# Package 'filematrix' 

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Type Package
Title File-Backed Matrix Class with Convenient Read and Write Access
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Description 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,
exactly as with usual R matrices.
Supports very large matrices.
Tested on multi-terabyte matrices.
Allows for more than $2^{\wedge} 32$ rows or columns.
Allows for quick addition of extra columns to a filematrix.
Cross-platform as the package has R code only.
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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 $2^{\wedge} 32$ 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 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: $\mathrm{fm}[1: 3,2: 4]$ and $\mathrm{fm}[, 2: 4]$.
The values can also be accessed as a vector with single indexing.
For example: $\mathrm{fm}[3: 7]$ and $\mathrm{fm}[4: 7]=1: 4$.
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: $\operatorname{dim}(f m)$ and $\operatorname{dim}(f m)=c(10,100)$.
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
```

```
## 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
```


## Arguments

x
A filematrix object (filematrix).
$i, j \quad$ Row/column indices specifying elements to extract or replace.
value $\quad$ 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 $\mathrm{fm}[\mathrm{i}:(\mathrm{i}+\mathrm{num}-1)$, 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
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 $f m=\operatorname{cbind}(f m$, 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,
```

```
    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. <br> 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 $2^{\wedge} 32$ are supported. |
| ncol | Number of columns in the matrix. Values over $2^{\wedge} 32$ 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. <br> Default values are 8 for "double", 4 for "integer", and 1 for "logical" and "raw". <br> 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. |


| skipColumns | Number of columns before matrix values begin. Can be zero. <br> sliceSize |
| :--- | :--- |
| The text file with matrix is read in chuncks of sliceSize rows. This is a per- <br> 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. <br> 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 <br> operations by multiple R instances accessing the same filematrix or different <br> 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 $f m[$,$] and vector f m[]$ 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.

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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)
```


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