Package 'RaJIVE'

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Title Robust Angle Based Joint and Individual Variation Explained

Version 1.0

Description A robust alternative to the aJIVE (angle based Joint and Individual Variation Explained) method (Feng et al 2018: <doi:10.1016/j.jmva.2018.03.008>) for the estimation of joint and individual components in the presence of outliers in multi-source data. It decomposes the multi-source data into joint, individual and residual (noise) contributions. The decomposition is robust to outliers and noise in the data. The method is illustrated in Ponzi et al (2021) <doi:10.48550/arXiv.2101.09110>.

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Depends R (>= 3.1.0)

Suggests knitr, rmarkdown, testthat (>= 2.1.0), cowplot, reshape2, dplyr

Imports ggplot2, doParallel, foreach

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Repository CRAN

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Contents

`ajive.data.sim` *Simulation of data blocks*

Description

Simulates blocks of data with joint and individual structures

Usage

```
ajive.data.sim(
  K = 3,
  rankJ = 2,
  rankA = c(20, 15, 10),
  n = 100,
  pks,
  dist.type = 1,
  noise = 1
)
```

Arguments

`K`	Integer. Number of data blocks.
`rankJ`	Integer. Joint rank.
`rankA`	Vector of Integers. Individual Ranks.
`n`	Integer. Number of data points.
`pks`	Vector of Integers. Number of variables in each block.
`dist.type`	Integer. 1 for normal, 2 for uniform, 3 for exponential
`noise`	Integer. Standard deviation in dist

Value

Xsim a list of simulated data matrices and true rank values

Examples

```
n <- 20
p1 <- 10
p2 <- 8
p3 <- 5
JrankTrue <- 2
initial_signal_ranks <- c(5, 2, 2)
 Y <- ajive.data.sim(K =3, rankJ = JrankTrue,
 rankA = initial_signal_ranks,n = n,
 pks = c(p1, p2, p3), dist.type = 1)
```

---

data_heatmap *Decomposition Heatmaps*

Description

Visualization of the RaJIVE decomposition, it shows heatmaps of the decomposition obtained by RaJIVE

Usage

```
data_heatmap(data, show_color_bar = TRUE, title = "", xlab = "", ylab = "")
```

Arguments

data	List. The initial data blocks.
show_color_bar	Boolean.
title	Character.
xlab	Character.
ylab	Character

decomposition_heatmaps_robustH *Decomposition Heatmaps*

Description

Visualization of the RaJIVE decomposition, it shows heatmaps of the decomposition obtained by RaJIVE

Usage

```
decomposition_heatmaps_robustH(blocks, jive_results_robust)
```

Arguments

blocks	List. The initial data blocks.
jive_results_robust	List. The RaJIVE decomposition.

Value

The heatmap of the decomposition

Examples

```
n <- 10
pks <- c(20, 10)
Y <- ajive.data.sim(K =2, rankJ = 2, rankA = c(7, 4), n = n,
                pks = pks, dist.type = 1)
initial_signal_ranks <-  c(7, 4)
data.ajive <- list((Y$sim_data[[1]]), (Y$sim_data[[2]]))
ajive.results.robust <- Rajive(data.ajive, initial_signal_ranks)
decomposition_heatmaps_robustH(data.ajive, ajive.results.robust)
```

get_block_loadings *Block Loadings*

Description

Gets the block loadings from the Rajive decomposition

Usage

```
get_block_loadings(ajive_output, k, type)
```

Arguments

ajive_output	List. The decomposition from Rajive
k	Integer. The index of the data block
type	Character. Joint or individual

Value

The block loadings

Examples

```
n <- 10
pks <- c(20, 10)
Y <- ajive.data.sim(K =2, rankJ = 2, rankA = c(7, 4), n = n,
                    pks = pks, dist.type = 1)
initial_signal_ranks <-  c(7, 4)
data.ajive <- list((Y$sim_data[[1]]), (Y$sim_data[[2]]))
ajive.results.robust <- Rajive(data.ajive, initial_signal_ranks)
get_block_loadings(ajive.results.robust, 2, 'joint')
```

---

get_block_scores *Block Scores*

Description

Gets the block scores from the Rajive decomposition

Usage

```
get_block_scores(ajive_output, k, type)
```

Arguments

ajive_output	List. The decomposition from Rajive
k	Integer. The index of the data block
type	Character. Joint or individual

Value

The block scores

Examples

```
n <- 10
pks <- c(20, 10)
Y <- ajive.data.sim(K =2, rankJ = 2, rankA = c(7, 4), n = n,
                 pks = pks, dist.type = 1)
initial_signal_ranks <-  c(7, 4)
data.ajive <- list((Y$sim_data[[1]]), (Y$sim_data[[2]]))
ajive.results.robust <- Rajive(data.ajive, initial_signal_ranks)
get_block_scores(ajive.results.robust, 2, 'joint')
```

`get_final_decomposition_robustH` *Computes the final JIVE decomposition.*

Description

Computes X = J + I + E for a single data block and the respective SVDs.

Usage

```
get_final_decomposition_robustH(X, joint_scores, sv_threshold, full = TRUE)
```

Arguments

`X`	Matrix. The original data matrix.
`joint_scores`	Matrix. The basis of the joint space (dimension n x joint_rank).
`sv_threshold`	Numeric vector. The singular value thresholds from the initial signal rank estimates.
`full`	Boolean. Do we compute the full J, I matrices or just svd

---

`get_individual_decomposition_robustH` *Computes the individual matrix for a data block.*

Description

Computes the individual matrix for a data block.

Usage

```
get_individual_decomposition_robustH(
  X,
  joint_scores,
  sv_threshold,
  full = TRUE
)
```

Arguments

X	Matrix. The original data matrix.
joint_scores	Matrix. The basis of the joint space (dimension n x joint_rank).
sv_threshold	Numeric vector. The singular value thresholds from the initial signal rank estimates.
full	Boolean. Do we compute the full J, I matrices or just the SVD (set to FALSE to save memory).

---

get_individual_rank *Individual Rank*

Description

Gets the individual ranks from the Rajive decomposition

Usage

```
get_individual_rank(ajive_output, k)
```

Arguments

ajive_output	List. The decomposition from Rajive
k	Integer. The index of the data block.

Value

The individual ranks

Examples

```
n <- 10
pks <- c(20, 10)
Y <- ajive.data.sim(K =2, rankJ = 2, rankA = c(7, 4), n = n,
                 pks = pks, dist.type = 1)
initial_signal_ranks <-  c(7, 4)
data.ajive <- list((Y$sim_data[[1]]), (Y$sim_data[[2]]))
ajive.results.robust <- Rajive(data.ajive, initial_signal_ranks)
get_individual_rank(ajive.results.robust, 2)
```

get_joint_decomposition_robustH *Computes the individual matrix for a data block*

Description

Computes the individual matrix for a data block

Usage

```
get_joint_decomposition_robustH(X, joint_scores, full = TRUE)
```

Arguments

X	Matrix. The original data matrix.
joint_scores	Matrix. The basis of the joint space (dimension n x joint_rank).
full	Boolean. Do we compute the full J, I matrices or just the SVD (set to FALSE to save memory).

get_joint_rank *Joint Rank*

Description

Gets the joint rank from the Rajive decomposition

Usage

```
get_joint_rank(ajive_output)
```

Arguments

ajive_output	List. The decomposition from Rajive

Value

The joint rank

Examples

```
n <- 10
pks <- c(20, 10)
Y <- ajive.data.sim(K =2, rankJ = 2, rankA = c(7, 4), n = n,
                pks = pks, dist.type = 1)
initial_signal_ranks <-  c(7, 4)
data.ajive <- list((Y$sim_data[[1]]), (Y$sim_data[[2]]))
ajive.results.robust <- Rajive(data.ajive, initial_signal_ranks)
get_joint_rank(ajive.results.robust)
```

---

get_joint_scores_robustH

Computes the joint scores.

Description

Estimate the joint rank with the wedin bound, compute the signal scores SVD, double check each joint component.

Usage

```
get_joint_scores_robustH(
  blocks,
  block_svd,
  initial_signal_ranks,
  sv_thresholds,
  n_wedin_samples = 1000,
  n_rand_dir_samples = 1000,
  joint_rank = NA
)
```

Arguments

`blocks`	List. A list of the data matrices.
`block_svd`	List. The SVD of the data blocks.
`initial_signal_ranks`	Numeric vector. Initial signal ranks estimates.
`sv_thresholds`	Numeric vector. The singular value thresholds from the initial signal rank estimates.
`n_wedin_samples`	Integer. Number of wedin bound samples to draw for each data matrix.
`n_rand_dir_samples`	Integer. Number of random direction bound samples to draw.
`joint_rank`	Integer or NA. User specified joint_rank. If NA will be estimated from data.

get_random_direction_bound_robustH

Estimate the wedin bound for a data matrix.

Description

Samples from the random direction bound. Returns on the scale of squared singular value.

Usage

```
get_random_direction_bound_robustH(n_obs, dims, num_samples = 1000)
```

Arguments

n_obs	The number of observations.
dims	The number of features in each data matrix
num_samples	Integer. Number of vectors selected for resampling procedure.

Value

rand_dir_samples

get_svd_robustH

Computes the robust SVD of a matrix Using robRsvd

Description

Computes the robust SVD of a matrix Using robRsvd

Usage

```
get_svd_robustH(X, rank = NULL)
```

Arguments

X	Matrix. X matrix.
rank	Integer. Rank of SVD decomposition

Value

List. The SVD of X.

get_sv_threshold *The singular value threshold.*

Description

Computes the singular value threshold for the data matrix (half way between the rank and rank + 1 singluar value).

Usage

```
get_sv_threshold(singular_values, rank)
```

Arguments

singular_values	Numeric. The singular values.
rank	Integer. The rank of the approximation.

get_wedin_bound_samples *Gets the wedin bounds*

Description

Gets the wedin bounds

Usage

```
get_wedin_bound_samples(X, SVD, signal_rank, num_samples = 1000)
```

Arguments

X	Matrix. The data matrix.
SVD	List. The SVD decomposition of the matrix. List with entries 'u', 'd', and 'v'from the svd function.
signal_rank	Integer.
num_samples	Integer. Number of vectors selected for resampling procedure.

`Rajive` *Robust Angle based Joint and Individual Variation Explained*

Description

Computes the robust aJIVE decomposition with parallel computation.

Usage

```
Rajive(
  blocks,
  initial_signal_ranks,
  full = TRUE,
  n_wedin_samples = 1000,
  n_rand_dir_samples = 1000,
  joint_rank = NA
)
```

Arguments

`blocks`	List. A list of the data matrices.
`initial_signal_ranks`	Vector. The initial signal rank estimates.
`full`	Boolean. Whether or not to store the full J, I, E matrices or just their SVDs (set to FALSE to save memory).
`n_wedin_samples`	Integer. Number of wedin bound samples to draw for each data matrix.
`n_rand_dir_samples`	Integer. Number of random direction bound samples to draw.
`joint_rank`	Integer or NA. User specified joint_rank. If NA will be estimated from data.

Value

The aJIVE decomposition.

Examples

```
n <- 50
pks <- c(100, 80, 50)
Y <- ajive.data.sim(K =3, rankJ = 3, rankA = c(7, 6, 4), n = n,
                    pks = pks, dist.type = 1)
initial_signal_ranks <-  c(7, 6, 4)
data.ajive <- list((Y$sim_data[[1]]), (Y$sim_data[[2]]), (Y$sim_data[[3]]))
ajive.results.robust <- Rajive(data.ajive, initial_signal_ranks)
```

RobRSVD.all *Computes the robust SVD of a matrix*

Description

Computes the robust SVD of a matrix

Usage

```
RobRSVD.all(data, nrank = min(dim(data)), svdinit = svd(data))
```

Arguments

data	Matrix. X matrix.
nrank	Integer. Rank of SVD decomposition
svdinit	List. The standard SVD.

Value

List. The SVD of X.

showVarExplained_robust *Proportions of variance explained*

Description

Gets the variance explained by each component of the Rajive decomposition

Usage

```
showVarExplained_robust(ajiveResults, blocks)
```

Arguments

ajiveResults	List. The decomposition from Rajive
blocks	List. The initial data blocks

Value

The proportion of variance explained by each component

Examples

```
n <- 10
pks <- c(20, 10)
Y <- ajive.data.sim(K =2, rankJ = 2, rankA = c(7, 4), n = n,
                 pks = pks, dist.type = 1)
initial_signal_ranks <-  c(7, 4)
data.ajive <- list((Y$sim_data[[1]]), (Y$sim_data[[2]]))
ajive.results.robust <- Rajive(data.ajive, initial_signal_ranks)
showVarExplained_robust(ajive.results.robust, data.ajive)
```

---

sim_dist *Simulation of single data block from distribution*

Description

Simulation of single data block from distribution

Usage

```
sim_dist(num, n, p)
```

Arguments

num	Integer. Type of distribution. 1 for normal, 2 for uniform, 3 for exponential
n	Integer. Number of data points.
p	Integers. Number of variables in block.

---

svd_reconstruction *Reconstruces the original matrix from its robust SVD.*

Description

Computes UDV^T to get the approximate (or full) X matrix.

Usage

```
svd_reconstruction(decomposition)
```

Arguments

decomposition	List. List with entries 'u', 'd', and 'v'from the svd function.

Value

Matrix. The original matrix.

truncate_svd *Truncates a robust SVD.*

Description

Removes columns from the U, D, V matrix computed form an SVD.

Usage

```
truncate_svd(decomposition, rank)
```

Arguments

decomposition	List. List with entries 'u', 'd', and 'v'from the svd function.
rank	List. List with entries 'u', 'd', and 'v'from the svd function.

Value

The trucated robust SVD of X.

wedin_bound_resampling *Resampling procedure for the wedin bound*

Description

Resampling procedure for the wedin bound

Usage

```
wedin_bound_resampling(X, perp_basis, right_vectors, num_samples = 1000)
```

Arguments

X	Matrix. The data matrix.
perp_basis	Matrix. Either U_perp or V_perp: the remaining left/right singluar vectors of X after estimating the signal rank.
right_vectors	Boolean. Right multiplication or left multiplication.
num_samples	Integer. Number of vectors selected for resampling procedure.

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