Package 'toscca'

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Title What the Package Does (One Line, Title Case)

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Description What the package does (one paragraph).

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boostrapCCA *Bootstrap to get empirical intervals of canonical correlation.*

Description

Bootstrap to get empirical intervals of canonical correlation.

Usage

```
boostrapCCA(
  A,
  B,
  nonzero_a,
  nonzero_b,
  cancor,
  folds = 10,
  n = 100,
  ci_quant = 0.01,
  silent = TRUE,
  toPlot = FALSE,
  parallel_logic = TRUE,
  nuisanceVar = 0,
  testStatType = "CC"
)
```

Arguments

A, B	Data matrices.
nonzero_a, nonzero_b	Numeric. Scalar or vector over the number of nonzeroes allowed for a correlation estimate.
cancor	Numeric. Scalar or vector: anonical correlation estimate(s).
folds	Numeric. Number of folds for the cross-validation process.
n	Numeric. Times of bootstrapping.
ci_quant	Numeric. Between 0 and 1. quantile of intervale.
silent	Logical. If FALSE, a progress bar will appear on the console. Default is FALSE.
toPlot	Logical. If TRUE, plot will be generated automatically showing the estimated canonical weights. Default is TRUE.
parallel_logic	Logical. If TRUE, cross-validation is done in parallel.Default is FALSE.
nuisanceVar	Data with nuisance variables. For statistic type.
testStatType	Character. Choice of statistic. Options are CC (default), Wilks and Roy.
K	Numeric. Number of components to be computed.
draws	Numeric. Number of permutations for each component.

Details

For a exploratory analysis nonzero_a and nonzero_b can be vectors. The algorithm will then search for the best combination of sparsity choice nonzero_a and nonzero_b for each component.

Value

Matrix with permutation estimates.

`CCAtStat` *Get the estatistic for the permutations.*

Description

Get the estatistic for the permutations.

Usage

```
CCAtStat(cancor, A, B, C = 0, type = c("CC", "Wilks", "Roy"))
```

Arguments

`cancor`	Numeric. Canonical Correlation estimate.
`A`	An nxp matrix.
`B`	An nxq matrix.
`C`	An nxs matrix. Confounding variables.
`type`	Character. Choice of statistic: Canonical correlation, Wilks'statistic or Roy's statistic.

Value

Statistic

`cpev.fun` *Calculated cummulative percentage of explained variance.*

Description

Calculated cummulative percentage of explained variance.

Usage

```
cpev.fun(mat, matK)
```

Arguments

`mat`	An nxp matrix.
`maxK`	An nxk matrix. Each column corresponds to a latent variable.

Value

Scalar.

`eigenDecompostion` *Performs eigen decomposition of a matrix in PS space.*

Description

Performs eigen decomposition of a matrix in PS space.

Usage

```
eigenDecompostion(A)
```

Arguments

`A`	A square matrix nxn.

Value

Matrix. Positive definite matrix.

`getCanSubspace` *Performs matrix residualisation over estimated canonical vectors. There are three types: basic (subtracts scaled estimated latent variable from data), null (uses the null space of the estimated canonical vector to construct a new matrix) and LV (uses SVD to residualise).*

Description

Performs matrix residualisation over estimated canonical vectors. There are three types: basic (subtracts scaled estimated latent variable from data), null (uses the null space of the estimated canonical vector to construct a new matrix) and LV (uses SVD to residualise).

Usage

```
getCanSubspace(mat, vec)
```

Arguments

`mat`	An nxp matrix.
`vec`	A vector of dimensions nxk.

Details

For nxp matrix

$$\mathbf{A}$$

and pxk vector

$$\alpha$$

, the canonical is compute as $\mathbf{A}_{sub} = \mathbf{A}\alpha(\alpha^T\alpha)\alpha^T$.

Value

An nxk matrix.

`initialiseCanVar` *Initialised the canonical vector for the iterative process based on positive eigen values. Then, SVD is performed on that PS matrix.*

Description

Initialised the canonical vector for the iterative process based on positive eigen values. Then, SVD is performed on that PS matrix.

Usage

```
initialiseCanVar(A, B)
```

Arguments

`A`	An nxp matrix.
`B`	An nxq matrix.

Value

An pzp vector.

`KFoldSCCA` *Sparse Canonical Correlation Analysis. Computation of CC via NIPALS with soft thresholding.*

Usage

```
KFoldSCCA(
  A,
  B,
  nonzero_a,
  nonzero_b,
  alpha_init = c("eigen", "random", "uniform"),
  folds = 1,
  parallel_logic = FALSE,
  silent = FALSE,
  toPlot = TRUE,
  ATest_res = NULL,
  BTest_res = NULL
)
```

Arguments

`A, B`	Data matrices.
`nonzero_a, nonzero_b`	Numeric. Scalar or vector over the number of nonzeroes allowed for a correlation estimate.
`silent`	Logical. If FALSE, a progress bar will appear on the console. Default is FALSE.

alphaInit	Character. Type initialisation for α. Default is "eigen".
iter	Numeric. Maximum number of iterations. Default is 20.
tol	Numeric. Tolerance threshold. Default is 10^6.

Value

a list with the following elements:

- alphaCanonical vector for matrix $\mathbf{A}$, for each combination of sparsity value specified.
- betaCanonical vector for matrix $\mathbf{B}$, for each combination of sparsity value specified.
- cancorMax. canonical correlation estimate.
- nonzero_a,nonzero_bOptimal nonzero values for each canonical vector.

Sparse Canonical Correlation Analysis. Computation of CC via NIPALS with soft thresholding.

MSCCA *Sparse Canonical Correlation Analysis. Computation of CC via NIPALS with soft thresholding.*

Usage

```
MSCCA(
  A,
  B,
  nonzero_a,
  nonzero_b,
  K = 1,
  alpha_init = c("eigen", "random", "uniform"),
  folds = 1,
  silent = FALSE,
  toPlot = TRUE,
  typeResid = "basic",
  combination = TRUE,
  parallel_logic = FALSE
)
```

Arguments

A, B	Data matrices.
nonzero_a, nonzero_b	Numeric. Scalar or vector over the number of nonzeroes allowed for a correlation estimate.
K	Numeric. Number of components to be computed.
alpha_init	Character. Type initialisation for α . Default is "eigen".
folds	Numeric. Number of folds for the cross-validation process.
silent	Logical. If FALSE, a progress bar will appear on the console. Default is FALSE.
toPlot	Logical. If TRUE, plot will be generated automatically showing the estimated canonical weights. Default is TRUE.
typeResid	Character. Choice of residualisation technique. Options are basic (default), null and LV.
combination	Logical. If TRUE, the algorithm will search for the best combination of sparsity choice nonzero_a and nonzero_b for each component. This should be used for exploratory analysis. Default is FALSE.
parallel_logic	Logical. If TRUE, cross-validation is done in parallel.Default is FALSE.

Value

a list with the following elements:

- alphaCanonical vector for matrix $\mathbf{A}$, for each combination of sparsity value specified.
- betaCanonical vector for matrix $\mathbf{B}$, for each combination of sparsity value specified.
- cancorMax. canonical correlation estimate.

This function performs CCA on matrices $\mathbf{A}$ and $\mathbf{B}$ via Non-Iterative PArtial Least Squares (NIPALS) algorithm imposing sparsity over a fixed number of variables especified.

For a exploratory analysis nonzero_a and nonzero_b can be vectors. The algorithm will then search for the best combination of sparsity choice nonzero_a and nonzero_b for each component.

myHeatmap *Plot heatmap*

Description

This function generated a heatmap. Withing the package it is used to provide with a visualisation of the exploration of optimal sparsity levels.

Usage

```
myHeatmap(mat, palette = "Teal", coln = 12, xlab = "", ylab = "", axes = FALSE)
```

Arguments

mat	Matrix containing values along a grid.
palette	Choice of palette. Default is Teal.
coln	Number of columns for the grid distribution. Default is 12.
xlab	Lable for X axis.
ylab	Lable for Y axis.
axes	Logical. Have axes between 0 and 1. Default is FALSE.

Value

Grid plot.

permcvscca *Permutation testing for MSCCA*

Description

This function performs permutation testing on CC estimates.

Usage

```
permcvscca(
  A,
  B,
  nonzero_a,
  nonzero_b,
  K,
  alpha_init = c("eigen", "random", "uniform"),
  folds = 1,
  toPlot = FALSE,
  draws = 20,
  cancor,
  bootCCA = NULL,
  silent = TRUE,
  parallel_logic = TRUE,
  nuisanceVar = 0,
  testStatType = "CC"
)
```

Arguments

A, B	Data matrices.
nonzero_a, nonzero_b	Numeric. Scalar or vector over the number of nonzeroes allowed for a correlation estimate.
K	Numeric. Number of components to be computed.
folds	Numeric. Number of folds for the cross-validation process.
toPlot	Logical. If TRUE, plot will be generated automatically showing the estimated canonical weights. Default is TRUE.
draws	Numeric. Number of permutations for each component.
cancor	Numeric. Scalar or vector: anonical correlation estimate(s).
silent	Logical. If FALSE, a progress bar will appear on the console. Default is FALSE.
parallel_logic	Logical. If TRUE, cross-validation is done in parallel.Default is FALSE.
nuisanceVar	Data with nuisance variables. For statistic type.
testStatType	Character. Choice of statistic. Options are CC (default), Wilks and Roy.
combination	Logical. If TRUE, the algorithm will search for the best combination of sparsity choice nonzero_a and nonzero_b for each component. This should be used for exploratory analysis. Default is FALSE.

Details

For a exploratory analysis nonzero_a and nonzero_b can be vectors. The algorithm will then search for the best combination of sparsity choice nonzero_a and nonzero_b for each component.

Value

Matrix with permutation estimates.

---

powerMethod *Performs power method.*

Description

Performs power method.

Usage

```
powerMethod(mat, vec, tol = 10^(-6), maxIter = 500, silent = TRUE)
```

Arguments

mat	A square matrix nxn.
vec	A vector of dimensions nx1.
tol	Convergence criterion. Default is 10^(-6).
silent	Logical. If TRUE, convergence performance will be printed.
matIter	Maximum iterations. Default is 500.

Value

List: vec: eigen vector; lambda: eigen value; t: total iterations.

progressBar *Progress bar*

Description

Shows progress of a process.

Usage

```
progressBar(end, round)
```

Arguments

end	maximum number of times a process will run.
round	current round

Value

Display in consol of current status.

residualisation *Performs matrix residualisation over estimated canonical vectors. There are three types: basic (subtracts scaled estimated latent variable from data), null (uses the null space of the estimated canonical vector to construct a new matrix) and LV (uses SVD to residualise).*

Description

Performs matrix residualisation over estimated canonical vectors. There are three types: basic (subtracts scaled estimated latent variable from data), null (uses the null space of the estimated canonical vector to construct a new matrix) and LV (uses SVD to residualise).

Usage

```
residualisation(
  mat,
  vec,
  spaceMat = NULL,
  type = c("LV", "null", "basic"),
  na.allow = TRUE
)
```

Arguments

mat	An nxp matrix.
vec	A vector of dimensions nxk.
spaceMat	Only for "null" type residualisation. Default is NULL.
type	Character. It can be LV, null or basic depending on which type of residualisation will be performed.
na.allow	Logical. If TRUE, NAs will be allowed.

Value

Matrix.

`scaledResidualMat` *Performs scalling for matrix residualisation based on calculated coefficients.*

Description

Performs scalling for matrix residualisation based on calculated coefficients.

Usage

```
scaledResidualMat(A)
```

Arguments

`A`	An nxp matrix.

Value

scaled matrix.

`SCCA` *Sparse Canonical Correlation Analysis. Computation of CC via NIPALS with soft thresholding.*

Usage

```
SCCA(
  alphaInit,
  A,
  B,
  nonzero_a,
  nonzero_b,
  iter = 20,
  tol = 10^(-6),
  silent = FALSE
)
```

Arguments

`alphaInit`	Character. Type initialisation for α .
`A, B`	Data matrices.
`nonzero_a, nonzero_b`	Numeric. Scalar or vector over the number of nonzeroes allowed for a correlation estimate.

iter	Numeric. Maximum number of iterations. Default is 20.
tol	Numeric. Tolerance threshold. Default is 10^6.
silent	Logical. If FALSE, a progress bar will appear on the console. Default is FALSE.

Value

a list with the following elements:

- alphaCanonical vector for matrix $\mathbf{A}$, for each combination of sparsity value specified.
- betaCanonical vector for matrix $\mathbf{B}$, for each combination of sparsity value specified.
- cancorMax. canonical correlation estimate.
- cancor_allCall canonical correlations calculated for each sparsity levels.

Sparse Canonical Correlation Analysis. Computation of CC via NIPALS with soft thresholding.

---

standardVar *Stardardise a matrix*

Description

This function stardardises a matrix or a vector and gives the option to centre or normalise (only vectors).

Usage

```
standardVar(mat, centre = TRUE, normalise = FALSE)
```

Arguments

centre	Logical, if true, cetre to mean zero.
normalise	Logical, if true, performs vector normalisation.
X	Matrix or vector to be standardise.

Value

A matrix or vector with the preferred standardarisation

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