

# Package ‘PSA’

October 6, 2024

**Title** Principal Nested Simplices Analysis

**Version** 0.0.0.9000

**Description** What the package does (one paragraph).

**License** GPL-3

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.2

**Imports** stats,  
utils,  
graphics,  
ggplot2 (>= 3.4.0),  
dendextend,  
compositions

**Depends** R (>= 4.0.0)

**Suggests** testthat (>= 3.0.0)

**Enhance** ggtern

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compare_analysis	<i>Comparison of PSA and Benchmark Methods</i>
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**Description**

A wrapper function of PSA and benchmark methods for convenience

**Usage**

```
compare_analysis(X)
```

**Arguments**

X                      a data matrix

**Value**

a list of data matrix X and outcomes of PSA-S, PSA-O, PCA, log-ratio PCA, and power transform PCA with power 1/2.

**See Also**

[psa\(\)](#), [comp\\_pca\(\)](#), [comp\\_apca\(\)](#), [comp\\_power\\_pca\(\)](#)

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comp_apca	<i>Log-Ratio Principal Component Analysis for PSA Comparison</i>
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**Description**

A wrapper function of princomp.acomp() for comparison to PSA. Zeros are substituted by half of the overall nonzero minimum. Manipulates result of princomp.acomp() in a similar format to psa().

**Usage**

```
comp_apca(X)
```

**Arguments**

X                      a data matrix.

**Value**

A list

- pts.approx a list of lower dimensional representations with respect to the original basis
- scores a matrix of scores.
- rss a vector of residual sums of squares.
- modes a list of modes of variation. The  $r$ th element of the list is the difference of rank  $r$  approximations to rank  $r - 1$  approximations.
- loadings a matrix of loading vectors.
- center mean of the data

**See Also**

[comp\\_pca\(\)](#), [psa\(\)](#)

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 comp\_pca

*Principal Component Analysis for PSA Comparison*


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**Description**

A wrapper function of `princomp()` for comparison to PSA. Manipulates result of `princomp()` in a similar format to `psa()`.

**Usage**

```
comp_pca(X)
```

**Arguments**

`X` a data matrix.

**Value**

A list

- `pts.approx` a list of lower dimensional representations with respect to the original basis
- `scores` a matrix of scores.
- `rss` a vector of residual sums of squares.
- `modes` a list of modes of variation. The  $r$ th element of the list is the difference of rank  $r$  approximations to rank  $r - 1$  approximations.
- `loadings` a matrix of loading vectors.
- `center` mean of the data

**See Also**

[psa\(\)](#)

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 comp\_power\_pca

*Power Transform Principal Component Analysis for PSA Comparison*


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**Description**

A wrapper function of `princomp()` applied to power transformed data.

**Usage**

```
comp_power_pca(X, alpha = 1/2)
```

**Arguments**

- `x` a data matrix.
- `alpha` a scalar  $\alpha > 0$  for power transform.

**Value**

A list

- `pts.approx` a list of lower dimensional representations with respect to the original basis
- `scores` a matrix of scores.
- `rss` a vector of residual sums of squares.
- `modes` a list of modes of variation. The  $r$ th element of the list is the difference of rank  $r$  approximations to rank  $r - 1$  approximations.
- `loadings` a matrix of loading vectors.
- `center mean` of the data

**See Also**

[comp\\_pca\(\)](#), [psa\(\)](#)

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loading\_bar

*Loading Bar Plots for PSA*

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**Description**

Creating a bar plot summarizing a loading vector

**Usage**

```
loading_bar(v, max.k = 12)
```

**Arguments**

- `v` a loading vector
- `max.k` maximum number of elements to display

**Value**

a ggplot object

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plotdendrogram2	<i>Plot a dendrogram of the merges.</i>
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**Description**

Converts results to a dendrogram (see [stats::as.dendrogram\(\)](#)) object and plots it.

**Usage**

```
plotdendrogram2(
  result,
  plot = TRUE,
  edge.root = TRUE,
  edgePar = list(p.lty = 0, t.cex = 1, t.adj = c(1, 1)),
  horiz = TRUE,
  nodeheight = "step",
  colors = NULL,
  ...
)
```

**Arguments**

result	The dendrogram.input value from <a href="#">psa()</a> .
plot	logical; if TRUE, plot the dendrogram output.
edge.root	logical; if TRUE, draw an edge to the root node.
edgePar	a list of plotting parameters for edges. See <a href="#">graphics::segments()</a> .
horiz	logical; if TRUE, draw the dendrogram horizontally.
nodeheight	a string specifying node height. "step" for heights given by the number of vertices after the merge or "rmse scores" for heights given by the RMSE of the scores of the merge.
colors	a vector of colors of nodes
...	Passed to <a href="#">graphics::plot()</a> .

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psa	<i>Principal Nested Simplices Analysis</i>
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**Description**

Estimate PSA-S or PSA-O of given data matrix.

**Usage**

```
psa(type, X, testweights = seq(0, 1, length.out = 100))
```

**Arguments**

type	s for PSA-S or o for PSA-O.
X	A data matrix.
testweights	A vector of weights to try.

**Value**

A list

- vertices a list of matrix representing vertices of the lower dimensional subsimplex. The  $r$ th element of the list corresponds to the rank  $r-1$  subsimplex.
- pts a list of lower dimensional representations with respect to the reduced basis vertices
- pts.approx a list of lower dimensional representations with respect to the original basis
- scores a matrix of scores.
- rss a vector of residual sums of squares.
- modes a list of modes of variation. The  $r$ th element of the list is the difference of rank  $r$  approximations to rank  $r-1$  approximations.
- loadings a matrix of loading vectors.
- const.info a data frame of merged vertices and merging weight at each merge.
- dendrogram.input additional information to apply `plotdendrogram2()`.

**See Also**

[plotdendrogram2\(\)](#)

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to\_simplex

*Map data to the simplex*

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**Description**

Map data to the simplex while preserving proportions.

**Usage**

```
to_simplex(X)
```

**Arguments**

`X` a data matrix

**Value**

a matrix of the same dimension as `X`

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