# Package 'mmapcharr'

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Title Memory-Map Character Files
Version 0.3.1
<b>Date</b> 2025-11-27
<b>Description</b> Uses memory-mapping to enable the random access of elements of a text file of characters separated by characters as if it were a simple R(cpp) matrix.
Encoding UTF-8
License GPL-3
ByteCompile TRUE
<b>Depends</b> R (>= 3.3.0)
Imports methods, Rcpp
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Suggests testthat
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<pre>URL https://github.com/privefl/mmapcharr</pre>
<pre>BugReports https://github.com/privefl/mmapcharr/issues</pre>
Collate 'RcppExports.R' 'extract.R' 'file-dim.R' 'mmapchar.R' 'mmapcharr-package.r' 'utils.R'
NeedsCompilation yes
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dim\_file

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File dimensions

## **Description**

Number of lines and columns of file (and extra 'return' characters).

#### Usage

```
dim_file(file)
```

#### **Arguments**

file

Path to file.

#### Value

The number of lines and columns of file (and extra 'return' characters).

## **Examples**

```
tmpfile <- tempfile()
write(0:9, tmpfile, ncolumns = 2)
dim_file(tmpfile)</pre>
```

Extract

Create an Implementation of [ For Custom Matrix-Like Types

## **Description**

extract is a function that converts different index types such as negative integer vectors or logical vectors passed to the [function as i (e.g. X[i]) or i and j (e.g. X[i, j]) into positive integer vectors. The converted indices are provided as the i parameter of extract\_vector or i and j parameters of extract\_matrix to facilitate implementing the extraction mechanism for custom matrix-like types.

## Usage

```
Extract(extract_vector, extract_matrix)
```

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## Arguments

extract\_vector A function in the form of function(x, i) that takes a subset of x based on a single vector of indices i and returns a vector.extract\_matrix A function in the form of function(x, i, j) that takes a subset of x based on two vectors of indices i and j and returns a matrix.

#### **Details**

The custom type must implement methods for dim for this function to work. Implementing methods for nrow and ncol is not necessary as the default method of those generics calls dim internally.

This idea initially comes from package crochet.

## Value

A function in the form of function(x, i, j, ..., drop = TRUE) that is meant to be used as a method for [ for a custom type.

mmapchar-class Class mmapchar
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## **Description**

A reference class for storing and accessing matrix-like data stored on disk in files containing only characters (digits) separated by a character.

#### Usage

```
mmapchar(file, code)
```

#### **Arguments**

```
file Path of the file.

code Integer vector of size 256 to access integers instead of rawToChar(as.raw(0:255),
multiple = TRUE). See mmapcharr:::CODE_012 and mmapcharr:::CODE_DIGITS.
```

#### **Examples**

```
test_file <- system.file("testdata/test-windows.txt", package = "mmapcharr")
test <- mmapchar(test_file, code = mmapcharr:::CODE_012)
test[, 1:3]
test[]
readLines(test_file)</pre>
```

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Methods for the mmapchar class

## Description

Methods for the mmapchar class

Accessor methods for class mmapchar. You can use positive and negative indices, logical indices (that are recycled) and also a matrix of indices (but only positive ones).

Dimension and type methods for class mmapchar.

## Usage

```
## S4 method for signature 'mmapchar'
x[i, j, ..., drop = TRUE]
## S4 method for signature 'mmapchar'
dim(x)
## S4 method for signature 'mmapchar'
length(x)
```

## Arguments

X	A mmapchar object.
i	A vector of indices (or nothing). You can use positive and negative indices, logical indices (that are recycled) and also a matrix of indices (but only positive ones).
j	A vector of indices (or nothing). You can use positive and negative indices, logical indices (that are recycled).
	Not used. Just to make nargs works.
drop	Whether to delete the dimensions of a matrix which have one dimension equals to 1.

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Size of line

## **Description**

Number of elements of each line of a file.

## Usage

```
nelem(file)
```

nline 5

## **Arguments**

file

Path to file.

## Value

The number of elements of each line of a file.

## **Examples**

```
tmpfile <- tempfile()
write(1:10, tmpfile, ncolumns = 2)
nline(tmpfile)</pre>
```

nline

Number of lines

## Description

Number of lines of a file.

## Usage

```
nline(file)
```

## **Arguments**

file

Path to file.

#### Value

The number of lines of the file.

## Examples

```
tmpfile <- tempfile()
write(1:5, tmpfile, ncolumns = 1)
nline(tmpfile)</pre>
```

## **Index**