

Pysparse Documentation

Release 1.0.2

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INTRODUCTION TO PYSPARSE

PySparse extends the Python interpreter by a set of sparse matrix types holding double precision values. PySparse also includes modules that implement

- Iterative Krylov methods for solving linear systems of equations,
- Diagonal (Jacobi) and SSOR preconditioners,
- Interfaces to direct solvers for sparse linear systems of equations (SuperLU and UMFPACK),
- A Jacobi-Davidson eigenvalue solver for the symmetric, generalised matrix eigenvalue problem (JDSYM),
- Low-level C classes to represent and manipulate sparse matrices,
- · High-level Python classes with operator overloading to perform usual operations on matrices.

Most of the above modules are implemented as C extension modules for maximum performance.

PySparse uses NumPy for handling dense vectors and matrices and uses SuperLU and UMFPACK for factorizing general sparse matrices.

1.1 Module Overview

1.1.1 spmatrix

The spmatrix module is the foundation of the Pysparse package. It extends the Python interpreter by three new types named ll_mat, csr_mat and sss_mat. These types represent sparse matrices in the LL-, the CSR- and SSS-formats respectively (see *Sparse Matrix Formats*). For all three formats, double precision values (C type double) are used to represent the nonzero entries. The common way to use the spmatrix module is to first build a matrix in the LL-format. The LL-matrix is manipulated until it has its final shape and content. Afterwards it may be converted to either the CSR- or SSS-format, which needs less memory and allows for fast matrix-vector multiplications. A ll_mat object can be created from scratch, by reading data from a file (in MatrixMarket format) or as a result of matrix operation (as e.g. a matrix-matrix multiplication). The ll_mat object supports manipulating (reading, writing, add-updating) single entries or sub-matrices. On the other hand, csr_mat and sss_mat matrices are not constructed directly, instead they are created by converting ll_mat objects. Once created, csr_mat and sss_mat objects cannot be manipulated. Their purpose is to support efficient matrix-vector multiplications.

1.1.2 itsolvers

The itsolvers module provides a set of iterative methods for solving linear systems of equations. The iterative methods are callable like ordinary Python functions. All these functions expect the same parameter list, and all function

return values also follow a common standard. Any user-defined iterative solvers should also follow these conventions, since other PySparse modules rely on them (e.g. the jdsym module).

Currently the itsolvers module contains the following iterative methods: PCG, MINRES, QMRS, BICGSTAB and CGS.

1.1.3 precon

The precon module provides preconditioners, which can be used e.g. for the iterative methods implemented in the itsolvers module or the JDSYM eigensolver (in the jdsym module). In the PySparse framework, any Python object that has the following properties can be used as a preconditioner:

- a shape attribute, which returns a 2-tuple describing the dimension of the preconditioner,
- a precon method, that accepts two vectors x and y, and applies the preconditioner to x and stores the result in y. Both x and y are double precision, rank-1 NumPy arrays of appropriate size.

The precon module currently implements m-step Jacobi and m-step SSOR preconditioners.

1.1.4 superlu

The superlu module interfaces the SuperLU library to make it usable by Python code. SuperLU is a software package written in C for the direct solution of a general linear system of equations. SuperLU computes LU-factorizations of general non-symmetric, sparse matrices with partial pivoting. It is also applicable to rectangular systems of equations.

1.1.5 umfpack

The umfpack module interfaces the UMFPACK factorization package. UMFPACK computes the LU factorization of a general matrix with partial pivoting. It is also applicable to rectangular systems.

The main difference between the superlu and umfpack modules resides in the way the factorization is performed internally. SuperLU works with the concept of *supernodes* leading naturally to parallelism in the factorization. If available, a custom-built SuperLU library for multi-core processors can be supplied to Pysparse in place of the default library. Both factorization packages rely intensively on the BLAS to operate on dense sub-matrices. Provided the BLAS library supplied to Pysparse was compiled with multi-threading, some level of parallelism will also be available in UMFPACK. However, a rough empirical observation is that UMFPACK is often faster than SuperLU on monoprocessor machines.

1.1.6 jdsym

The jdsym module provides an implementation of the JDSYM algorithm, that is conveniently callable from Python. The JDSYM algorithm computes solutions of large sparse symmetric (genralised or standard) eigenvalue problems. JDSYM is an implementation of the Jacobi-Davidson method, optimized for symmetric matrices.

1.2 Prerequisites

- NumPy
- Optionally, a custom-built UMFPACK and/or SuperLU

1.3 Installing Pysparse

python setup.py install

1.4 Testing Pysparse

From the Test directory, testSuperLU runs a series of tests to exercise the various options of the SuperLU direct solver:

\$ python testSu	perlu.py					
Test	RelErr	Tol	nnz(A)	nnz(L+U)	Fact	Solve
poild-dflt	1.78e-12	2.25e-07	99999 99999	199998	0.06	0.00
poild-size	1.78e-12	2.25e-07	99999	199998	0.05	0.00
poild-relx	1.78e-12	2.25e-07	99999	200758	0.06	0.00
poild-trsh	1.78e-12	2.25e-07	99999	199998	0.06	0.00
poild-prm0	1.44e-12	2.25e-07	99999	199998	0.04	0.00
poild-prm1	1.78e-12	2.25e-07	99999	199998	0.07	0.00
poild-prm2	1.78e-12	2.25e-07	99999	199998	0.06	0.00
poild-prm3	1.44e-12	2.25e-07	99999	200000	0.06	0.00
poi2d-dftl	2.55e-16	3.60e-12	119600	1952434	0.47	0.02
poi2d-size	3.39e-16	3.60e-12	119600	1952434	0.42	0.02
poi2d-relx	2.60e-16	3.60e-12	119600	2000252	0.48	0.02
poi2d-trsh	2.55e-16	3.60e-12	119600	1952434	0.48	0.02
poi2d-prm0	2.69e-15	3.60e-12	119600	16000398	5.48	0.11
poi2d-prm1	7.00e-16	3.60e-12	119600	3506336	0.89	0.03
poi2d-prm2	2.55e-16	3.60e-12	119600	1952434	0.47	0.02
poi2d-prm3	2.24e-15	3.60e-12	119600	3472176	0.82	0.03
spdgs-trsh	4.44e-16	2.22e-14	29998	39998	0.01	0.00
spdgs-prm0	4.44e-16	2.22e-14	29998	40000	0.01	0.00
spdgs-prm1	4.44e-16	2.22e-14	29998	40002	0.01	0.00
spdgs-prm3	4.44e-16	2.22e-14	29998	40002	0.01	0.00

Ran 20 tests in 12.675s

OK

There is a corresponding test script for UMFPACK, testUmfpack:

\$pyt	<pre>\$python testUmfpack.py</pre>							
	RelErr	Tol	nnz(A)	nnz(L)	nnz (U)	Fact	Solve	
	1.44e-12	2.25e-07	 149998	99999	99999	0.13	0.01	
	1.44e-12	2.25e-07	149998	99999	99999	0.13	0.01	
	1.44e-12	2.25e-07	149998	99999	99999	0.12	0.01	
	1.44e-12	2.25e-07	149998	99999	99999	0.12	0.01	
	1.44e-12	2.25e-07	149998	99999	99999	0.12	0.01	
	1.44e-12	2.25e-07	149998	99999	99999	0.14	0.01	
	1.55e-17	3.60e-12	199200	1081911	1081911	0.54	0.03	
	1.55e-17	3.60e-12	199200	1081911	1081911	0.53	0.03	
	2.50e-17	3.60e-12	199200	1081911	1081911	0.53	0.03	
	2.50e-17	3.60e-12	199200	1081911	1081911	0.53	0.03	
	1.55e-17	3.60e-12	199200	1081911	1081911	0.53	0.03	
	1.64e-17	3.60e-12	199200	1489438	2166768	1.00	0.04	

4.44e-16	2.22e-14	29998	19999	19999	0.03	0.00
4.44e-16	2.22e-14	29998	19999	19999	0.02	0.00
4.44e-16	2.22e-14	29998	19999	19999	0.02	0.00
4.44e-16	2.22e-14	29998	19999	19999	0.02	0.00
4.44e-16	2.22e-14	29998	19999	19999	0.03	0.00
4.44e-16	2.22e-14	29998	19999	19999	0.02	0.00

Ran 18 tests in 8.486s

1.5 Generating the Documentation

To re-generate the documentation, Sphinx version 0.5 or higher must be installed. The jsMath package must also be installed. Edit \$PYSPARSE/Doc/pysparse/source/conf.py to specify the location of jsMath.

To re-generate the html documentation,

```
cd $PYSPARSE/Doc/pysparse
make html
```

where Doc is a subdirectory of the top Pysparse directory. You can then point your browser to the file build/index.html.

Similarly, to re-generate the pdf documentation,

```
cd $PYSPARSE/Doc/pysparse
make latex
cd build/latex
make all-pdf
```

This creates Pysparse.pdf in the current directory. Obviously, you need to have a working LaTeX distribution installed.

SPARSE MATRIX FORMATS

This section describes the sparse matrix storage schemes available in Pysparse. It also covers sparse matrix creation, population and conversion.

- Linked-list format (LL): a convenient format for creating and populating a sparse matrix, whether symmetric or general.
- Compressed sparse row format (CSR): a format designed to speed up matrix-vector products, but not well suited to matrix population and manipulation.
- Sparse Skyline format (SSS): a format for symmetric matrices designed to speed up matrix-vector products, but not well suited to matrix population and manipulation.

2.1 Linked-List Format

The linked-list format allows insertion and lookup of nonzero elements in moderate time and without having to move too much data around. Internally, the nonzero entries of a matrix are stored row by row in a linked list. Within a given row, column indices are sorted in ascending order.

In Pysparse, matrices in linked-list format are created by using the 11_mat class.

This format resembles a sorted version of the coordinate format but with a data structure that lends itself to fast insertion, removal and lookup.

Typically, a new matrix should be created as an ll_mat and populated. If necessary, it can then be converted to compressed sparse row or sparse skyline format using the to_csr() and to_sss() methods.

The data structure for a matrix in linked-list format has the following components:

- val The double precision array val of length nalloc contains the non-zero entries of matrix.
- col The integer array col of length nalloc contains the column indices of the non-zero entries stored in val.
- **link** the integer array link of length nalloc stores the pointer (index) to the next non-zero entry of the same row. A value of -1 indicates that there is no next entry.
- root The integer array root of length n contains the pointers to the first entry of each row. The other entries of the same row can be located by following the link array.
- **free** The integer free points to the first entry of the *free list*, i.e. a linked list of unoccupied spots in the val and col arrays. This list is populated when non-zero entries are removed from the matrix.

Here n is the number of rows of the matrix and nalloc is number of allocated elements in the arrays val, col and link. Note that the number of nonzero entries stored is less than or equal to nalloc, but the val, col and link arrays can be enlarged dynamically if necessary.

2.2 Compressed Sparse Row Format

In CSR format, a sparse matrix is represented via three arrays:

- va The double precision array va of length nnz contains the non-zero entries of the matrix, stored row by row.
- ja The integer array ja of length nnz contains the column indices of the non-zero entries stored in va.
- ia The integer array ia of length n + 1 contains the pointers (indices) to the beginning of each row in the arrays va and ja. The last element of ia always has the value nnz + 1.

Here n is the number of rows of the matrix and nnz is its number of nonzero entries.

This format is particularly interesting for computing matrix-vector products. Even though the order of the entries is not prescribed in this format, we sort the entries of each row by ascending column indices. This enables us to use more efficient algorithms for certain operations.

2.3 Sparse Skyline Format

The SSS format is closely related to the CSR format. It is often used for sparse *symmetric* matrices. The diagonal is stored in a separate (full) vector and the strict lower triangle is stored in CSR format:

- va The double precision array va of length nnz contains the non-zero entries of the strict lower triangle, stored row by row.
- ja The integer array ja of length nnz contains the column indices of the non-zero entries stored in va.
- ia The integer array ia of length n + 1 contains the pointers (indices) to the beginning of each row in the arrays va and ja. The last element of ia always has the value nnz + 1.
- da The double precision array da of length n stores all diagonal entries of the matrix.

Here n is the order of the matrix and nnz is the number of nonzero entries in the strict lower triangle.

We sort the entries of each row by ascending column indices, like we do with the CSR format. The SSS format has the advantage over the CSR format, that it requires roughly half of the storage space and that the matrix-vector multiplication can be implemented more efficiently

LOW-LEVEL SPARSE MATRIX TYPES

3.1 The spmatrix Module

The spmatrix module is the foundation of the PySparse package. It extends the Python interpreter by three new types named ll_mat, csr_mat and sss_mat. These types represent sparse matrices in the LL-, the CSR- and SSS-formats respectively (see *Sparse Matrix Formats*). For all three formats, double precision values (C type double) are used to represent the non-zero entries.

The common way to use the spmatrix module is to first build a matrix in the LL-format. The LL-matrix is manipulated until it has its final shape and content. Afterwards it may be converted to either the CSR- or SSS-format, which needs less memory and allows for fast matrix-vector multiplications.

A 11_mat object can be created from scratch, by reading data from a file (in MatrixMarket format) or as a result of matrix operation (as e.g.a matrix-matrix multiplication). The 11_mat object supports manipulating (reading, writing, add-updating) single entries or sub-matrices.

csr_mat and sss_mat are not constructed directly, instead they are created by converting ll_mat objects. Once created, csr_mat and sss_mat objects cannot be manipulated. Their purpose is to support efficient matrix-vector multiplications.

3.1.1 spmatrix module functions

11 mat (*n*, *m*, *sizeHint=1000*)

Creates a ll_mat object, that represents a general, all zero $m \times n$ matrix. The optional sizeHint parameter specifies the number of non-zero entries for which space is allocated initially.

If the total number of non-zero elements of the final matrix is known (approximately), this number can be passed as sizeHint. This will avoid costly memory reallocations.

11_mat_sym (*n*, *sizeHint=1000*)

Creates a ll_mat object, that represents a *symmetric*, all zero $n \times n$ matrix. The optional sizeHint parameter specifies, how much space is initially allocated for the matrix.

11_mat_from_mtx(fileName)

Creates a ll_mat object from a file named fileName, which must be in MatrixMarket Coordinate format. Depending on the file content, either a symmetric or a general sparse matrix is generated.

matrixmultiply(A, B)

Computes the matrix-matrix multiplication $\mathbf{C} := \mathbf{AB}$ and returns the result \mathbf{C} as a new 11_mat object representing a general sparse matrix. The parameters \mathbf{A} and \mathbf{B} are expected to be objects of type 11_mat.

dot(A, B)

Computes the dot-product $C := A^TB$ and returns the result C as a new ll_mat object representing a general sparse matrix. The parameters A and B are expected to be objects of type ll_mat.

3.1.2 11 mat objects

11_mat objects represent matrices stored in the LL format, which are described in *Sparse Matrix Formats*. 11_mat objects come in two flavours: general matrices and symmetric matrices. For symmetric matrices only the non-zero entries in the lower triangle are stored. Write operations to the strictly upper triangle are prohibited for the symmetric format. The issym attribute of an 11_mat object can be queried to find out whether or not the symmetric storage format is used.

The entries of a matrix can be accessed conveniently using two-dimensional array indices. In the Python language, subscripts can be of any type (as it is customary for dictionaries). A two-dimensional index can be regarded as a 2-tuple (the brackets do not have to be written, so A[1,2] is the same as A[(1,2)]). If both tuple elements are integers, then a single matrix element is referenced. If at least one of the tuple elements is a slice (which is also a Python object), then a submatrix is referenced.

Subscripts have to be decoded at runtime. This task includes type checks, extraction of indices from the 2-tuple, parsing of slice objects and index bound checks. Following Python conventions, indices start with 0 and wrap around (so -1 is equivalent to the last index).

The following code creates an empty 5×5 matrix A, sets all diagonal elements to their respective row/column index and then copies the value of A[0,0] to A[2,1]:

```
>>> from pysparse import spmatrix
>>> A = spmatrix.ll_mat(5, 5)
>>> for i in range(5):
...         A[i,i] = i+1
>>> A[2,1] = A[0,0]
>>> print A
ll_mat(general, [5,5], [(0,0): 1, (1,1): 2, (2,1): 1, (2,2): 3, (3,3): 4, (4,4): 5])
```

The Python slice notation can be used to conveniently access sub-matrices.

```
>>> print A[:2,:]  # the first two rows
ll_mat(general, [2,5], [(0,0): 1, (1,1): 2])
>>> print A[:,2:5]  # columns 2 to 4
ll_mat(general, [5,3], [(2,0): 3, (3,1): 4, (4,2): 5])
>>> print A[1:3,2:5]  # submatrix from row 1 col 2 to row 2 col 4
ll_mat(general, [2,3], [(1,0): 3])
```

The slice operator always returns a new ll_mat object, containing a **copy** of the selected submatrix.

Write operations to slices are also possible:

```
>>> B = 11_mat(2, 2)  # create 2-by-2

>>> B[0,0] = -1; B[1,1] = -1  # diagonal matrix

>>> A[:2,:2] = B  # assign it to upper

>>> print A

11_mat(general, [5,5], [(0,0): -1, (1,1): -1, (2,1): 1,

(2,2): 3, (3,3): 4, (4,4): 5])
```

Fancy Indexing

There is flexibility in the way submatrices of 11_mat objects can be accessed. In particular, rows and columns can be permuted arbitrarily and submatrices need not be composed of consecutive rows or indices. Let's look at an example.

Below, the poisson1d() function assembles a Poisson matrix. We come back to Poisson matrices later in this section.

```
>>> from pysparse import poisson
>>> n = 5
>>> A = poisson.poisson1d(n)
>>> print A # Original matrix
11_mat(general, [5,5]):
2.000000 -1.000000 ------
-1.000000 2.000000 -1.000000 -----
 ----- -1.000000 2.000000 -1.000000 -----
----- -1.000000 2.000000 -1.000000
----- -1.000000 2.000000
>>> print A[n-1:1:-1,1:n-1] # Rows 2 through n-1 in reverse order,
                      # second through one before last col
11_mat(general, [3,3]):
 ----- -1.000000
----- -1.000000 2.000000
-1.000000 2.000000 -1.000000
>>> print A[::-1,:] # Reverse row order
11_mat(general, [5,5]):
----- -1.000000 2.000000
        ----- -1.000000 2.000000 -1.000000
----- -1.000000 2.000000 -1.000000 ------
-1.000000 2.000000 -1.000000 -----
2.000000 -1.000000 ------
>>> print A[:,::-1] # Reverse col order (same as above b/c A is symmetric)
11_mat(general, [5,5]):
------ -1.000000 2.000000
----- -1.000000 2.000000 -1.000000
----- -1.000000 2.000000 -1.000000 -----
-1.000000 2.000000 -1.000000 -----
2.000000 -1.000000 ------
>>> print A[::-1,::-1] # Reverse row and col order (same as original matrix)
11_mat(general, [5,5]):
2.000000 -1.000000 ------ -----
-1.000000 2.000000 -1.000000 -----
----- -1.000000 2.000000 -1.000000 ------
----- -1.000000 2.000000 -1.000000
----- -1.000000 2.000000
>>> print A[1:3,3:]  # Rows 1 and 2, cols 3 and up
ll_mat(general, [2,2]):
_____
-1.000000 -----
>>> print A[::2,::2] # Every other row and col
11_mat(general, [3,3]):
2.000000 -----
----- 2.000000 -----
----- 2.000000
```

Keep in mind that as always with Python slices, the final index is never included. Note also that slicing always returns a general matrix. Even though it might be symmetric, both triangles are stored. Finally, slicing should be applied to

general matrices. If applied to symmetric matrices, only a partial result is returned.

Fancy indexing can also be done with Python lists:

or with integer Numpy arrays:

Finally, fancy indexing can be used to assign the same numerical value to a submatrix:

Notice however that although the slice [0:3] appears to amount to the list [0,1,2], the assignments A[:3,:3]=7 and A.put([7,7,7], [0,1,2], [0,1,2]) produce **very different** results.

Warning: For large-scale matrices, fancy indexing is most efficient when both index sets have the same type: two Python slices or two Python lists. When the index sets have different types, index arrays are built internally and this results in a performance hit.

11_mat Object Attributes and Methods

```
class 11_mat()
```

A general sparse matrix class in linked-list format which also allows the representation of symmetric matrices. Only the lower triangle of a symmetric matrix is kept in memory for efficiency.

shape

Returns a 2-tuple containing the shape of the matrix A, i.e. the number of rows and columns.

nnz

Returns the number of non-zero entries stored in matrix **A**. If **A** is stored in symmetric format, only the number of non-zero entries in the lower triangle (including the diagonal) are returned.

issym

Returns true (a non-zero integer) if matrix **A** is stored in the symmetric LL format, i.e. only the non-zero entries in the lower triangle are stored. Returns false (zero) if matrix **A** is stored in the general LL format.

matvec(x, y)

Computes the sparse matrix-vector product $\mathbf{y} := \mathbf{A}\mathbf{x}$ where \mathbf{x} and \mathbf{y} are double precision, rank-1 NumPy arrays of appropriate size.

$matvec_transp(x, y)$

Computes the transposed sparse matrix-vector product $\mathbf{y} := \mathbf{A}^T \mathbf{x}$ where \mathbf{x} and \mathbf{y} are double precision, rank-1 NumPy arrays of appropriate size. For sss_mat objects matvec_transp is equivalent to matvec.

to_csr()

This method converts a sparse matrix in linked list format to compressed sparse row format. Returns a newly allocated csr_mat object, which results from converting matrix **A**.

to sss()

This method converts a sparse matrix in linked list format to sparse skyline format. Returns a newly allocated <code>sss_mat</code> object, which results from converting matrix **A**. This function works for <code>ll_mat</code> objects in both the symmetric and the general format. If **A** is stored in general format, only the entries in the lower triangle are used for the conversion. No check whether **A** is symmetric is performed.

export_mtx (fileName, precision=6)

Exports the matrix A to file named fileName. The matrix is stored in MatrixMarket Coordinate format. Depending on the properties of the ll_mat object A the generated file either uses the symmetric or a general MatrixMarket Coordinate format. The optional parameter precision specifies the number of decimal digits that are used to express the non-zero entries in the output file.

shift (sigma, M)

Performs the daxpy operation $A \leftarrow A + \sigma M$. The parameter σ is expected to be a Python Float object. The parameter M is expected to an object of type ll_mat of compatible shape.

copy (

Returns a new 11_mat object that is a (deep) copy of the 11_mat object A. So:

```
>>> B = A.copy()
```

is equivalent to:

```
>>> B = A[:,:]
```

On the other hand:

```
>>> B = A.copy()
```

is *not* the same as:

```
>>> B = A
```

The latter version only returns a reference to the same object and assigns it to B. Subsequent changes to A will therefore also be visible in B.

update add mask (B, ind0, ind1, mask0, mask1)

This method is provided for efficiently assembling global finite element matrices. The method adds the matrix \mathbf{B} to entries of matrix \mathbf{A} . The indices of the entries to be updated are specified by the integer arrays ind0 and ind1. The individual updates are enabled or disabled using the mask0 and mask1 arrays.

The operation is equivalent to the following Python code:

All five parameters are NumPy arrays. ${\bf B}$ is a rank-2 array. The four remaining parameters are rank-1 arrays. Their length corresponds to either the number of rows or the number of columns of ${\bf B}$.

This method is not supported for ll_mat objects of symmetric type, since it would generally result in an non-symmetric matrix. update_add_mask_sym must be used in that case. Attempting to call this method using a ll_mat object of symmetric type will raise an exception.

update_add_mask_sym(B, ind, mask)

This method is provided for efficiently assembling symmetric global finite element matrices. The method adds the matrix ${\bf B}$ to entries of matrix ${\bf A}$. The indices of the entries to be updated are specified by the integer array ind. The individual updates are enabled or disabled using the mask array.

The operation is equivalent to the following Python code:

The three parameters are all NumPy arrays. $\bf B$ is a rank-2 array representing a square matrix. The two remaining parameters are rank-1 arrays. Their length corresponds to the order of matrix $\bf B$.

update add at (val, irow, jcol)

Add in place the elements of the vector val at the indices given by the two arrays irow and jcol. The operation is equivalent to:

```
for i in range(len(val)):
    A[irow[i], jcol[i]] += val[i]
```

generalize()

Convert ll_mat object to non-symmetric form in place.

compress()

Frees memory by reclaiming unused space in the internal data structure. Returns the number of elements freed.

norm(p)

Returns the p-norm of a matrix, where p is a string. If p='1', the 1-norm is returned, if p='inf', the infinity-norm is returned, and if p='fro', the Frobenius norm is returned.

Note: The 1 and infinity norm are not yet implemented for symmetric matrices.

keys()

Return a list of tuples (i, j) of the nonzero matrix entries.

values()

Return a list of the nonzero matrix entries as floats.

items()

Return a list of tuples (indices, value) of the nonzero entries keys and values. The indices are themselves tuples (i, j) of row and column values.

scale (sigma)

Scale each element in the matrix by the constant sigma.

take (val, irow, jcol)

Extract elements at positions (irow[i], jcol[i]) and place them in the array val. In other words:

```
for i in range(len(val)): val[i] = A[irow[i], jcol[i]]
```

put (val, irow, icol)

put takes the opposite tack to take. Place the values in val at positions given by irow and jcol:

```
for i in range(len(val)): A[irow[i], jcol[i]] = val[i]
```

Here, irow and jcol can be Python lists or integer Numpy arrays. If either irow or jcol is omitted, it is replaced with [0, 1, 2, ...]. Similarly, val can be a Python list, an integer Numpy array or a single scalar. If val is a scalar v, it has the same effect as if it were the constant list or array [v, v, ..., v].

delete rows (mask)

Delete rows in place. If mask[i] == 0, the i-th row is deleted. This operation does not simply zero out rows, they are *removed*, i.e., the resulting matrix is *smaller*.

delete cols(mask)

Similar to delete_rows only with columns.

delete rowcols (mask)

If mask[i] == 0 both the i-th row and the i-th column are deleted.

find()

Returns a triple (val, irow, jcol) of Numpy arrays containing the matrix in coordinate format. There is a nonzero element with value val[i] in position (irow[i], jcol[i]).

3.1.3 csr_mat and sss_mat Objects

csr_mat objects represent matrices stored in the CSR format, which are described in *Sparse Matrix Formats*. sss_mat objects represent matrices stored in the SSS format (c.f. *Sparse Matrix Formats*). The only way to create a csr_mat or a sss_mat object is by conversion of a ll_mat object using the to_csr() or the to_sss() method respectively. The purpose of the csr_mat and the sss_mat objects is to provide fast matrix-vector multiplications for sparse matrices. In addition, a matrix stored in the CSR or SSS format uses less memory than the same matrix stored in the LL format, since the link array is not needed.

csr_mat and sss_mat objects do not support two-dimensional indices to access matrix entries or sub-matrices. Again, their purpose is to provide fast matrix-vector multiplication.

csr_mat and sss_mat Object Attributes and Methods

class csr mat()

A general sparse matrix class in compressed sparse row format which also allows the representation of symmetric matrices. Only the lower triangle of a symmetric matrix is kept in memory for efficiency.

class sss mat()

A general sparse matrix class in sparse skyline format which also allows the representation of symmetric matrices. Only the lower triangle of a symmetric matrix is kept in memory for efficiency.

shape

Returns a 2-tuple containing the shape of the matrix **A**, i.e. the number of rows and columns.

nnz

Returns the number of non-zero entries stored in matrix A. If A is an sss_mat object, the non-zero entries in the strictly upper triangle are not counted.

matvec(x, y)

Computes the sparse matrix-vector product $\mathbf{y} := \mathbf{A}\mathbf{x}$ where \mathbf{x} and \mathbf{y} are double precision, rank-1 NumPy arrays of appropriate size.

matvec transp(x, y)

Computes the transposed sparse matrix-vector product $\mathbf{y} := \mathbf{A}^T \mathbf{x}$ where \mathbf{x} and \mathbf{y} are double precision, rank-1 NumPy arrays of appropriate size. For sss_mat objects matvec_transp is equivalent to matvec.

3.2 Example: 2D-Poisson matrix

This section illustrates the use of the spmatrix module to build the well known 2D-Poisson matrix resulting from a $n \times n$ square grid:

```
from pysparse import spmatrix
def poisson2d(n):
   n2 = n * n
   L = spmatrix.ll_mat(n2, n2, 5*n2-4*n)
    for i in range(n):
        for j in range(n):
            k = i + n * j
            L[k,k] = 4
            if i > 0:
               L[k, k-1] = -1
            if i < n-1:
               L[k, k+1] = -1
            if j > 0:
               L[k,k-n] = -1
            if j < n-1:
               L[k, k+n] = -1
    return L
```

Using the symmetric variant of the ll_mat object, this gets even shorter:

```
def poisson2d_sym(n):
    n2 = n*n
    L = spmatrix.ll_mat_sym(n2, 3*n2-2*n)
    for i in range(n):
        k = i + n*j
        L[k,k] = 4
        if i > 0:
        L[k,k-1] = -1
        if j > 0:
        L[k,k-n] = -1
    return L
```

To illustrate the use of the slice notation to address sub-matrices, let's build the 2D Poisson matrix using the diagonal and off-diagonal blocks:

```
def poisson2d_sym_blk(n):
    n2 = n*n
    L = spmatrix.ll_mat_sym(n2, 2*n2-2*n)
    I = spmatrix.ll_mat_sym(n, n)
    P = spmatrix.ll_mat_sym(n, 2*n-1)
    for i in range(n):
        I[i,i] = -1
    for i in range(n):
        P[i,i] = 4
    if i > 0: P[i,i-1] = -1
    for i in range(0, n*n, n):
        L[i:i+n,i:i+n] = P
        if i > 0: L[i:i+n,i-n:i] = I
    return L
```

3.3 Vectorization

The put method of ll_mat objects allows us to operate on entire arrays at a time. This is advantageous because the loop over the elements of an array is performed at C level instead of in the Python script.

If you need to put the same value in many places, put lets you specify this value as a floating-point number instead of an array, e.g.:

```
A.put(4.0, range(n), range(n))
is perfectly equivalent to:
```

```
A.put(4*numpy.ones(n), range(n), range(n))
```

Moreover, if the second index set is omitted, it defaults to range (n) where n is the appropriate matrix dimension. So the above is again perfectly equivalent to:

```
A.put (4.0, range(n))
```

For illustration, let's rewrite the poisson2d, poisson2d_sym and poisson2d_sym_blk constructors.

The put method can be used in poisson2d as so:

```
from pysparse import spmatrix
import numpy

def poisson2d_vec(n):
    n2 = n*n
    L = spmatrix.ll_mat(n2, n2, 5*n2-4*n)
    d = numpy.arange(n2, dtype=numpy.int)
    L.put(4.0, d)
    L.put(-1.0, d[:-n], d[n:])
    L.put(-1.0, d[n:], d[:-n])
    for i in xrange(n):
        di = d[i*n:(i+1)*n]
        L.put(-1.0, di[1:], di[:-1])
        L.put(-1.0, di[:-1], di[1:])
    return L
```

And similarly in the symmetric version:

```
def poisson2d_sym_vec(n):
    n2 = n*n
    L = spmatrix.ll_mat_sym(n2, 3*n2-2*n)
    d = numpy.arange(n2, dtype=numpy.int)
    L.put(4.0, d)
    L.put(-1.0, d[n:], d[:-n])
    for i in xrange(n):
        di = d[i*n:(i+1)*n]
        L.put(-1.0, di[:-1], di[1:])
    return L
```

The time differences to construct matrices with and without vectorization can be dramatic. The following timings were generated on a 2.4GHz Intel Core2 Duo processor:

3.3. Vectorization 17

```
In [1]: from pysparse import poisson
In [2]: import poisson_vec
In [3]: timeit -n10 -r3 L = poisson.poisson2d(100)
10 loops, best of 3: 38.2 ms per loop
In [4]: %timeit -n10 -r3 L = poisson_vec.poisson2d_vec(100)
10 loops, best of 3: 4.26 ms per loop
In [5]: %timeit -n10 -r3 L = poisson.poisson2d(300)
10 loops, best of 3: 352 ms per loop
In [6]: %timeit -n10 -r3 L = poisson_vec.poisson2d_vec(300)
10 loops, best of 3: 31.7 ms per loop
In [7]: timeit -n10 -r3 L = poisson.poisson2d(500)
10 loops, best of 3: 980 ms per loop
In [8]: %timeit -n10 -r3 L = poisson_vec.poisson2d_vec(500)
10 loops, best of 3: 86.4 ms per loop
In [9]: %timeit -n10 -r3 L = poisson.poisson2d(1000)
10 loops, best of 3: 4.02 s per loop
In [10]: %timeit -n10 -r3 L = poisson_vec.poisson2d_vec(1000)
10 loops, best of 3: 333 ms per loop
and for the symmetric versions:
In [18]: timeit -n10 -r3 L = poisson.poisson2d_sym(100)
10 loops, best of 3: 22.6 ms per loop
In [19]: %timeit -n10 -r3 L = poisson_vec.poisson2d_sym_vec(100)
10 loops, best of 3: 2.48 ms per loop
In [20]: timeit -n10 -r3 L = poisson.poisson2d_sym(300)
10 loops, best of 3: 202 ms per loop
In [21]: %timeit -n10 -r3 L = poisson_vec.poisson2d_sym_vec(300)
10 loops, best of 3: 20 ms per loop
In [22]: timeit -n10 -r3 L = poisson.poisson2d_sym(500)
10 loops, best of 3: 561 ms per loop
In [23]: %timeit -n10 -r3 L = poisson_vec.poisson2d_sym_vec(500)
10 loops, best of 3: 53.8 ms per loop
In [24]: \frac{1}{24}: \frac{1}{24
10 loops, best of 3: 2.26 s per loop
In [25]: %timeit -n10 -r3 L = poisson_vec.poisson2d_sym_vec(1000)
10 loops, best of 3: 205 ms per loop
```

From these numbers, it is obvious that vectorizing is crucial, especially for large matrices. The gain in terms of time seems to be a factor of at least four or five. Note that the last system has order one million.

Finally, the block version could be written as:

```
def poisson2d_vec_sym_blk(n):
    n2 = n*n
    L = spmatrix.ll_mat_sym(n2, 3*n2-2*n)
    D = spmatrix.ll_mat_sym(n, 2*n-1)
    d = numpy.arange(n, dtype=numpy.int)
    D.put(4.0, d)
    D.put(-1.0, d[1:], d[:-1])
```

```
P = spmatrix.ll_mat_sym(n, n-1)
P.put(-1,d)
for i in xrange(n-1):
    L[i*n:(i+1)*n, i*n:(i+1)*n] = D
    L[(i+1)*n:(i+2)*n, i*n:(i+1)*n] = P
# Last diagonal block
L[-n:,-n:] = D
return L
```

Here, put is sufficiently efficient that the benefit of constructing the matrix by blocks is not apparent anymore. The slicing and block notation can nevertheless be used for clarity. It could also be implemented as a combination of find and put, at the expense of memory consumption.

```
In [9]: %timeit -n10 -r3 L = poisson.poisson2d_sym_blk(1000)
10 loops, best of 3: 246 ms per loop
In [10]: %timeit -n10 -r3 L = poisson_vec.poisson2d_sym_blk_vec(1000)
10 loops, best of 3: 232 ms per loop
```

3.4 Matlab Implementation

Let's compare the performance of three python codes above with the following Matlab functions:

The Matlab function poisson2d is equivalent to the Python function with the same name

The function poisson2d_blk is an adaption of the Python function poisson2d_sym_blk (except for exploiting the symmetry, which is not directly supported in Matlab).

```
function L = poisson2d_blk(n)
    e = ones(n,1);
    P = spdiags([-e 4*e -e], [-1 0 1], n, n);
    I = -speye(n);
    L = sparse(n*n);
    for i = 1:n:n*n
        L(i:i+n-1,i:i+n-1) = P;
        if i > 1, L(i:i+n-1,i-n:i-1) = I; end
        if i < n*n - n, L(i:i+n-1,i+n:i+2*n-1) = I; end
end</pre>
```

The function poisson2d_kron demonstrates one of the most efficient ways to generate the 2D Poisson matrix in Matlab.

```
function L = poisson2d_kron(n)
    e = ones(n,1);
    P = spdiags([-e 2*e -e], [-1 0 1], n, n);
    L = kron(P, speye(n)) + kron(speye(n), P);
```

The Matlab functions above were place in a Matlab script names poisson.m which takes n as argument. It then calls poisson2d_poisson2d_blk and poisson2d_kron successively, surrounding each call by tic and toc. The tests were performed on a 2.4GHz Intel Core2 Duo running Matlab 7.6.0.324 (R2008a).

The results are as follows:

```
>> poisson(100)
poisson2d
              Elapsed time is 1.731940 seconds.
poisson2d_blk Elapsed time is 0.804837 seconds.
poisson2d_kron Elapsed time is 0.019118 seconds.
>> poisson(300)
poisson2d
              Elapsed time is 145.979044 seconds.
poisson2d_blk Elapsed time is 32.785585 seconds.
poisson2d_kron Elapsed time is 0.215165 seconds.
>> poisson(500)
poisson2d
              Elapsed time is 2318.512099 seconds.
poisson2d_blk Elapsed time is 292.355093 seconds.
poisson2d_kron Elapsed time is 0.627137 seconds.
>> poisson(1000)
poisson2d_kron Elapsed time is 2.317660 seconds.
```

It is striking to see how slow the straightforward poisson2d version is in Matlab. As we see in the next section, the Python version is faster by several orders of magnitude.

3.5 Comparison with Matlab

First, consider the simple Poisson2D function. The *table below* summarizes the results of the previous section by giving timing ratios between the Python and Matlab Poisson constructors.

n	Matlab	Python	Python_vec	Matlab/Python	Matlab/Python_vec
100	1.73	0.0382	0.00426	45.53	406.1
300	145.98	0.3520	0.0317	414.72	4605.0
500	2318.51	0.9800	0.0864	2365.8	26834.6
1000	$ \infty $	4.02	0.333	∞	$ \infty $

Table 3.1: Matlab vs. Python: Construction of 2D Poisson matrices.

Unfortunately, since Matlab does not explicitly support symmetric matrices, we cannot compare the other functions. For information only, we compare the block version of the Python constructor with the Kronecker-product version of the Matlab constructor. The results are in the *next table*.

Table 3.2: Matlab vs. Python: Construction of 2D Poisson matrices—Fastest methods.

n	Matlab	Python_vec	Matlab/Python_vec
100	0.01912	0.0025	7.65
300	0.2152	0.0219	9.83
500	0.6271	0.0631	9.94
1000	2.318	0.232	9.99

PRECONDITIONERS

4.1 The precon Module

The precon module provides preconditioners, which can be used e.g. for the iterative methods implemented in the in the itsolvers module or the JDSYM eigensolver (in the jdsym module).

In the Pysparse framework, any Python object that has the following properties can be used as a preconditioner:

- a shape attribute, which returns a 2-tuple describing the dimension of the preconditioner,
- a precon method, that accepts two vectors x and y, and applies the preconditioner to x and stores the result in y. Both x and y are double precision, rank-1 NumPy arrays of appropriate size.

The precon module implements two new object types jacobi and ssor, representing Jacobi and SSOR preconditioners.

4.1.1 precon Module Functions

jacobi(A, omega=1.0, steps=1)

Creates a jacobi object, representing the Jacobi preconditioner. The parameter A is the system matrix used for the Jacobi iteration. The matrix needs to be subscriptable using two-dimensional indices, so e.g.an ll_mat object would work. The optional parameter ω , which defaults to 1.0, is the weight parameter. The optional steps parameter (defaults to 1) specifies the number of iteration steps.

ssor(A, omega=1.0, steps=1)

Creates a ssor object, representing the SSOR preconditioner. The parameter $\bf A$ is the system matrix used for the SSOR iteration. The matrix $\bf A$ has to be an object of type sss_mat. The optional parameter ω , which defaults to 1.0, is the relaxation parameter. The optional steps parameter (defaults to 1) specifies the number of iteration steps.

4.1.2 jacobi and ssor Objects

Both jacobi and ssor objects provide the shape attribute and the precon method, that every preconditioner object in the PySparse framework must implement. Apart from that, there is nothing noteworthy to say about these objects.

4.1.3 Example: Diagonal Preconditioner

The diagonal preconditioner is just a special case of the Jacobi preconditioner, with omega=1.0 and steps=1, which happen to be the default values of these parameters.

It is however easy to implement the diagonal preconditioner using a Python class:

```
class diag_prec:
    def __init__(self, A):
        self.shape = A.shape
        n = self.shape[0]
        self.dinv = numpy.empty(n)
        for i in xrange(n):
            self.dinv[i] = 1.0 / A[i,i]
    def precon(self, x, y):
        numpy.multiply(x, self.dinv, y)

So:
>>> D1 = precon.jacobi(A, 1.0, 1)
and:
>>> D2 = diag_prec(A)
```

yield functionally equivalent preconditioners. D1 is probably faster than D2, because it is fully implemented in C.

ITERATIVE SOLVERS

5.1 The itsolvers Module

The itsolvers module provides a set of iterative methods for solving linear systems of equations.

The iterative methods are callable like ordinary Python functions. All these functions expect the same parameter list, and all function return values also follow a common standard.

Any user-defined iterative solvers should also follow these conventions, since other PySparse modules rely on them (e.g. the jdsym module; see *Eigenvalue Solver*).

Let's illustrate the calling conventions, using the PCG method.

info, iter, relres = pcg(A, b, x, tol, maxit[, K])()

Solve a linear system $\mathbb{A} \times \mathbb{A} = \mathbb{b}$ with the preconditioned conjugate gradient algorithm. Overwrite \times with the best estimate of the solution found.

Parameters

- A The coefficient matrix of the linear system of equations. A must provide the shape attribute and the matvec and matvec_transp methods for multiplying with a vector.
- **b** The right-hand-side of the linear system as a rank-1 NumPy array.
- ${\bf x}$ A rank-1 NumPy array. Upon entry, ${\bf x}$ holds the initial guess. On exit, ${\bf x}$ holds an approximate solution of the linear system.
- **tol** A float value representing the requested error tolerance. The exact meaning of this parameter depends on the actual iterative solver.

maxit An integer that specifies the maximum number of iterations to be executed.

K A preconditioner object that supplies the shape attribute and the precon method.

Returns

- **info** an integer that contains the exit status of the iterative solver. info >= 0 indicates, that x holds an acceptable solution, and info < 0 indicates an error condition. info has one of the following values:
- +2. iteration converged, residual is as small as seems reasonable on this machine,
- +1. iteration converged, b = 0, so the exact solution is x = 0.
- +0. iteration converged, relative error appears to be less than tol.
- -1. iteration did not converge, maximum number of iterations was reached.
- -2. iteration did not converge, the system involving the preconditioner was ill-conditioned.
- -3. iteration did not converge, an inner product of the form $\mathbf{x}^T \mathbf{K}^{-1} \mathbf{x}$ was not positive, so the preconditioning matrix \mathbf{K} does not appear to be positive definite.
- -4. iteration did not converge, the matrix A appears to be very ill-conditioned.
- -5. iteration did not converge, the method stagnated.
- -6. iteration did not converge, a scalar quantity became too small or too large to continue computing.

iter the number of iterations performed.

relres the relative residual at the approximate solution computed by the iterative method. What this actually is depends on the actual iterative method used.

The iterative solvers may accept additional parameters, which are passed as keyword arguments.

Note that not all iterative solvers check for all above error conditions.

5.1.1 itsolvers Module Functions

The module functions defined in the precon module implement various iterative methods (PCG, MINRES, QMRS and CGS). The parameters and return values conform to the conventions described above.

5.1.2 Example: Solving the Poisson System

Let's solve the Poisson system

$$\mathbf{L}\mathbf{x} = \mathbf{1},\tag{5.1}$$

using the PCG method. L is the 2D Poisson matrix, introduced in *Low-Level Sparse Matrix Types*, and 1 is a vector with all entries equal to one.

The Python solution for this task looks as follows:

```
from pysparse import spmatrix, precon, itsolvers
import numpy
n = 300
L = poisson2d_sym_blk(n)
b = numpy.ones(n*n)
x = numpy.empty(n*n)
info, iter, relres = itsolvers.pcg(L.to_sss(), b, x, 1e-12, 2000)
```

The code makes use of the Python function poisson2d_sym_blk, which was defined in *Low-Level Sparse Matrix Types*.

Incorporating e.g. a SSOR preconditioner is straightforward:

```
from pysparse import spmatrix, precon, itsolvers
import numpy
n = 300
L = poisson2d_sym_blk(n)
b = numpy.ones(n*n)
x = numpy.empty(n*n)
S = L.to_sss()
```

```
Kssor = precon.ssor(S)
info, iter, relres = itsolvers.pcg(S, b, x, 1e-12, 2000, Kssor)
```

The Matlab solution (without preconditioner) may look as follows:

```
n = 300;
L = poisson2d_kron(n);
[x,flag,relres,iter] = pcg(L, ones(n*n,1), 1e-12, 2000, ...
[], [], zeros(n*n,1));
```

5.1.3 Performance comparison with Matlab and native C

Todo

Update the timings below.

Warning: The timings below are Roman's old benchmarks. I don't know on which machine they were run. We should update them.

To evaluate the performance of the Python implementation we solve the 2D Poisson system (5.1) using the PCG method. The Python timings are compared with results of a Matlab and a native C implementation.

The native C and the Python implementation use the same core algorithms for PCG method and the matrix-vector multiplication. On the other hand, C reads the matrix from an external file instead of building it on the fly. In contrast to the Python implementation, the native C version does not suffer from the overhead generated by the runtime argument parsing and calling overhead.

Table 5.1: Performance comparison of Python, Matlab and native C implementations to solve the linear system (1) without preconditioning. The execution times are given in seconds. Assembly is the time for constructing the matrix (or reading it from a file in the case of native C). Solve is the time spent in the PCG solver. Total is the sum of Assembly and Solve. Matlab version 6.0 Release 12 was used for these timings.

Function	Size	Assembly	Solve	Total
Python	Python n=100		1.12	1.15
	n=300	0.21	49.65	49.86
	n=500	0.62	299.39	300.01
Native C	n=100	0.30	0.96	1.26
	n=300	3.14	48.38	51.52
	n=500	10.86	288.67	299.53
Matlab	n=100	0.21	8.85	9.06
	n=300	2.05	387.26	389.31
	n=500	6.23	1905.67	1911.8

This table shows the execution times for the Python, the Matlab and the native C implementation for solving the linear system (5.1). Matlab is not only slower when building the matrix, also the matrix-vector multiplication seems to be implemented inefficiently. Considering *Solve*, the performance of Python and native C is comparable. The Python overhead is under a factor of 4.

DIRECT SOLVERS

6.1 The Low-Level C Modules

6.1.1 The superlu Module

The superlu module interfaces the SuperLU library to make it usable by Python code. SuperLU is a software package written in C, that is able to compute an LU-factorisation of a general non-symmetric sparse matrix with partial pivoting.

The superlu module exports a single function, called factorize.

```
factorize(A, **kwargs)
```

The factorize function computes an LU-factorisation of the matrix A.

Parameters A A csr_mat object that represents the matrix to be factorized.

Keywords diag_pivot_thresh the partial pivoting threshold, in the interval [0,1]. diag_pivot_thresh=0 corresponds to no pivoting. diag_pivot_thresh=1 corresponds to partial pivoting (default: 1.0).

drop_tol the drop tolerance, in the interval [0,1]. drop_tol=0 corresponds to the exact factorization (default: 0.0).

relax the degree of relaxing supernodes (default: 1).

panel_size the maximum number of columns that form a panel (default: 10).

permc spec the matrix ordering used to control sparsity of the factors:

- 1. natural ordering
- 2. MMD applied to the structure of $\mathbf{A}^T \mathbf{A}$
- 3. MMD applied to the structure of $A^T + A$
- 4. COLAMD, approximate minimum degree column ordering (default: 2).

Return type an object of type superlu_context. This object encapsulates the L and U factors of A (see below).

Note: The drop_tol has no effect in SuperLU version 2.0 and below. In SuperLU version 3.0 and above, the default value of permc_spec is 3.

superlu_context Object Attributes and Methods

class superlu_context()

An abstract encapsulation of the LU factorization of a matrix by SuperLU.

shape

A 2-tuple describing the dimension of the matrix factorized. It is equal to A. shape.

nnz

The nnz attribute holds the total number of nonzero entries stored in both the L and U factors.

```
solve (b, x, trans)
```

The solve method accepts two rank-1 NumPy arrays b and x of appropriate size and assigns the solution of the linear system $\mathbf{A}\mathbf{x} = \mathbf{b}$ to x. If the optional parameter trans is set to the string 'T', the transposed system $\mathbf{A}^T\mathbf{x} = \mathbf{b}$ is solved instead.

Example: Solving a 2D Poisson System with SuperLU

Let's now solve the 2D Poisson system Ax = 1 using an LU factorization. Here, A is the 2D Poisson matrix, introduced in Low-Level Sparse Matrix Types and 1 is a vector with all entries equal to one.

The Python solution for this task looks as follows:

```
from pysparse import spmatrix, superlu
import numpy
n = 100
A = poisson2d_sym_blk(n)
b = numpy.ones(n*n)
x = numpy.empty(n*n)
LU = superlu.factorize(A.to_csr(), diag_pivot_thresh=0.0)
LU.solve(b, x)
```

The code makes use of the Python function poisson2d_sym_blk(), which was defined in *Low-Level Sparse Matrix Types*.

Example: An Incomplete LU Factorization Preconditioner

Warning: SuperLU 3.0 and above accept a drop_tol argument although the source files mention that incomplete factorization is *not implemented*. Therefore, changing drop_tol has no effect on the factorization at the moment and we must wait for it to be implemented. In the meantime, we can still demonstrate in this section how to implement an incomplete factorization preconditioner in Pysparse, even though in the present situation, it will be a *complete* factorization preconditioner!

Versions of SuperLU above 3.0 accept the drop_tol argument that allows the computation of incomplete factors, realizing a tradeoff between computational cost and factor density. The following example show how to use an incomplete LU factorization as a preconditioner in any of the iterative methods of the itsolvers module:

```
from pysparse import poisson, superlu, itsolvers
import numpy

class ILU_Precon:
    """
    A preconditioner based on an
    incomplete LU factorization.

Input: A matrix in CSR format.
    Keyword argument: Drop tolerance.
    """

def __init__(self, A, drop=1.0e-3):
    self.LU = superlu.factorize(A, drop_tol=drop)
    self.shape = self.LU.shape
```

```
def precon(self, x, y):
    self.LU.solve(x,y)

n = 300
A = poisson.poisson2d_sym_blk(n).to_csr()  # Convert right away
b = numpy.ones(n*n)
x = numpy.empty(n*n)

K = ILU_Precon(A)
info, niter, relres = itsolvers.pcg(A, b, x, 1e-12, 2000, K)
```

Note: Note that the 2D Poisson matrix is symmetric and positive definite, although barely. Indeed its smallest eigenvalue is $2(1-\cos(\pi/(n+1))) \approx (\pi/(n+1))^2$. Therefore, a Cholesky factorization would be more appropriate. In the future, we intend to interface the Cholmod library.

6.1.2 The umfpack Module

The umfpack module interfaces the UMFPACK library to make it usable by Python code. UMFPACK is a software package written in C, that is able to compute an LU factorization of a general non-symmetric sparse matrix with partial pivoting.

Note: The major difference with the superlu modules is that umfpack receives a matrix in ll_mat format instead of *csr_mat* format.

The umfpack module exports a single function, called factorize.

factorize(A, **kwargs)

The factorize function computes an LU-factorisation of the matrix A.

Parameters A A 11 mat object that represents the matrix to be factorized.

Keywords

strategy Pivoting strategy. Possible values are:

- "UMFPACK STRATEGY AUTO"
- "UMFPACK_STRATEGY_UNSYMMETRIC"
- "UMFPACK_STRATEGY_SYMMETRIC"
- "UMFPACK_STRATEGY_2BY2"

tol2by2 Tolerance for the 2-by-2 strategy.

scale Scaling used during factorization. Possible values are:

- "UMFPACK SCALE NONE"
- "UMFPACK_SCALE_SUM"
- "UMFPACK_SCALE_MAX"

tolpivot Relative pivot tolerance for threshold partial pivoting with row interchanges.

tolsympivot If diagonal pivoting is attempted, this parameter controls when the diagonal is selected in a given pivot column.

Return type an object of type umfpack_context. This object encapsulates the L and U factors of A (see below).

umfpack_context Object Attributes and Methods

class umfpack_context()

An abstract encapsulation of the LU factorization of a matrix by UMFPACK.

shape

A 2-tuple describing the dimension of the matrix factorized. It is equal to A. shape.

nnz

The nnz attribute holds the total number of nonzero entries stored in the input matrix. It is equal to A.nnz. To obtain the number of nonzero element in the factors, see lunz().

```
solve (b, x, method, irsteps)
```

Parameters b The right-hand side of the system Ax = b as a Numpy array.

x A Numpy array to hold the solution of $\mathbf{A}x = b$.

method (optional) Different systems may be solved by setting the method argument appropriately. See the documentation of the pysparseUmfpackSolver below for more details.

irsteps (optional) The number of iterative refinement steps to perform.

lu()

Return the factors, permutation and scaling information. See the documentation of the pysparseUmfpackSolver below for more details.

lunz()

Return the number of nonzeros in factors, i.e., in L + U.

Example: Solving a 2D Poisson System with UMFPACK

We now solve again the 2D Poisson system Ax = 1 using an LU factorization. Here, A is the 2D Poisson matrix, introduced in Low-Level Sparse Matrix Types and 1 is a vector with all entries equal to one.

The Python solution using UMFPACK looks as follows:

```
from pysparse import spmatrix, umfpack
import numpy
n = 100
A = poisson2d_sym_blk(n)
b = numpy.ones(n*n)
x = numpy.empty(n*n)
LU = umfpack.factorize(A, strategy="UMFPACK_STRATEGY_SYMMETRIC")
LU.solve(b, x)
```

The code makes use of the Python function poisson2d_sym_blk(), which was defined in *Low-Level Sparse Matrix Types*.

6.2 Higher-Level Python Interfaces

This section anticipates on *Higher-Level Sparse Matrix Classes* and shows usage of higher-level interfaces to the LU factorization packages.

6.2.1 The Abstract directSolver Module

A framework for solving sparse linear systems of equations using a direct factorization.

```
class PysparseDirectSolver (A, **kwargs)
```

PysparseDirectSolver is a generic class and should be subclassed.

```
solve (b, **kwargs)
```

6.2.2 The pysparseSuperLU Module: A Higher-Level SuperLU Interface

A framework for solving sparse linear systems of equations using an LU factorization, by means of the supernodal sparse LU factorization package SuperLU ([DEGLL99], [DGL99], [LD03]).

This package is appropriate for factorizing sparse square unsymmetric or rectangular matrices.

See [SLU] for more information.

References:

class PysparseSuperLUSolver (A, **kwargs)

Bases: directSolver.PysparseDirectSolver

PysparseSuperLUSolver is a wrapper class around the SuperLu library for the factorization of full-rank n-by-m matrices. Only matrices with real coefficients are currently supported.

Parameters A The matrix to be factorized, supplied as a PysparseMatrix instance.

Keywords

symmetric a boolean indicating that the user wishes to use symmetric mode. In symmetric mode, permc_spec=2 must be chosen and diag_pivot_thresh must be small, e.g., 0.0 or 0.1. Since the value of diag_pivot_thresh is up to the user, setting symmetric to True does *not* automatically set permc_spec and diag_pivot_thresh to appropriate values.

diag_pivot_thresh a float value between 0 and 1 representing the threshold for partial pivoting (0 = no pivoting, 1 = always perform partial pivoting). Default: 1.0.

drop_tol the value of a drop tolerance, between 0 and 1, if an incomplete factorization is desired (0 = exact factorization). This keyword does not exist if using SuperLU version 2.0 and below. In more recent version of SuperLU, the keyword is accepted but has no effect. Default: 0.0

relax an integer controlling the degree of relaxing supernodes. Default: 1.

panel_size an integer specifying the maximum number of columns to form a panel. Default: 10.

permc_spec an integer specifying the ordering strategy used during the factorization.

- 1. natural ordering,
- 2. MMD applied to the structure of $\mathbf{A}^T \mathbf{A}$
- 3. MMD applied to the structure of $A^T + A$
- 4. COLAMD.

Default: 2.

LU

A superlu_context object encapsulating the factorization.

sol

The solution of the linear system after a call to solve ().

factorizationTime

The CPU time to perform the factorization.

solutionTime

The CPU time to perform the forward and backward sweeps.

lunz

The number of nonzero elements in the factors L and U together after a call to fetch_lunz().

fetch_factors()

Not yet available.

fetch_lunz()

Retrieve the number of nonzeros in the factors L and U together. The result is stored in the member lunz of the class instance.

```
solve (rhs, transpose=False)
```

Solve the linear system $A \times = rhs$, where A is the input matrix and rhs is a Numpy vector of appropriate dimension. The result is placed in the sol member of the class instance.

If the optional argument transpose is True, the transpose system $A^T x = rhs$ is solved.

6.2.3 Example: The 2D Poisson System with SuperLU

The solution of a 2D Poisson system with PysparseSuperLUSolver may look like this:

```
from pysparse.pysparseMatrix import PysparseMatrix
from pysparse.pysparseSuperLU import PysparseSuperLUSolver
from poisson_vec import poisson2d_sym_blk_vec
from numpy import ones
from numpy.linalg import norm
n = 200
A = PysparseMatrix( matrix=poisson2d_sym_blk_vec(n) )
x_exact = ones(n*n)/n
b = A * x_exact
LU = PysparseSuperLUSolver(A)
LU.solve(b)
print 'Factorization time: ', LU.factorizationTime
print 'Solution time: ', LU.solutionTime
print 'Error: ', norm(LU.sol - x_exact)/norm(x_exact)
The above script produces the output:
Factorization time: 0.494116
Solution time: 0.017096
```

Note that this example uses the vectorized Poisson constructors of Low-Level Sparse Matrix Types.

6.2.4 The pysparseUmfpack Module: A Higher-Level UMFPACK Interface

A framework for solving sparse linear systems of equations using an LU factorization, by means of the unsymmetric multifrontal sparse LU factorization package UMFPACK ([D04a], [D04b], [DD97]).

This package is appropriate for factorizing sparse square unsymmetric or rectangular matrices.

See [UMF] for more information.

Error: 2.099685128150953e-14

References:

```
class PysparseUmfpackSolver(A, **kwargs)
```

```
Bases: directSolver.PysparseDirectSolver
```

PysparseUmfpackSolver is a wrapper class around the UMFPACK library for the factorization of full-rank n-by-m matrices. Only matrices with real coefficients are currently supported.

```
Parameters A A PysparseMatrix instance representing the matrix to be factorized.

Keywords strategy string that specifies what kind of ordering and pivoting strategy UMF-PACK should use. Valid values are 'auto', 'unsymmetric', 'symmetric' and '2by2'.

Default: 'auto'

tol2by2 tolerance for the 2 by 2 strategy. Default: 0.1
```

scale string that specifies the scaling UMFPACK should use. Valid values are 'none', 'sum', and 'max'. Default: 'sum'.

tolpivot relative pivot tolerance for threshold partial pivoting with row interchanges. Default: 0.1

tolsympivot if diagonal pivoting is attempted, this parameter is used to control when the diagonal is selected in a given pivot column. Default: 0.0

LU

An umfpack_context object encapsulating the factorization.

sol

The solution of the linear system after a call to solve ().

L

The L factor of the input matrix.

U

The U factor of the input matrix.

Р

The row permutation used for the factorization.

Q

The column permutation used for the factorization.

R

The row scaling used during the factorization. See the documentation of fetch_factors().

factorizationTime

The CPU time to perform the factorization.

solutionTime

The CPU time to perform the forward and backward sweeps.

do_recip

Nature of the row scaling. See fetch_factors().

lnz

The number of nonzero elements in the factor L.

unz

The number of nonzero elements in the factor U from which the diagonal was removed.

nz_udiag

The number of nonzero elements on the diagonal of the factor U.

fetch_factors()

Retrieve the L and U factors of the input matrix along with the permutation matrices P and Q and the row scaling matrix R such that

$$PRAQ = LU.$$

The matrices P, R and Q are stored as Numpy arrays. L and U are stored as PysparseMatrix instances and are lower triangular and upper triangular, respectively.

R is a row-scaling diagonal matrix such that

- •the i-th row of A has been divided by R[i] if do_recip = True,
- •the i-th row of A has been multiplied by R[i] if do_recip = False.

fetch_lunz()

Retrieve the number of nonzeros in the factors. The results are stored in the members lnz, unz and nz_udiag of the class instance.

```
solve (rhs, **kwargs)
```

Solve the linear system $A \times = \text{rhs}$. The result is placed in the sol member of the class instance.

Parameters rhs a Numpy vector of appropriate dimension.

Keywords method specifies the type of system being solved:

```
\mathbf{A}x = b (default)
"UMFPACK_A"
                            \mathbf{A}^T x = b
"UMFPACK At"
                            \mathbf{P}^T \mathbf{L} x = b
"UMFPACK Pt L"
"UMFPACK L"
                            \mathbf{L}x = b
"UMFPACK_Lt_P"
                            \mathbf{L}^T \mathbf{P} x = b
                            \mathbf{L}^T x = b
"UMFPACK_Lt"
                            \mathbf{U}\mathbf{Q}^Tx = b
"UMFPACK_U_Qt"
"UMFPACK U"
                            \mathbf{U}x = b
                            \mathbf{Q}\mathbf{U}^Tx = b
"UMFPACK_Q_Ut"
                            \mathbf{U}^T x = b
"UMFPACK_Ut"
```

irsteps number of iterative refinement steps to attempt. Default: 2

6.2.5 Example: The 2D Poisson System with UMFPACK

The solution of a 2D Poisson system with PysparseUmfpackSolver may look like this:

```
from poisson_vec import poisson2d_sym_blk_vec
from numpy import ones
from numpy.linalg import norm

n = 200
A = PysparseMatrix( matrix=poisson2d_sym_blk_vec(n) )
x_exact = ones(n*n)/n
b = A * x_exact
LU = PysparseUmfpackSolver(A)
LU.solve(b)
print 'Factorization time: ', LU.factorizationTime
print 'Solution time: ', LU.solutionTime
print 'Error: ', norm(LU.sol - x_exact)/norm(x_exact)
```

This script produces the output:

Factorization time: 0.520043 Solution time: 0.031086 Error: 1.10998989668e-15

EIGENVALUE SOLVER

7.1 The jdsym Module

The jdsym module provides an implementation of the JDSYM algorithm, that is conveniently callable from Python. JDSYM is an eigenvalue solver to compute eigenpairs of a generalised matrix eigenvalue problem of the form

$$\mathbf{A}\mathbf{x} = \lambda \mathbf{M}\mathbf{x} \tag{7.1}$$

or a standard eigenvalue problem of the form

$$\mathbf{A}\mathbf{x} = \lambda \mathbf{x} \tag{7.2}$$

where **A** is symmetric and **M** is symmetric positive definite.

The module exports a single function:

jdsym (A, M, K, kmax, tau, jdtol, itmax, linsolver, **kwargs)

Implements Jacobi-Davidson iterative method to identify a given number of eigenvalues near a target value.

Parameters A the matrix A in (7.1) or (7.2). A must provide the shape attribute and the matvec and matvec_transp methods.

M the matrix M in (7.1). M must provide the shape attribute and the matvec and matvec_transp methods. If the standard eigenvalue problem (7.2) is to be solved, M should be set to None.

K a preconditioner object that supplies the shape attribute and the precon method. If no preconditioner is to used, then the None value can be passed for this parameter.

kmax the number of eigenpairs to be computed.

tau the target value τ . Eigenvalues in the vicinity of τ will be computed.

jdtol the convergence tolerance for eigenpairs (λ, \mathbf{x}) . The converged eigenpairs have a residual $\|\mathbf{A}\mathbf{x} - \lambda \mathbf{M}\mathbf{x}\|_2$ less than jdtol.

itmax an integer that specifies the maximum number of Jacobi-Davidson iterations to perform.

linsolver a function that implements an iterative method for solving linear systems of equations. The function linsolver is required to conform to the standards mentioned in *Iterative Solvers*.

Keywords jmax the maximum dimension of the search subspace (default: 25).

jmin the dimension of the search subspace after a restart (default: 10).

blksize the block size used in the JDSYM algorithm (default: 1).

blkwise is an integer that affects the convergence criterion if blksize is larger than 1 (default: 0).

V0 a NumPy array of rank one or two. It specifies the initial search subspace (default: a randomly generated initial search subspace).

optype is an integer specifying the operator type used in the correction equation. If optype=1, the non-symmetric version is used. If optype=2, the symmetric version is used (default: 2).

linitmax the maximum number of steps taken in the inner iteration (iterative linear solver) (default: 200).

eps_tr the tracking parameter (default: 1.0e-3).

toldecay is a float value that influences the dynamic adaptation of the stopping criterion of the inner iteration (default: 1.5).

clvl verbosity level. The higher the clvl parameter, the more output is sent to the standard output. clvl=0 produces no output (default: 0).

strategy is an integer specifying shifting and sorting strategy of JDSYM. strategy=0 enables the default JDSYM algorithm. strategy=1 enables JDSYM to avoid convergence to eigenvalues smaller than τ (default: 0).

projector is used to keep the search subspace and the eigenvectors in a certain subspace. The parameter projector can be any Python object that has a shape attribute and a project method. The project method takes a vector (a rank-1 NumPy array) as its sole argument and projects that vector in-place. This parameter can be used to implement the DIRPROJ and SAUG methods (default: no projection).

Returns

kconv the number of converged eigenpairs.

lambda a rank-1 NumPy array containing the converged eigenvalues.

Q a rank-2 NumPy array containing the converged eigenvectors. The i-th eigenvector is accessed by Q[:,i].

it an integer indicating the number of Jacobi-Davidson steps (outer iteration steps) performed.

7.1.1 Example: Maxwell Problem

Todo

Update the timings below.

Warning: The timings below are Roman's old benchmarks. We should run them again.

The following code illustrates the use of the jdsym module. Two matrices $\bf A$ and $\bf M$ are read from files. A Jacobi preconditioner from $\bf A-\tau M$ is built. Then the JDSYM eigensolver is called, calculating 5 eigenvalues near 25.0 and the associated eigenvalues to an accuracy of 10^{-10} . We set strategy=1 to avoid convergence to the high-dimensional null space of $(\bf A, M)$:

```
from pysparse import spmatrix, itsolvers, jdsym, precon
A = spmatrix.ll_mat_from_mtx('edge6x3x5_A.mtx')
M = spmatrix.ll_mat_from_mtx('edge6x3x5_B.mtx')
tau = 25.0

Atau = A.copy()
Atau.shift(-tau, M)
K = precon.jacobi(Atau)
```

This code takes 33.71 seconds to compute the five eigenpairs. A native C version, using the same computational kernels, takes 35.64 for the same task. We expected the Python version to be slower due to the overhead generated when calling the matrix-vector multiplication and the preconditioner, but surprisingly the Python code was even a bit faster.

New in version 1.0.1.

HIGHER-LEVEL SPARSE MATRIX CLASSES

8.1 The pysparseMatrix module

class PysparseMatrix (**kwargs)

Bases: sparseMatrix.SparseMatrix

A PysparseMatrix is a class wrapper for the pysparse spmatrix sparse matrix type. This class facilitates matrix populating and allows intuitive operations on sparse matrices and vectors.

Currently accepted keywords include

nrow	The number of rows of the matrix
ncol	The number of columns of the matrix
size	The common number of rows and columns, for a square matrix
bandwidth	The bandwidth (if creating a band matrix)
matrix	The starting <i>spmatrix</i> if there is one
sizeHint	A guess on the number of nonzero elements of the matrix
symmetric	A boolean indicating whether the matrix is symmetric.

addAt (vector, id1, id2)

Add elements of vector to the positions in the matrix corresponding to (id1,id2)

```
>>> L = PysparseMatrix(size = 3)
>>> L.put([3.,10.,numpy.pi,2.5], [0,0,1,2], [2,1,1,0])
>>> L.addAt((1.73,2.2,8.4,3.9,1.23), (1,2,0,0,1), (2,2,0,0,2))
>>> print L
12.300000 10.000000 3.000000
--- 3.141593 2.960000
2.500000 --- 2.200000
```

addAtDiagonal (vector)

Add the components of vector vector to the diagonal elements of the matrix.

copy()

Returns a (deep) copy of a sparse matrix

exportMmf (filename)

Exports the matrix to a Matrix Market file of the given filename.

find()

Returns three Numpy arrays to describe the sparsity pattern of self in so-called coordinate (or triplet) format:

```
>>> L = PysparseMatrix(size = 3)
    >>> L.put([3.,10.,numpy.pi,2.5], [0,0,1,2], [2,1,1,0])
    >>> (val, irow, jcol) = L.find()
    >>> val
    array([ 10. , 3. , 3.14159265, 2.5
                                                                      1)
    >>> irow
    array([0, 0, 1, 2])
    >>> jcol
    array([1, 2, 1, 0])
getMatrix()
    Returns the underlying ll mat sparse matrix of self
    Returns the number of nonzero elements of self
getNumpyArray()
    Convert a sparse matrix to a dense Numpy matrix.
getShape()
    Returns the shape (nrow, ncol) of a sparse matrix
isSymmetric()
    Returns True is self is a symmetric matrix or False otherwise
matvec(x)
    This method is required for scipy solvers.
put (value, id1, id2)
    Put elements of value at positions of the matrix corresponding to (id1, id2)
    >>> L = PysparseMatrix(size = 3)
    >>> L.put( [3.,10.,numpy.pi,2.5], [0,0,1,2], [2,1,1,0] )
    >>> print L
        --- 10.000000 3.000000
               3.141593 ---
     2.500000
                 ---
    >>> L.put(2*numpy.pi, range(3), range(3))
    >>> print L
     6.283185 10.000000 3.000000
        --- 6.283185
                           ___
     2.500000
                            6.283185
    If value is a scalar, it has the same effect as the vector of appropriate length with all values equal to
    range (ncol).
```

value. If idl is omitted, it is replaced with range (nrow). If idl is omitted, it is replaced with

putDiagonal (vector)

Put elements of vector along diagonal of matrix

```
>>> L = PysparseMatrix(size = 3)
>>> L.putDiagonal([3.,10.,numpy.pi])
>>> print L
3.000000
   --- 10.000000 ---
                    3.141593
            ___
>>> L.putDiagonal([10.,3.])
>>> print L
10.000000
   --- 3.000000
                    3.141593
>>> L.putDiagonal(2.7182)
```

```
>>> print L
2.718200 --- ---
--- 2.718200 ---
--- 2.718200 ---
2.718200
```

take (*id1*, *id2*)

Extract elements at positions (irow[i], jcol[i]) and place them in the array val. In other words:

```
for i in range(len(val)): val[i] = A[irow[i], jcol[i]]
```

```
takeDiagonal()
```

Extract the diagonal of a matrix and place it in a Numpy array.

8.1.1 Creating an Identity Matrix

```
class PysparseIdentityMatrix (size)
```

Bases: pysparseMatrix.PysparseMatrix

Represents a sparse identity matrix for pysparse.

```
>>> print PysparseIdentityMatrix(size = 3)
1.000000 --- ---
--- 1.000000 ---
1.000000
```

8.1.2 Creating Sparse Matrices from Diagonals

class PysparseSpDiagsMatrix (size, vals, pos, **kwargs)

Bases: pysparseMatrix.PysparseMatrix

Represents a banded matrix with specified diagonals.

Example: Create a tridiagonal matrix with 1's on the diagonal, 2's above the diagonal, and -2's below the diagonal.

Note that since the pos[k]-th diagonal has size-|pos[k]| elements, only that many first elements of vals[k] will be inserted.

If the banded matrix is requested to be symmetric, elements above the main diagonal are not inserted.

8.1.3 Fancy Indexing

Fancy indexing carries over to PysparseMatrix objects and is used exactly in the same way as with ll_mat objects. Refer to Section *Low-Level Sparse Matrix Types* for details.

OTHER SPARSE MATRIX PACKAGES FOR PYTHON

- Scipy provides a sparse matrix module featuring several storage formats.
- CVXOPT is a package for convex optimization and allows creation and manipulation of sparse matrices in compressed sparse row format.

Pysparse Documentation, Release 1.0.2				
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Parts of this documentation are scavenged from Roman Geus' original Pysparse website.

Some pages of this documentation display equations via the jsMath package. They should look reasonably good with most setups but the best rendering is obtained by installing the TeX fonts. Please refer to http://www.math.union.edu/~dpvc/jsMath/users/welcome.html.

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CHAPTER

ELEVEN

TODO LIST

Todo

Update the timings below.

(The original entry is located in itsolvers.rst, line 158 and can be found here.)

Todo

Update the timings below.

(The original entry is located in jdsym.rst, line 106 and can be found here.)

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