

Package ‘rmumps’

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Type Package

Title Wrapper for MUMPS Library

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Description Some basic features of 'MUMPS' (Multifrontal Massively Parallel sparse direct Solver) are wrapped in a class whose methods can be used for sequentially solving a sparse linear system (symmetric or not) with one or many right hand sides (dense or sparse). There is a possibility to do separately symbolic analysis, LU (or LDL^t) factorization and system solving. Third part ordering libraries are included and can be used: 'PORD', 'METIS', 'SCOTCH'. 'MUMPS' method was first described in Amestoy et al. (2001) <doi:10.1137/S0895479899358194> and Amestoy et al. (2006) <doi:10.1016/j.parco.2005.07.004>.

License GPL (>= 2)

Depends methods

Imports Rcpp (>= 0.12.0)

LinkingTo Rcpp

SystemRequirements GNU Make

NeedsCompilation yes

Biarch yes

Suggests testthat, Matrix, slam

BugReports <https://github.com/sgsokol/rmumps/issues>

URL <http://www.mumps-solver.org/>, <https://github.com/sgsokol/rmumps/>

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rmumps-package	<i>Rcpp port of MUMPS library for LU or LDL^t factorization of sparse matrices</i>
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Description

Creates a MUMPS compatible object storing a sparse matrix. Gives a possibility to do separately symbolic analysis, factorization and system solving.

Details

Create a new Rmumps object with `A <- Rmumps$new(aspase)` then solve a linear system with one or many right hand sides `x <- solve(A, b)`. Cf. [Rmumps](#)

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References

MUMPS official site <http://mumps.enseeiht.fr>
Sokol S (2025). `_Rmumps: Rcpp port of MUMPS_`. rmumps package version 5.2.1-35, <URL: <http://CRAN.R-project.org/package=rmumps>>.

Examples

```
## Not run:
A <- Rmumps$new(aspase)
x <- solve(A, b)

## End(Not run)
```

Rcpp_Rmumps-class

Rcpp Exported Class Wrapping MUMPS library

Description

This class can be used for storing sparse matrix and solving corresponding linear system with one or many right hand sides. There is a possibility to do separately symbolic analysis, LU factorization and system solving.

Fields

sym: integer (read only), 0=non symmetric matrix, 1=symmetric with pivots on diagonal or 2=general symmetric

copy: logical, copy or not rhs and matrix values

mrhs: numeric matrix, multiple rhs (always overwritten with solution)

rhs: numeric vector, single rhs (always overwritten with solution)

Methods

new(asp, sym=0, copy=TRUE): constructor from Matrix::dgTMatrix class (or from convertible to it) and slam::simple_triplet_matrix class

new(i, j, x, n, copy=TRUE): constructor from triade rows, cols, vals

symbolic(): do symbolic analysis (stored internally)

numeric(): do LU or LDL^t factorization (stored internally)

solve(b): solve single rhs (if b is a vector) or multiple rhs if b is a matrix (can be dense or sparse).
Return the solution(s).

solvect(b): same as solve() but solves with transposed matrix

det(): Return determinant of the matrix

inv(): Return inverse of the matrix

set_mat_data(x): updates matrix entries (x must be in the same order as in previous calls)

set_icntl(iv, ii): set ICNTL parameter vector

get_icntl(): get ICNTL parameter vector

set_cntl(v, iv): set CNTL parameter vector

get_cntl(): get CNTL parameter vector

get_infos(): get a named list of information vectors: info, rinfo, infog and rinfog

`dim()`: Return a dimension vector of the matrix
`nrow()`: Return a row number of the matrix
`ncol()`: Return a column number of the matrix
`print()`: Print summary information on the matrix
`show()`: Print summary information on the matrix
`set_keep()`: Set KEEP array elements (undocumented feature of MUMPS)
`get_keep()`: Get a copy of KEEP array elements (length=500)
`set_permutation(perm)`: Set permutation type which can impact storage and factorization performances. Parameter `perm` can take one of the following predefined integer values `RMUMPS_PERM_AMD`, `RMUMPS_PERM_AMF`, `RMUMPS_PERM_SCOTCH`, `RMUMPS_PERM_PORD`, `RMUMPS_PERM_METIS`, `RMUMPS_PERM_QAMD`. This method should be called once and before symbolic analysis of the matrix. If it is called afterward, a new symbolic and numeric factorization will be performed when one of other methods (e.g. `solve()`) will request them. In other words, previous symbolic and numeric factorizations are canceled by this method.
`get_permutation()`: get permutation type currently set in the object
`mumps_version()`: Return a string with MUMPS version used in `rmumps`

Note

When creating a symmetric matrix (`sym=1` or `sym=2`), the upper (or lower) part of the input matrix must be zeroed.

For meaning of entries in MUMPS vectors `cntl`, `icntl`, `info`, `rinfo`, `infog` and `rinfog` cf. original documentation of MUMPS project.

No need to call `symbolic()` and `numeric()` methods before a `solve()` call.

If in constructor, a parameter `copy` is set to `FALSE`, no rhs neither matrix copying is done. The solution is written "in place" thus overwriting rhs (watch out side effects)

For a detailed error diagnostic (e.g. when factorizing a singular matrix), use method `get_infos()` and cf. MUMPS documentation on the official MUMPS site).

Author(s)

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References

MUMPS official site <http://mumps.enseiht.fr>

Sokol S (2020). `_Rmumps`: Rcpp port of MUMPS. `rmumps` package version 5.2.1-X, <URL: <http://CRAN.R-project.org/package=rmumps>>.

Examples

```
## Not run:
# prepare random sparse matrix
library(Matrix)
library(rmumps)
```

```

n=2000
a=Matrix(0, n, n)
set.seed(7)
ij=sample(1:(n*n), 15*n)
a[ij]=runif(ij)
diag(a)=0
diag(a)=-rowSums(a)
a[1,1]=a[1,1]-1
am=Rmumps$new(a)
b=as.double(a%*(1:n)) # rhs for an exact solution vector 1:n
# following time includes symbolic analysis, LU factorization and system solving
system.time(x<-solve(am, b))
bb=2*b
# this second time should be much shorter
# as symbolic analysis and LU factorization are already done
system.time(xx<-solve(am, bb))
# compare to Matrix corresponding times
system.time(xm<-solve(a, b))
system.time(xxm<-solve(a, bb))
# compare to Matrix precision
range(x-1:n) # mumps
range(xm-1:n) # Matrix

# matrix inversion
system.time(aminv <- solve(am))
system.time(ainv <- solve(a)) # the same in Matrix

# symmetric matrix
asy=as(a+t(a), "symmetricMatrix")
bs=as.double(asy%*(1:n)) # rhs for 1:n solution
au=asy
# Here, we keep only diagonal and upper values of asy matrix.
# It could be also diagonal and lower values.
au[row(au)>col(au)]=0
ams=Rmumps$new(au, sym=1)
system.time(xs<-solve(ams, bs)) # rmumps
system.time(xsm<-solve(asy, bs)) # Matrix
# compare to Matrix precision
range(xs-1:n) # mumps
range(xsm-1:n) # Matrix

# clean up by hand to avoid possible interference between gc() and
# Rcpp object destructor after unloading this namespace
rm(am, ams)
gc()

## End(Not run)

```

Description

Integer constants defining permutation types and exported from rmumps are following:

- RMUMPS_PERM_AMD
- RMUMPS_PERM_AMF
- RMUMPS_PERM_SCOTCH
- RMUMPS_PERM_PORD
- RMUMPS_PERM_METIS
- RMUMPS_PERM_QAMD
- RMUMPS_PERM_AUTO

They are all regrouped in a named vector RMUMPS_PERM where names are items above and values are corresponding constants.

Examples

```
am=rmumps::Rmumps$new(slam::as.simple_triplet_matrix(diag(1:3)))
am$set_permutation(RMUMPS_PERM_SCOTCH)
am$solve(1:3)
```

Rmumps__del_ptr	<i>Delete via Pointer</i>
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Description

This is a C wrapper to Rmumps::~~Rmumps() destructor. Available in R too. In C++ code can be used as rmumps::Rmumps__del_ptr(pm)

Usage

```
Rmumps__del_ptr(pm)
```

Arguments

pm	pointer of type XPtr<Rmumps>, object to be deleted
----	--

Rmumps__get_permutation

Get Permutation Parameter

Description

This is a C wrapper to `Rmumps::get_permutation()` method. Available in R too. In C++ code can be used as `rmumps::Rmumps__get_permutation(pm)`

Usage

```
Rmumps__get_permutation(pm)
```

Arguments

pm	pointer of type <code>XPtr<Rmumps></code> , object having sparse matrix permuted according to some method.
----	--

Value

integer defining permutation method used before matrix decomposition.

Rmumps__ptr_ijv

Construct via Triplet Pointers

Description

This is a C wrapper to `Rmumps::Rmumps(i, j, v, n, nz, sym)` constructor. Available in R too. In C++ code can be used as `rmumps::Rmumps__ptr_ijv(pi, pj, pa, n, nz, sym)`

Usage

```
Rmumps__ptr_ijv(pi, pj, pa, n, nz, sym)
```

Arguments

pi	pointer of type <code>XPtr<int></code> , vector of i-indeces for sparse triplet
pj	pointer of type <code>XPtr<int></code> , vector of j-indeces for sparse triplet
pa	pointer of type <code>XPtr<double></code> , vector of values for sparse triplet
n	integer, size of the matrix (n x n)
nz	integer, number of non zeros in the matrix
sym	integer, 0 means general (non symmetric) matrix, 1 - symmetric with pivotes on the main diagonal, 2 - general symmetric (pivotes may be anywhere)

Value

pointer of type `XPtr<Rmumps>` pointing to newly created object. To avoid memory leakage, it is user's responsibility to call `Rmumps__del_ptr(pm)` in a due moment (where `pm` is the returned pointer).

`Rmumps__set_mat_ptr` *Set Matrix via Pointer*

Description

This is a C wrapper to `Rmumps::set_mat_ptr(a)` method. Available in R too. In C++ code can be used as `rmumps::Rmumps__set_mat_ptr(pm)`. Using this method invalidates previous numeric decomposition (but not symbolic one).

Usage

```
Rmumps__set_mat_ptr(pm, pa)
```

Arguments

<code>pm</code>	pointer of type <code>XPtr<Rmumps></code> , object having sparse matrix to be replaced with second parameter
<code>pa</code>	pointer of type <code>XPtr<double></code> , value vector from sparse triplet providing a new matrix. Structure of the new matrix must be identical to the old one. That's why there is no need to provide <code>i</code> and <code>j</code> for the new triplet.

`Rmumps__set_permutation`
Set Permutation Parameter

Description

This is a C wrapper to `Rmumps::set_permutation(permutation)` method. Available in R too. In C++ code can be used as `rmumps::Rmumps__set_permutation(pm, permutation)`

Usage

```
Rmumps__set_permutation(pm, permutation)
```

Arguments

<code>pm</code>	pointer of type <code>XPtr<Rmumps></code> , object having sparse matrix permuted according to a chosen method.
<code>permutation</code>	integer one of predefined constants (cf. RMUMPS_PERM). Setting a new permutation invalidates current symbolic and numeric matrix decompositions.

Rmumps__solveptr	<i>Solve via Pointer</i>
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Description

This is a C wrapper to `Rmumps::solveptr()` method. Available in R too. In C++ code can be used as `rmumps::Rmumps__solveptr(pobj, pb, lrhs, nrhs)`

Usage

```
Rmumps__solveptr(pobj, pb, lrhs, nrhs)
```

Arguments

pobj	pointer of type <code>XPtr<Rmumps></code> , object having sparse matrix
pb	pointer of type <code>XPtr<double></code> , vector or dense matrix of rhs
lrhs	integer, leading dimension in pb
nrhs	integer, number of rhs to solve.

Rmumps__triplet	<i>Explore via Triplet</i>
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Description

This is a C wrapper to `Rmumps::triplet()` method. Available in R too. In C++ code can be used as `rmumps::Rmumps__triplet(pm)`

Usage

```
Rmumps__triplet(pm)
```

Arguments

pm	pointer of type <code>XPtr<Rmumps></code> , object having sparse matrix to be explored
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Value

a list with sparse triplet described with fields `i`, `j`, `v`

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