

Package ‘proxyC’

December 31, 2018

Type Package

Title Computes Proximity in Large Sparse Matrices

Version 0.1.0

Description Computes proximity between rows or columns of large matrices efficiently in C++.

Functions are optimized for large sparse matrices using the Armadillo and Intel TBB libraries.

Among several built-in similarity/distance measures, computation of correlation, cosine similarity and Euclidean distance is particularly fast.

Encoding UTF-8

LazyData true

LinkingTo Rcpp, RcppParallel, RcppArmadillo (>= 0.7.600.1.0)

BugReports <https://github.com/koheiw/proxyC/issues>

System Requirements C++11

License GPL-3

Depends R (>= 3.1.0), methods

Imports Matrix (>= 1.2), Rcpp (>= 0.12.12), RcppParallel

Suggests testthat, proxy

Collate 'RcppExports.R' 'proxy.R'

RoxxygenNote 6.1.1

NeedsCompilation yes

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Repository CRAN

Date/Publication 2018-12-31 22:10:02 UTC

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simil	<i>Compute similiarty/distance between raws or columns of large matrices</i>
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Description

Fast similarity/distance computation function for large sparse matrices. You can floor small similiarty value to save computation time and storage space by an arbitrary threshold (`min_simil`) or rank (`rank`). Please increase the number of threads for better performance using [setThreadOptions](#).

Usage

```
simil(x, y = NULL, margin = 1, method = c("cosine", "correlation",
  "jaccard", "ejaccard", "dice", "edice", "hamman", "simple matching",
  "faith"), min_simil = NULL, rank = NULL)

dist(x, y = NULL, margin = 1, method = c("euclidean", "chisquared",
  "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski"),
  p = 2)
```

Arguments

<code>x</code>	a matrix or Matrix object
<code>y</code>	if a matrix or Matrix object is provided, proximity between documents or features in <code>x</code> and <code>y</code> is computed.
<code>margin</code>	integer indicating margin of similiarty/distance computation. 1 indicates rows or 2 indicates columns.
<code>method</code>	method to compute similarity or distance
<code>min_simil</code>	the minimum similiarty value to be recorded.
<code>rank</code>	an integer value specifying top-n most similiarty values to be recorded.
<code>p</code>	weight for minkowski distance

Examples

```
mt <- Matrix::rsparsematrix(100, 100, 0.01)
simil(mt, method = "cosine")[1:5, 1:5]
mt <- Matrix::rsparsematrix(100, 100, 0.01)
dist(mt, method = "euclidean")[1:5, 1:5]
```

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