is *not* converted to ML format. Instead, `EpetraMatrix2MLMatrix` defines a suitable `getrow` function (and other minor data structures) that allows **ML** to work with **A**.

Let `agg_object` a `ML_Aggregate` pointer, created using

```
ML_Aggregate_Create(&agg_object);
```

At this point, users have to create the multilevel hierarchy, define the aggregation schemes, the smoothers, the coarse solver, and create the solver. Then, we can finally create the `MLEpetra::MultiLevelOperator` object

```
MLEpetra::MultiLevelOperator Mlop(ml_handle,comm,map,map);
```

(`map` being the `Epetra_Map` used to create the matrix) and set the preconditioning operator of our `AZTECOO` solver,

```
Epetra_LinearProblem Problem(A,&x,&b);
Aztec00 Solver(Problem);
solver.SetPrecOperator(&Mlop);
```

where `x` and `b` are `Epetra_MultiVector`'s defining solution and right-hand side. The linear problem can now be solved as, for instance,

```
Solver.SetAztecOption( AZ_solver, AZ_gmres );
solver.Iterate(Niters, 1e-12);
```

## 12 Using ML without Epetra

### 12.1 Creating a ML matrix: Single Processor

Matrices are created by defining some size information, a matrix-vector product and a `getrow` function (which is used to extract matrix information). We note that `EPETRA` and `AZTEC` users do not need to read this (or the next) section as there are special functions to convert `EPETRA` objects and `AZTEC` matrices to **ML** matrices (see Section 5.2). Further, functions for some common matrix storage formats (CSR & MSR) already exist within **ML** and do not need to be rewritten<sup>13</sup>.

Size information is indicated via `ML_Init_Amatrix`. The third parameter in the Figure 4 invocation indicates that a matrix with `nlocal` rows is being defined. The fourth parameter gives the vector length of vectors that can be multiplied with this matrix. Additionally, a data pointer, `A_data`, is associated with the matrix. This pointer is passed back into the matrix-vector product and `getrow` functions that are supplied by the user. Finally, the number '0' indicates at what level within the multigrid hierarchy the matrix is to be stored. For discussions within this document, this is always '0'. It should be noted that there appears to be some redundant information. In particular, the number of rows and the vector length in `ML_Init_Amatrix` should be the same number as the discretization matrices are square. Cases where these 'apparently' redundant parameters might be set differently are not discussed in this document.

The function `ML_Set_Amatrix_Matvec` associates a matrix-vector product with the discretization matrix. The invocation in Figure 4 indicates that the matrix-vector product function `user_matvec` is associated with the matrix located at level '0' of the multigrid hierarchy. The signature of `user_matvec` is

---

<sup>13</sup>The functions `CSR_matvec`, `CSR_getrows`, `MSR_matvec` and `MSR_getrows` can be used.



```
int user_matvec(ML_Operator *Amat, int in_length, double p[], int out_length,
               double ap[])
```

where `A.mat` is an internal `ML` object, `p` is the vector to apply to the matrix, `in_length` is the length of this vector, and `ap` is the result after multiplying the discretization matrix by the vector `p` and `out_length` is the length of `ap`. The function `ML_Get_MyMatvecData(Amat)` can be used to get a pointer back to the user's data (i.e. the data pointer given with the `ML_Init_Amatrix` invocation).

Finally, `ML_Set_Amatrix_Getrow` associates a getrow function with the discretization matrix. This getrow function returns nonzero information corresponding to specific rows. The invocation in Figure 4 indicates that a user supplied function `user_getrow` is associated with the matrix located at level '0' of the multigrid hierarchy and that this matrix contains `nlocal_allcolumns` columns and that no communication (NULL) is used (discussed in the next section). It again appears that some redundant information is being asked as the number of columns was already given. However, when running in parallel this number will include ghost node information and is usually different from the number of rows. The signature of `user_getrow` is

```
int user_getrow(ML_Operator *Amat, int N_requested_rows, int requested_rows[],
               int allocated_space, int columns[], double values[], int row_lengths[])
```

where `Amat` is an internal `ML` object, `N_requested_rows` is the number of matrix rows for which information is returned, `requested_rows` are the specific rows for which information will be returned, `allocated_space` indicates how much space has been allocated in `columns` and `values` for nonzero information. The function `ML_Get_MyGetrowData(Amat)` can be used to get a pointer back to the user's data (i.e. the data pointer given with the `ML_Init_Amatrix` invocation). On return, the user's function should take each row in order within `requested_rows` and place the column numbers and the values corresponding to nonzeros in the arrays `columns` and `values`. The length of the `i`th requested row should appear in `row_lengths[i]`. If there is not enough allocated space in `columns` or `values`, this routine simply returns a '0', otherwise it returns a '1'.

To clarify, these functions, one concrete example is given corresponding to the matrix:

$$\begin{pmatrix} 2 & -1 & & & \\ -1 & 2 & -1 & & \\ & -1 & 2 & -1 & \\ & & -1 & 2 & -1 \\ & & & -1 & 2 \end{pmatrix}. \quad (5)$$

To implement this matrix, the following functions are defined:

```
int Poisson_getrow(ML_Operator *Amat, int N_requested_rows, int requested_rows[],
                  int allocated_space, int columns[], double values[], int row_lengths[])
{
    int count = 0, i, start, row;

    for (i = 0; i < N_requested_rows; i++) {
        if (allocated_space < count+3) return(0);
        start = count;
```



```

        row = requested_rows[i];
        if ( (row >= 0) || (row <= 4) ) {
            columns[count] = row; values[count++] = 2.;
            if (row != 0) { columns[count] = row-1; values[count++] = -1.; }
            if (row != 4) { columns[count] = row+1; values[count++] = -1.; }
        }
        row_lengths[i] = count - start;
    }
    return(1);
}
}

and

int Poisson_matvec(ML_Operator *Amat, int in_length, double p[], int out_length,
                  double ap[])
{
    int i;

    for (i = 0; i < 5; i++ ) {
        ap[i] = 2*p[i];
        if (i != 0) ap[i] -= p[i-1];
        if (i != 4) ap[i] -= p[i+1];
    }
    return 0;
}

```

Finally, these matrix functions along with size information are associated with the fine grid discretization matrix via

```

ML_Init_Amatrix      (ml_object, 0,  5, 5, NULL);
ML_Set_Amatrix_Getrow(ml_object, 0,  Poisson_getrow, NULL, 5);
ML_Set_Amatrix_Matvec(ml_object, 0,  Poisson_matvec);

```

Notice that in these simple examples `Amat` was not used. In the next section we give a parallel example which makes use of `Amat`. The complete sample program can be found in the file `mlguide.c` within the **ML** code distribution.

## 12.2 Creating a ML matrix: Multiple Processors

Creating matrices in parallel requires a bit more work. In this section local versus global indexing as well as communication are discussed. In the description, we reconsider the previous example (5) partitioned over two processors. The matrix row indices (ranging from 0 to 4) are referred to as global indices and are independent of the number of processors being used. On distributed memory machines, the matrix is subdivided into pieces that are assigned to individual processors. **ML** requires matrices be partitioned by rows (i.e. each row is assigned to a processor which holds the entire data for that row). These matrix pieces are stored on each processor as smaller local matrices. Thus, global indices in the original matrix get mapped to local indices on each processor. In our example, we will assign global rows 0 and 4 to processor 0 and store them locally as rows 1 and 0 respectively. Global columns 0, 1, 3, and 4 are stored locally as columns 1, 3, 2, and 0. This induces the local matrix

$$\begin{pmatrix} 2 & & -1 & \\ & 2 & & -1 \end{pmatrix}.$$

Likewise, processor 1 is assigned global rows 1, 2, and 3 which are stored locally as rows 0, 1, and 2 respectively. Global columns 0 - 4 are stored locally as columns 3, 0, 1, 2, and 4 inducing the local matrix

$$\begin{pmatrix} 2 & -1 & & & -1 \\ -1 & 2 & -1 & & \\ & -1 & 2 & -1 & \\ & & & & \end{pmatrix}.$$

At the present time, there are some restrictions as to what type of mappings can be used. In particular, all global rows stored on a processor must be mapped from 0 to  $k - 1$  where  $k$  is the number of rows assigned to this processor. This row mapping induces a partial column mapping. Any additional columns must be mapped with consecutive increasing numbers starting from  $k$ .

**ML** has no notion of global indices and uses only the local indices. In most cases, another package or application already mapped the global indices to local indices and so **ML** works with the existing local indices. Specifically, the parallel version of `user_getrow` and `user_matvec` should correspond to each processor's local matrix. This means that when giving the column information with `ML_Set_Amatrix_Getrow`, the total number of columns in the local matrix should be given and that when row  $k$  is requested, `user_getrow` should return the  $k^{th}$  local row using local column indices. Likewise, the matrix-vector product takes a local input vector and multiplies it by the local matrix. It is important to note that this local input vector does not contain ghost node data (i.e. the input vector is of length `nlocal` where `nlocal` is the number of matrix rows). Thus, `user_matvec` must perform the necessary communication to update ghost variables. When invoking `ML_Init_Amatrix`, the local number of rows should be given for the number of rows and the vector length<sup>14</sup>. A specific communication function must also be passed into **ML** when supplying the `getrow` function so that **ML** can determine how local matrices on different processors are 'glued' together. The signature of the communication function is

```
int user_comm(double x[], void *Adata)
```

where `A.data` is the user-defined data pointer specified in the `ML_Init_Amatrix` and `x` is a vector of length `nlocal_allcolumns` specified in `ML_Set_Amatrix_Getrow`. This parameter should be set to the total number of matrix columns stored on this processor. On input, only the first `nlocal` elements of `x` are filled with data where `nlocal` is the number of rows/columns specified in `ML_Init_Amatrix`. On output, the ghost elements are updated to their current values (defined on other processors). Thus, after this function a local matrix-vector product could be properly performed using `x`. To make all this clear, we give the new functions corresponding to our two processor example.

```
int Poisson_getrow(ML_Operator *Amat, int N_requested_rows, int requested_rows[],
    int allocated_space, int cols[], double values[], int row_lengths[])
{
    int m = 0, i, row, proc, *itemp, start;

    itemp = (int *) ML_Get_MyGetrowData(Amat);
    proc = *itemp;
```

---

<sup>14</sup>In contrast to `ML_Set_Amatrix_Getrow` in which the number of local columns are given (including those that correspond to ghost variables), `ML_Init_Amatrix` does not include ghost variables and so both size parameters should be the number of local rows.

```

for (i = 0; i < N_requested_rows; i++) {
    row = requested_rows[i];
    if (allocated_space < m+3) return(0);
    values[m] = 2; values[m+1] = -1; values[m+2] = -1;
    start = m;
    if (proc == 0) {
        if (row == 0) {cols[m++] = 0; cols[m++] = 2;           }
        if (row == 1) {cols[m++] = 1; cols[m++] = 3;}
    }
    if (proc == 1) {
        if (row == 0) {cols[m++] = 0; cols[m++] = 1; cols[m++] = 4;}
        if (row == 1) {cols[m++] = 1; cols[m++] = 0; cols[m++] = 2;}
        if (row == 2) {cols[m++] = 2; cols[m++] = 1; cols[m++] = 3;}
    }
    row_lengths[i] = m - start;
}
return(1);
}

int Poisson_matvec(ML_Operator *Amat, int in_length, double p[], int out_length,
                  double ap[])
{
    int i, proc, *itemp;
    double new_p[5];

    itemp = (int *) ML_Get_MyMatvecData(Amat);
    proc = *itemp;

    for (i = 0; i < in_length; i++) new_p[i] = p[i];
    Poisson_comm(new_p, A_data);

    for (i = 0; i < out_length; i++) ap[i] = 2.*new_p[i];

    if (proc == 0) {
        ap[0] -= new_p[2];
        ap[1] -= new_p[3];
    }
    if (proc == 1) {
        ap[0] -= new_p[1]; ap[0] -= new_p[4];
        ap[1] -= new_p[2]; ap[1] -= new_p[0];
        ap[2] -= new_p[3]; ap[2] -= new_p[1];
    }
    return 0;
}

and

int Poisson_comm(double x[], void *A_data)
{
    int    proc, neighbor, length, *itemp;
    double send_buffer[2], recv_buffer[2];

    itemp = (int *) A_data;
    proc = *itemp;

    length = 2;
    if (proc == 0) {

```

```

    neighbor = 1;
    send_buffer[0] = x[0]; send_buffer[1] = x[1];
    send_msg(send_buffer, length, neighbor);
    recv_msg(recv_buffer, length, neighbor);
    x[2] = recv_buffer[1]; x[3] = recv_buffer[0];
}
else {
    neighbor = 0;
    send_buffer[0] = x[0]; send_buffer[1] = x[2];
    send_msg(send_buffer, length, neighbor);
    recv_msg(recv_buffer, length, neighbor);
    x[3] = recv_buffer[1]; x[4] = recv_buffer[0];
}
return 0;
}

```

Finally, these matrix functions along with size information are associated with the fine grid discretization matrix via

```

if      (proc == 0) {nlocal = 2; nlocal_allcolumns = 4;}
else if (proc == 1) {nlocal = 3; nlocal_allcolumns = 5;}
else      {nlocal = 0; nlocal_allcolumns = 0;}

ML_Init_Amatrix      (ml_object, 0,  nlocal, nlocal, &proc);
ML_Set_Amatrix_Getrow(ml_object, 0,  Poisson_getrow, Poisson_comm,
                        nlocal_allcolumns);
ML_Set_Amatrix_Matvec(ml_object, 0,  Poisson_matvec);

```

## 13 MLMEX: The MATLAB Interface for ML

MLMEX is ML's interface to the MATLAB environment. It allows access to a limited set of routines either using the ML\_Epetra or the MLAPI interfaces. It is designed to provide access to ML's aggregation and solver routines from MATLAB and does little else. MLMEX allows users to setup and solve arbitrarily many problems, so long as memory suffices. More than one problem can be setup simultaneously.

### 13.1 Configure and Make

In order to use MLMEX, ML must be configured with (at least) the following options:

```

../configure --enable-epetra --enable-teuchos --enable-ifpack \
--enable-aztecoo --enable-galeri --enable-amesos --enable-epetraext \
--enable-ml-matlab \
--with-matlab-exec=<directory with matlab binaries> \
--with-matlab-root=<root directory of matlab install>

```

Additional options (such as `--enable-zoltan`) can be specified as well.

On 64-bit Intel/AMD architectures, ML and all required Trilinos libraries must be compiled with the `-fPIC` option. This necessitates adding `CFLAGS=-fPIC CXXFLAGS=-fPIC FFLAGS=-fPIC` to the Trilinos configure line. Currently, MLMEX will not compile under Solaris.

The build system uses the `mexext` script from Mathworks to detect the appropriate file extension for the architecture. At least at present, this means that cross-compilation is not possible.

## 13.2 Using MLMEX

MLMEX is designed to be interfaced with via the MATLAB script `ml.m`. There are five modes in which MLMEX can be run:

1. Setup Mode — Performs the problem setup for ML. Depending on whether or not the `mlmex: interface` option is used, MLMEX creates either a `ML_Epetra` or `MLAPI` object. This call returns a problem handle used to reference the problem in the future.
2. Solve Mode — Given a problem handle and a right-hand side, ML solves the problem specified. Setup mode must be called before solve mode.
3. Cleanup Mode — Frees the memory allocated to internal ML objects. This can be called with a particular problem handle, in which case it frees that problem, or without one, in which case all MLMEX memory is freed.
4. Status Mode — Prints out status information on problems which have been set up. Like cleanup, it can be called with or without a particular problem handle.
5. Aggregate Mode — Uses MLMEX as an interface to ML's aggregation routines.

All of these modes, with the exception of status and cleanup take option lists which will be directly converted into `Teuchos::ParameterList` objects by MLMEX. This means that the options described in Section 6 for `ML_Epetra` and those in the `MLAPI` guide (for `MLAPI`) will work correctly for MLMEX.

### 13.2.1 Setup Mode

Setup mode is called as follows:

```
>> [h,oc]=ml('setup',A,['parameter',value,...])
```

The parameter `A` represents the sparse matrix to perform aggregation on and the parameter/value pairs represent standard ML options.

The routine returns a problem handle, `h`, and the operator complexity `oc` for the operator. In addition to the standard options, setup mode has one unique option of its own:

`mlmex: interface` [string] Whether to use `ML_Epetra` ('epetra') or `MLAPI` ('mlapi'). Default: 'epetra'.

### 13.2.2 Solve Mode

Solve mode is called as follows:

```
>> ml(h,A,b,['parameter',value,...])
```

The parameter `h` is a problem handle returned by the setup mode call, `A` is the sparse matrix with which to solve and `b` is the right-hand side. Parameter/value pairs are as above. However, a few additional MLAPI options, normally not supported by ML\_Epetra, are supported as well:

<code>krylov: tolerance</code>	[double] Tolerance for the solver. Default: 1e-9.
<code>krylov: max iterations</code>	[int] Maximum number of krylov iterations. Default 1550.
<code>krylov: type</code>	[string] Solver to use, e.g. 'cg', 'gmres', 'fixed point'. Default 'gmres'.
<code>krylov: output level</code>	[int] ML output level. Default 10.
<code>krylov: conv</code>	[int] ML convergence criterion. Default 'r0'.

All of these options are taken directly from MLAPI, so consult its manual for more information.

### 13.2.3 Cleanup Mode

Cleanup mode is called as follows:

```
>> ml('cleanup',[h])
```

The parameter `h` is a problem handle returned by the setup mode call and is optional. If `h` is provided, that problem is cleaned up. If the option is not provided all currently set up problems are cleaned up.

### 13.2.4 Status Mode

Status mode is called as follows:

```
>> ml('status',[h])
```

The parameter `h` is a problem handle returned by the setup mode call and is optional. If `h` is provided, status information for that problem is printed. If the option is not provided all currently set up problems have status information printed.

### 13.2.5 Aggregate Mode

Status mode is called as follows:

```
>> agg=ml('aggregate',A,['parameter',value,...])
```

The parameter `A` represents the sparse matrix to perform aggregation upon. The aggregates are returned and all memory is immediately freed. The purpose of this option is to allow access to ML's aggregation routines in MATLAB.

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