Package 'NPE'

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Title Non-Parametric Estimators

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Author Sumit Sethi	
Maintainer Sumit Sethi <sumit.sethi@nyu.edu></sumit.sethi@nyu.edu>	
Description Functions for calculating a variety of nonparametric estimators, including estimators of location (median, Hodges-Lehmann), of dispersion (Median Absolute Deviation), and ofdependency-covariance (Theil-Sen). It also implements the nonparametric statistics of the Wilcoxon Signed Rank test, the Mann-Whitney-Wilcoxon Rank Sum test, and the Kruskal-Wallis test. It also implements PCA.	n
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calc_pca

Performs a principle component analysis on given matrix *or* time series *using* RcppArmadillo.

Description

Performs a principle component analysis on given matrix or time series using RcppArmadillo.

Usage

```
calc_pca(mat_rix)
```

Arguments

mat_rix

A matrix or a time series.

Details

The function calc_pca() performs a principle component analysis on a matrix using RcppArmadillo.

Value

A matrix of variable loadings (i.e. a matrix whose columns contain the eigenvectors).

Examples

```
## Not run:
# Create a matrix of random returns
re_turns <- matrix(rnorm(5e6), nc=5)
# Compare calc_pca() with standard prcomp()
all.equal(drop(NPE::calc_pca(re_turns)),
    prcomp(re_turns))
# Compare the speed of RcppArmadillo with R code
library(microbenchmark)
summary(microbenchmark(
    rcpp=NPE::calc_pca(re_turns),
    rcode=prcomp(re_turns),
    times=10))[, c(1, 4, 5)] # end microbenchmark summary
## End(Not run)</pre>
```

calc_ranksWithTies

Calculate the ranks of the elements of a vector or a single-column time series using RcppArmadillo and boost.

Description

Calculate the ranks of the elements of a *vector* or a single-column *time series* using RcppArmadillo and boost.

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Usage

```
calc_ranksWithTies(vec_tor)
```

Arguments

vec_tor

A vector or a single-column time series.

Details

The function calc_ranks() calculates the ranks of the elements of a *vector* or a single-column *time series*. It *averages* the ranks in case fo ties. It uses the boost function boost::sort::parallel_stable_sort for sorting array in parallel fashion.

Value

A double vector with the ranks of the elements of the vector.

Examples

```
## Not run:
# Create a vector of random data
da_ta <- round(runif(7), 2)</pre>
# Calculate the ranks of the elements in two ways
all.equal(rank(da_ta), drop(NPE::calc_ranksWithTies(da_ta)))
# Create a time series of random data
da_ta <- xts::xts(runif(7), seq.Date(Sys.Date(), by=1, length.out=7))</pre>
# Calculate the ranks of the elements in two ways
all.equal(rank(coredata(da\_ta)), \ drop(NPE::calc\_ranksWithTies(da\_ta)))
\# Compare the speed of this function with RcppArmadillo and R code
da_ta <- runif(7)</pre>
library(microbenchmark)
summary(microbenchmark(
  rcpp=calc_ranks(da_ta),
  rcode=rank(da_ta),
  boost=calc_ranksWithTies(da_ta)
  times=10))[, c(1, 4, 5)] # end microbenchmark summary
## End(Not run)
```

hle

Calculate the non parametric Hodges-Lehmann estimator of location for a vector or a single-column time series using RcppArmadillo and RcppParallel.

Description

Calculate the non parametric Hodges-Lehmann estimator of location for a *vector* or a single-column *time series* using RcppArmadillo and RcppParallel.

Usage

```
hle(vec_tor)
```

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Arguments

vec_tor

A vector or a single-column time series.

Details

The function hle() calculates the Hodges-Lehmann estimator of the *vector*, using RcppArmadillo and RcppParallel. The function hle() is very much faster than function wilcox.test() in R.

Value

A single double value representing Hodges-Lehmann estimator of the vector.

Examples

```
## Not run:
# Create a vector of random returns
re_turns <- rnorm(1e6)
# Compare hle() with wilcox.test()
all.equal(drop(NPE::hle(re_turns)),
    wilcox.test(re_turns, conf.int = TRUE))
# Compare the speed of RcppParallel with R code
library(microbenchmark)
summary(microbenchmark(
    rcpp=NPE::hle(re_turns),
    rcode=wilcox.test(re_turns, conf.int = TRUE),
    times=10))[, c(1, 4, 5)] # end microbenchmark summary
## End(Not run)</pre>
```

kruskalWalliceTest

Performs a Kruskal-Wallis rank sum test. using Rcpp and boost.

Description

Performs a Kruskal-Wallis rank sum test. using Rcpp and boost.

Usage

```
kruskalWalliceTest(x)
```

Arguments

Х

A List of numeric data vectors

Details

The function kruskalWalliceTest() performs a Kruskal-Wallis rank sum test of the null hypothesis that the location parameters of the distribution of x are the same in each group. The alternative is that they differ in at least in one.

Value

A double indicating p-value of the test.

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Examples

```
## Not run:
x <- c(2.9, 3.0, 2.5, 2.6, 3.2) # normal subjects
y <- c(3.8, 2.7, 4.0, 2.4)  # with obstructive airway disease
z <- c(2.8, 3.4, 3.7, 2.2, 2.0) # with asbestosis

# Carry out Kruskal wallice rank sum test on the elements in two ways
all.equal(kruskal.test(list(x, y, z))$p.value, drop(NPE::kruskalWalliceTest(list(x, y, z))))
# Compare the speed of Rcpp and R code
library(microbenchmark)
summary(microbenchmark)
summary(microbenchmark(
   rcpp=kruskalWalliceTest(list(x, y, z)),
   rcode=kruskal.test(list(x, y, z))$p.value,
   times=10))[, c(1, 4, 5)] # end microbenchmark summary

## End(Not run)</pre>
```

medianAbsoluteDeviation

Calculate the Median absolute deviation of a vector or a single-column time series using RcppArmadillo.

Description

Calculate the Median absolute deviation of a vector or a single-column time series using RcppArmadillo.

Usage

```
medianAbsoluteDeviation(vec_tor)
```

Arguments

vec_tor

A vector or a single-column time series.

Details

The function medianAbsoluteDeviation() calculates the median of the *vector*, using RcppArmadillo. The function medianAbsoluteDeviation() is several times faster than mad() in R.

Value

A single double value representing median absolue deviation of the vector.

Examples

```
## Not run:
# Create a vector of random returns
re_turns <- rnorm(1e6)
# Compare medianAbsoluteDeviation() with mad()
all.equal(drop(NPE::medianAbsoluteDeviation(re_turns)),
   mad(re_turns))
# Compare the speed of RcppArmadillo with R code</pre>
```

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```
library(microbenchmark)
summary(microbenchmark(
  rcpp=NPE::medianAbsoluteDeviation(re_turns),
  rcode=mad(re_turns),
  times=10))[, c(1, 4, 5)] # end microbenchmark summary
## End(Not run)
```

med_ian

Calculate the median of a vector or a single-column time series using RcppArmadillo.

Description

Calculate the median of a *vector* or a single-column *time series* using RcppArmadillo.

Usage

```
med_ian(vec_tor)
```

Arguments

vec_tor

A vector or a single-column time series.

Details

The function med_ian() calculates the median of the *vector*, using RcppArmadillo. The function med_ian() is several times faster than median() in R.

Value

A single double value representing median of the vector.

Examples

```
## Not run:
# Create a vector of random returns
re_turns <- rnorm(1e6)
# Compare med_ian() with median()
all.equal(drop(NPE::med_ian(re_turns)),
    median(re_turns))
# Compare the speed of RcppArmadillo with R code
library(microbenchmark)
summary(microbenchmark(
    rcpp=NPE::med_ian(re_turns),
    rcode=median(re_turns),
    times=10))[, c(1, 4, 5)] # end microbenchmark summary
## End(Not run)</pre>
```

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rolling_mad	Calculate the rolling median absolute deviation over a vector or a
	single-column time series using RcppArmadillo and RcppParallel.

Description

Calculate the rolling median absolute deviation over a *vector* or a single-column *time series* using RcppArmadillo and RcppParallel.

Usage

```
rolling_mad(vec_tor, look_back)
```

Arguments

vec_tor A *vector* or a single-column *time series*.

look_back The length of look back interval, equal to the number of elements of data used

for calculating the median.

Details

The function rolling_mad() calculates a vector of rolling medians, over a *vector* of data, using *RcppArmadillo* and *RcppParallel*.

Value

A column *vector* of the same length as the argument vect_tor.

Examples

```
## Not run:
# Create a vector of random returns
re_turns <- rnorm(1e6)
rolling_mad(re_turns)
## End(Not run)</pre>
```

 $rolling_median$

Calculate the rolling median over a vector or a single-column time series using RcppArmadillo and RcppParallel.

Description

Calculate the rolling median over a *vector* or a single-column *time series* using RcppArmadillo and RcppParallel.

Usage

```
rolling_median(vec_tor, look_back)
```

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Arguments

vec_tor A vector or a single-column time series.

look_back The length of look back interval, equal to the number of elements of data used

for calculating the median.

Details

The function rolling_median() calculates a vector of rolling medians, over a *vector* of data, using *RcppArmadillo* and *RcppParallel*. The function rolling_median() is faster than roll::roll_median() which uses Rcpp.

Value

A column *vector* of the same length as the argument vect_tor.

Examples

```
## Not run:
# Create a vector of random returns
re_turns <- rnorm(1e6)
# Compare rolling_median() with roll::roll_median()
all.equal(drop(NPE::rolling_median(re_turns, look_back=11)),
    roll::roll_median(re_turns, width=11))
# Compare the speed of RcppArmadillo with R code
library(microbenchmark)
summary(microbenchmark(
    parallel_rcpp=NPE::rolling_median(re_turns),
    rcpp=roll::roll_median(re_turns),
    times=10))[, c(1, 4, 5)] # end microbenchmark summary
## End(Not run)</pre>
```

theilSenEstimator

Calculate the non parametric Theil-Sen estimator of dependencycovariance for two vectors using RcppArmadillo

Description

Calculate the non parametric Theil-Sen estimator of dependency-covariance for two *vectors* using RcppArmadillo

Usage

```
theilSenEstimator(x, y)
```

Arguments

vector_x A *vector* independent (explanatory) data.

vector_y A vector dependent data.

Details

The function theilSenEstimator() calculates the Theil-Sen estimator of the *vector*, using RcppArmadillo . The function theilSenEstimator() is significantly faster than function WRS::tsreg() in R.

Value

A column *vector* containing two values i.e intercept and slope

Examples

```
## Not run:
# Create a vector of random returns
vector_x <- rnorm(10)
vactor_y <- rnorm(10)
# Compare theilSenEstimator() with tsreg()
# Compare the speed of RcppParallel with R code
library(microbenchmark)
summary(microbenchmark(
   rcpp=NPE::theilSenEstimator(vector_x, vector_y),
   rcode=WRS(vector_x, vector_y),
   times=10))[, c(1, 4, 5)] # end microbenchmark summary
## End(Not run)</pre>
```

wilcoxanMannWhitneyTest

Performs two sample Wilcoxan-Mann-Whitney rank sum test also known as Mann-Whitney U Test on vector or a single-column time series using RcppArmadillo and boost.

Description

Performs two sample Wilcoxan-Mann-Whitney rank sum test also known as Mann-Whitney U Test on *vector* or a single-column *time series* using RcppArmadillo and boost.

Usage

```
wilcoxanMannWhitneyTest(x, y, mu = 0, alternative = "two.sided",
  exact = FALSE, correct = TRUE)
```

Arguments

mu

x A *vector* or a single-column *time series*.

y A *vector* or a single-column *time series*.

A *double* specifing an optional parameter used to form null hypothesis. Default value is *zero*.

alternative a *character* string specifying the alternative hypothesis. It must be one of :

- "two.sided" two tailed test.
- "greater" greater(right) tailed test.
- "less" smaller(left) tailed test.

(The default is two.sided test.)

exact A boolean indicating whether an exact p-value should be computed.

correct A boolean indicating whether to apply continuity correction in normal approxi-

mation for the p-value.

Details

The function wilcoxanMannWhitneyTest() carries out the wilcoxan-Mann-Whitney signed rank test on x & y and returns the p-value of the test. By default (if exact is not specified), an exact p-value is computed if sample contains less than 50 finite values and there are no ties. Otherwise, a normal approximation is used.

Value

A double indicating p-value of the test.

Examples

```
## Not run:
# Create a vector of random data
x <- round(runif(10), 2)
y <- round(runif(10), 2)
# Carry out WMW signed rank test on the elements in two ways
all.equal(wilcox.test(x, y)$p.value, drop(NPE::wilcoxanMannWhitneyTest(x, y)))
# Compare the speed of Rcpp and R code
library(microbenchmark)
summary(microbenchmark(
  rcpp=wilcoxanMannWhitneyTest(x, y),
  rcode=wilcox.test(x, y),
  times=10))[, c(1, 4, 5)] # end microbenchmark summary</pre>
## End(Not run)
```

wilcoxanSignedRankTest

Performs one sample Wilcoxan ranked sum test on vector or a single-column time series using RcppArmadillo and boost.

Description

Performs one sample Wilcoxan ranked sum test on *vector* or a single-column *time series* using RcppArmadillo and boost.

Usage

```
wilcoxanSignedRankTest(x, mu = 0, alternative = "two.sided",
  exact = FALSE, correct = TRUE)
```

Arguments

x A *vector* or a single-column *time series*.

mu A double specifing an optional parameter used to form null hypothesis. Default

value is zero.

alternative a *character* string specifying the alternative hypothesis. It must be one of :

• "two.sided" two tailed test.

• "greater" greater(right) tailed test.

• "less" smaller(left) tailed test.

(The default is *two.sided* test.)

exact A boolean indicating whether an exact p-value should be computed.

correct A boolean indicating whether to apply continuity correction in normal approxi-

mation for the p-value.

Details

The function wilcoxanSignedRankTest() carries out the wilcoxan signed rank test on vec_tor and returns the p-value of the test. By default (if exact is not specified), an exact p-value is computed if sample contains less than 50 finite values and there are no ties. Otherwise, a normal approximation is used.

Value

A double indicating p-value of the test.

Examples

```
## Not run:
# Create a vector of random data
da_ta <- round(runif(7), 2)</pre>
# Carry out wilcoxan signed rank test on the elements in two ways
all.equal(wilcox.test(da_ta)$p.value, drop(HighFreq::WilcoxanSignedRankTest(da_ta)))
# Create a time series of random data
da_ta <- xts::xts(runif(7), seq.Date(Sys.Date(), by=1, length.out=7))</pre>
# Calculate the ranks of the elements in two ways
all.equal(wilcox.test(coredata(da_ta))$p.value, drop(NPE::wilcoxanSignedRankTest(da_ta)))
# Compare the speed of Rcpp and R code
da_ta <- runif(10)
library(microbenchmark)
summary(microbenchmark(
  rcpp=wilcoxanSignedRankTest(da_ta),
  rcode=wilcox.test(da_ta),
  times=10))[, c(1, 4, 5)] # end microbenchmark summary
## End(Not run)
```

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