

Package ‘nipals’

October 13, 2022

Title Principal Components Analysis using NIPALS or Weighted EMPCA,
with Gram-Schmidt Orthogonalization

Version 0.8

Date 2021-09-14

Description Principal Components Analysis of a matrix using Non-linear Iterative Partial Least Squares or weighted Expectation Maximization PCA with Gram-Schmidt orthogonalization of the scores and loadings. Optimized for speed. See Andrecut (2009) <[doi:10.1089/cmb.2008.0221](https://doi.org/10.1089/cmb.2008.0221)>.

Depends R (>= 3.4.0)

License GPL-3

Encoding UTF-8

LazyData true

Suggests knitr, rmarkdown, testthat

URL <https://kwstat.github.io/nipals/>

BugReports <https://github.com/kwstat/nipals/issues>

VignetteBuilder knitr

RxygenNote 7.1.1

NeedsCompilation no

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empca	<i>Principal component analysis by weighted EMPCA, expectation maximization principal component-analysis</i>
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Description

Used for finding principal components of a numeric matrix. Missing values in the matrix are allowed. Weights for each element of the matrix are allowed. Principal Components are extracted one at a time. The algorithm computes $x = TP'$, where T is the 'scores' matrix and P is the 'loadings' matrix.

Usage

```
empca(
  x,
  w,
  ncomp = min(nrow(x), ncol(x)),
  center = TRUE,
  scale = TRUE,
  maxiter = 100,
  tol = 1e-06,
  seed = NULL,
  fitted = FALSE,
  gramschmidt = TRUE,
  verbose = FALSE
)
```

Arguments

<code>x</code>	Numerical matrix for which to find principal components. Missing values are allowed.
<code>w</code>	Numerical matrix of weights.
<code>ncomp</code>	Maximum number of principal components to extract from <code>x</code> .
<code>center</code>	If TRUE, subtract the mean from each column of <code>x</code> before PCA.
<code>scale</code>	If TRUE, divide the standard deviation from each column of <code>x</code> before PCA.
<code>maxiter</code>	Maximum number of EM iterations for each principal component.
<code>tol</code>	Default 1e-6 tolerance for testing convergence of the EM iterations for each principal component.
<code>seed</code>	Random seed to use when initializing the random rotation matrix.
<code>fitted</code>	Default FALSE. If TRUE, return the fitted (reconstructed) value of <code>x</code> .
<code>gramschmidt</code>	Default TRUE. If TRUE, perform Gram-Schmidt orthogonalization at each iteration.
<code>verbose</code>	Default FALSE. Use TRUE or 1 to show some diagnostics.

Value

A list with components eig, scores, loadings, fitted, ncomp, R2, iter, center, scale.

Author(s)

Kevin Wright

References

Stephen Bailey (2012). Principal Component Analysis with Noisy and/or Missing Data. Publications of the Astronomical Society of the Pacific. <http://doi.org/10.1086/668105>

Examples

```
B <- matrix(c(50, 67, 90, 98, 120,
             55, 71, 93, 102, 129,
             65, 76, 95, 105, 134,
             50, 80, 102, 130, 138,
             60, 82, 97, 135, 151,
             65, 89, 106, 137, 153,
             75, 95, 117, 133, 155), ncol=5, byrow=TRUE)
rownames(B) <- c("G1","G2","G3","G4","G5","G6","G7")
colnames(B) <- c("E1","E2","E3","E4","E5")
dim(B) # 7 x 5
p1 <- empca(B)
dim(p1$scores) # 7 x 5
dim(p1$loadings) # 5 x 5

B2 = B
B2[1,1] = B2[2,2] = NA
p2 = empca(B2, fitted=TRUE)
```

nipals

Principal component analysis by NIPALS, non-linear iterative partial least squares

Description

Used for finding principal components of a numeric matrix. Missing values in the matrix are allowed. Principal Components are extracted one at a time. The algorithm computes $x = TP'$, where T is the 'scores' matrix and P is the 'loadings' matrix.

Usage

```
nipals(
  x,
  ncomp = min(nrow(x), ncol(x)),
```

```

    center = TRUE,
    scale = TRUE,
    maxiter = 500,
    tol = 1e-06,
    startcol = 0,
    fitted = FALSE,
    force.na = FALSE,
    gramschmidt = TRUE,
    verbose = FALSE
)

```

Arguments

<code>x</code>	Numerical matrix for which to find principal components. Missing values are allowed.
<code>ncomp</code>	Maximum number of principal components to extract from <code>x</code> .
<code>center</code>	If TRUE, subtract the mean from each column of <code>x</code> before PCA.
<code>scale</code>	If TRUE, divide the standard deviation from each column of <code>x</code> before PCA.
<code>maxiter</code>	Maximum number of NIPALS iterations for each principal component.
<code>tol</code>	Default 1e-9 tolerance for testing convergence of the NIPALS iterations for each principal component.
<code>startcol</code>	Determine the starting column of <code>x</code> for the iterations of each principal component. If 0, use the column of <code>x</code> that has maximum absolute sum. If a number, use that column of <code>x</code> . If a function, apply the function to each column of <code>x</code> and choose the column with the maximum value of the function.
<code>fitted</code>	Default FALSE. If TRUE, return the fitted (reconstructed) value of <code>x</code> .
<code>force.na</code>	Default FALSE. If TRUE, force the function to use the method for missing values, even if there are no missing values in <code>x</code> .
<code>gramschmidt</code>	Default TRUE. If TRUE, perform Gram-Schmidt orthogonalization at each iteration.
<code>verbose</code>	Default FALSE. Use TRUE or 1 to show some diagnostics.

Details

The R2 values that are reported are marginal, not cumulative.

Value

A list with components `eig`, `scores`, `loadings`, `fitted`, `ncomp`, `R2`, `iter`, `center`, `scale`.

Author(s)

Kevin Wright

References

- Wold, H. (1966) Estimation of principal components and related models by iterative least squares. In Multivariate Analysis (Ed., P.R. Krishnaiah), Academic Press, NY, 391-420.
- Andrecut, Mircea (2009). Parallel GPU implementation of iterative PCA algorithms. Journal of Computational Biology, 16, 1593-1599.

Examples

```
B <- matrix(c(50, 67, 90, 98, 120,
             55, 71, 93, 102, 129,
             65, 76, 95, 105, 134,
             50, 80, 102, 130, 138,
             60, 82, 97, 135, 151,
             65, 89, 106, 137, 153,
             75, 95, 117, 133, 155), ncol=5, byrow=TRUE)
rownames(B) <- c("G1","G2","G3","G4","G5","G6","G7")
colnames(B) <- c("E1","E2","E3","E4","E5")
dim(B) # 7 x 5
p1 <- nipals(B)
dim(p1$scores) # 7 x 5
dim(p1$loadings) # 5 x 5

B2 = B
B2[1,1] = B2[2,2] = NA
p2 = nipals(B2, fitted=TRUE)

# Two ways to make a biplot

# method 1
biplot(p2$scores, p2$loadings)

# method 2
class(p2) <- "princomp"
p2$sdev <- sqrt(p2$eig)
biplot(p2, scale=0)
```

Description

U.S. Crime rates per 100,00 people for 7 categories in each of the 50 U.S. states in 1977.

Usage

uscrime

Format

A data frame with 50 observations on the following 8 variables.

state U.S. state
murder murders
rape rapes
robbery robbery
assault assault
burglary burglary
larceny larceny
autotheft automobile thefts

Details

There are two missing values.

Source

Documentation Example 3 for PROC HPPRINCOMP. <http://documentation.sas.com/api/docsets/statpug/14.2/content/statphpprincmpexamples.html#example3>

References

SAS/STAT User's Guide: High-Performance Procedures. The HPPRINCOMP Procedure. http://support.sas.com/documentation/cdl/en/statug/63080/HTML/default/index.html#page/_statug_hpprincmp_sect001.htm

Examples

```
library(nipals)
head(uscrime)

# SAS deletes rows with missing values
dat <- uscrime[complete.cases(uscrime), ]
dat <- as.matrix(dat[, -1])
m1 <- nipals(dat) # complete-data method

# Traditional NIPALS with missing data
dat <- uscrime
dat <- as.matrix(dat[, -1])
m2 <- nipals(dat, gramschmidt=FALSE) # missing
round(crossprod(m2$loadings),3) # Prin Comps not quite orthogonal

# Gram-Schmidt corrected NIPALS
m3 <- nipals(dat, gramschmidt=TRUE) # TRUE is default
round(crossprod(m3$loadings),3) # Prin Comps are orthogonal
```

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