# Package 'filematrix'

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filematrix-package filematrix-class fm.create

2 filematrix-package

Index 8

filematrix-package

File-backed numeric matrix.

# **Description**

File-Backed Matrix Class with Convenient Read and Write Access

#### **Details**

Interface for working with large matrices stored in files, not in computer memory. Supports multiple non-character data types (double, integer, logical and raw) of various sizes (e.g. 8 and 4 byte real values). Access to parts of the matrix is done by indexing (e.g. fm[,1]), exactly as with usual R matrices. Supports very large matrices. Tested on multi-terabyte matrices. Allows for more than 2^32 rows or columns. Allows for quick addition of extra columns to a filematrix. Cross-platform as the package has R code only.

A new file.matrix object can be created with fm. create and fm. create.from.matrix. Existing file.matrix files can be opened with fm. open.

Once a file.matrix is created or opened it can be accessed as a regular matrix object in R. All changes to file.matrix object are written to the data files without extra buffering.

#### Note

Due to lack of 64 bit integer data type in R, the package uses double values for calculation of indices. The precision of double data type is sufficient for indexing matrices up to 8,192 terabytes in size.

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# See Also

See fm. create and filematrix for reference.

Run browseVignettes("filematrix") for the list of vignettes.

filematrix-class 3

filematrix-class

Manipulating file matrices (class "filematrix")

# **Description**

filematrix is a class for working with very large matrices stored in files, not held in computer memory. It is intended as a simple, efficient solution to handling big numeric data (i.e., datasets larger than memory capacity) in R.

A new filematrix can be created with fm.create. It can be created from an existing R matrix with fm.create.from.matrix. A text file with a matrix can be scanned and converted into a filematrix with fm.create.from.text.file. An existing filematrix can be opened for read/write access with fm.open or loaded fully in memory with fm.load.

A filematrix can be handled as an ordinary matrix in R.

It can be read from and written to via usual indexing with possible omission of indices.

For example: fm[1:3,2:4] and fm[,2:4].

The values can also be accessed as a vector with single indexing.

For example: fm[3:7] and fm[4:7] = 1:4.

A whole filematrix can be read memory as an ordinary R matrix with as.matrix function or empty indexing fm[].

The dimensions of filematrix can be obtained via dim, nrow and ncol functions and modified with dim function.

For example: dim(fm) and dim(fm) = c(10, 100).

The number of elements in filematrix is returned by length function.

A filematrix can have row and column names. They can be accessed using the standard functions rownames, colnames, and dimnames.

A filematrix can be closed after use with close command. Note, however, that there is no risk of losing modifications to a filematrix if an object is not closed, as all changes are written to disk without delay.

#### Usage

```
## S3 method for class 'filematrix'
x[i,j]
## S3 replacement method for class 'filematrix'
x[i,j] <- value

## S4 method for signature 'filematrix'
as.matrix(x)

## S4 method for signature 'filematrix'
dim(x)
## S4 replacement method for signature 'filematrix'
dim(x) <- value</pre>
```

4 filematrix-class

```
## S4 method for signature 'filematrix'
length(x)

## S4 method for signature 'filematrix'
rownames(x)
## S4 replacement method for signature 'filematrix'
rownames(x) <- value

## S4 method for signature 'filematrix'
colnames(x)
## S4 replacement method for signature 'filematrix'
colnames(x) <- value

## S4 method for signature 'filematrix'
dimnames(x)
## S4 replacement method for signature 'filematrix'
dimnames(x) <- value</pre>
```

# **Arguments**

x A filematrix object (filematrix).

i, j Row/column indices specifying elements to extract or replace.

value A new value to replace the indexed element(s).

# Value

length function returns the number of elements in the filematrix.

Functions colnames, rownames, and dimnames return the same values as their counterparts for the regular R matrices.

# Methods

```
isOpen Returns TRUE is the filematrix is open.
readAll(): Return the whole matrix.
    Same as fm[] or as.matrix(fm)
writeAll(value): Fill in the whole matrix.
    Same as fm[] = value
readSubCol(i, j, num): Read num values in column j starting with row i.
    Same as fm[i:(i+num-1), j]
writeSubCol(i, j, value): Write values in the column j starting with row i.
    Same as fm[i:(i+length(value)-1), j] = value
readCols(start, num): Read num columns starting with column start.
    Same as fm[, start:(start+num-1)]
writeCols(start, value): Write columns starting with column start.
    Same as fm[, start:(start+ncol(value)-1)] = value
```

fm.create 5

```
readSeq(start, len): Read len values from the matrix starting with start-th value.
    Same as fm[start:(start+len-1)]
writeSeq(start, value): Write values in the matrix starting with start-th value.
    Same as fm[start:(start+length(value)-1)] = value
appendColumns(mat) Increases filematrix by adding columns to the right side of the matrix. Matrix mat must have the same number of rows.
    Same as fm = cbind(fm, mat) for ordinary matrices.
```

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#### See Also

For function creating and opening file matrices see fm. create.

Run browseVignettes("filematrix") for the list of vignettes.

fm.create

Functions to Create a New, or Open an Existing Filematrix

# Description

Create a new or open existing filematrix object.

fm.create creates a new filematrix. If a filematrix with this name exists, it is overwritten (destroyed).

fm.create.from.matrix creates a new filematrix copy of an existing R matrix.

fm. open opens an existing filematrix for read/write access.

fm. load loads entire existing filematrix into memory as an ordinary R matrix.

fm.create.from.text.file reads a matrix from a text file into a new filematrix. The rows in the text file become columns in the filematrix. The transposition happens because the text files stores data by rows and filematrices store data by columns.

# Usage

```
fm.create(
    filenamebase,
    nrow = 0,
    ncol = 1,
    type = "double",
    size = NULL,
    lockfile = NULL)

fm.create.from.matrix(
    filenamebase,
    mat,
```

6 fm.create

```
size = NULL,
    lockfile = NULL)
fm.open(
    filenamebase,
    readonly = FALSE,
    lockfile = NULL)
fm.load(filenamebase, lockfile = NULL)
fm.create.from.text.file(
    textfilename,
    filenamebase,
    skipRows = 1,
    skipColumns = 1,
    sliceSize = 1000,
    omitCharacters = "NA",
    delimiter = "\t",
    rowNamesColumn = 1,
    type = "double",
    size = NULL)
## S4 method for signature 'filematrix'
close(con)
closeAndDeleteFiles(con)
```

# **Arguments**

filenamebase	Name without	extension	for the	files	storing	the filematrix.

The file <filenamebase>.bmat keeps the matrix values and <filenamebase>.desc.txt

stores the matrix dimensions, data type, and data type size. Names of rows and

columns, if defined, are stored in <filenamebase>.nmsrow.txt and <filenamebase>.nmscol.txt.

nrow Number of rows in the matrix. Values over 2^32 are supported.

Number of columns in the matrix. Values over 2^32 are supported.

type The type of values stored in the matrix. Can be either "double", "integer",

"logical", or "raw".

size Size of each item of the matrix in bytes.

Default values are 8 for "double", 4 for "integer", and 1 for "logical" and

"raw".

Do not change if not sure.

mat Regular R matrix, to be copied into a new filematrix.

readonly If TRUE, the values in the opened filematrix cannot be changed.

textfilename Name of the text file with matrix data, to be copied into a new filematrix.

skipRows Number of rows with column names. The matrix values are expected after first

skipRows rows of the file. Can be zero.

fm.create 7

skipColumns Number of columns before matrix values begin. Can be zero.

sliceSize The text file with matrix is read in chuncks of sliceSize rows. This is a per-

formance tuning parameter, it does not affect the outcome.

omitCharacters The text string representing missing values. Default value is NA.

delimiter The delimiter separating values in the text matrix file.

rowNamesColumn The row names are taken from the rowNamesColumn-th column of the text file.

By default, row names are extracted from the first column.

con A filematrix object.

lockfile Optional. Name of a lock file (file is overwritten). Used to avoid simultaneous

operations by multiple R instances accessing the same filematrix or different

filematrices on the same hard drive. Do not use if not sure.

#### **Details**

Once created or opened, a filematrix object can be accessed as an ordinary matrix using both matrix fm[,] and vector fm[] indexing. The indices can be integer (no zeros) or logical vectors.

#### Value

Returns a filematrix object. The object can be closed with close command or closed and deleted from disk with closeAndDeleteFiles command.

# Author(s)

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# See Also

For more on the use of filematrices see filematrix.

Run browseVignettes("filematrix") for the list of vignettes.

# **Examples**

```
# Create a 10x10 matrix
fm = fm.create(filenamebase=tempfile(), nrow=10, ncol=10)
# Change values in the top 3x3 corner
fm[1:3,1:3] = 1:9
# View the values in the top 4x4 corner
fm[1:4,1:4]
# Close and delete the filematrix
closeAndDeleteFiles(fm)
```

# **Index**

```
* classes
    filematrix-class, 3
* filematrix
    filematrix-package, 2
[.filematrix(filematrix-class), 3
[<-.filematrix(filematrix-class), 3
as.matrix,filematrix-method
        (filematrix-class), 3
close, filematrix-method (fm. create), 5
closeAndDeleteFiles, 7
closeAndDeleteFiles(fm.create), 5
colnames, filematrix-method
        (filematrix-class), 3
colnames<-,filematrix-method
        (filematrix-class), 3
dim,filematrix-method
        (filematrix-class), 3
dim<-,filematrix-method</pre>
        (filematrix-class), 3
dimnames, filematrix-method
        (filematrix-class), 3
dimnames<-,filematrix-method
        (filematrix-class), 3
filematrix, 2, 5, 7
filematrix (filematrix-class), 3
filematrix-class, 3
filematrix-package, 2
fm.create, 2, 3, 5, 5
fm.create.from.matrix, 2, 3
fm.create.from.text.file, 3
fm.load, 3
fm.load(fm.create), 5
fm. open, 2, 3
fm.open (fm.create), 5
length, filematrix-method
        (filematrix-class), 3
```

```
matrix, 2

rownames, filematrix-method
      (filematrix-class), 3

rownames<-, filematrix-method
      (filematrix-class), 3</pre>
```