

Package ‘ISR’

October 12, 2022

Title The Iterated Score Regression-Based Estimation Algorithm

Date 2022-04-22

Version 2022.4.22

Description Algorithm to handle with PCA-based missing data, where ISR is for PCA-based missing data with high correlation and DISR is for distributed PCA-based missing data. The philosophy of the package is described in Guo G. (2020) <[doi:10.1080/02331888.2020.1823979](https://doi.org/10.1080/02331888.2020.1823979)>.

License MIT + file LICENSE

Encoding UTF-8

LazyData true

RoxygenNote 7.1.1

Imports MASS, stats

Suggests testthat (>= 3.0.0)

Config/testthat/edition 3

Depends R (>= 3.5.0)

NeedsCompilation no

Author Guangbao Guo [aut, cre] (<<https://orcid.org/0000-0002-4115-6218>>),
Haoyue Song [aut],
Lixing Zhu [aut]

Maintainer Guangbao Guo <ggb11111111@163.com>

Repository CRAN

Date/Publication 2022-04-22 12:00:03 UTC

R topics documented:

CKD	2
DISR	3
HCV	4
ISR	4
Mean	5
MMLPCA	6

MNIPALS	7
MRPCA	8
orange	9
ozone	9
PM2.5	10
review	11
SR	12
Index	13

 CKD

CKD

Description

chronic kidney disease

Usage

```
data("CKD")
```

Format

The format is: num [1:400, 1:18] 48 7 62 48 51 60 68 24 52 53 ... - attr(*, "dimnames")=List of 2 ..\$: NULL ..\$: chr [1:18] "age" "bp" "sg" "al" ...

Details

There are 1010 missing values in the data set, accounting for 14.03 percent.

Source

Dr.P.Soundarapandian.M.D.,D.M (Senior Consultant Nephrologist), Apollo Hospitals, Managiri, Madurai Main Road, Karaikudi, Tamilnadu, Indi

References

Polat, H., Danaei-Mehr, H., and Cetin, A. (2017). Diagnosis of chronic kidney disease based on support vector machine by feature selection methods. Journal of Medical Systems, 41(4), 1-11.

Examples

```
data(CKD)
## maybe str(CKD) ; plot(CKD) ...
```

DISR

*Distributed iterated score regression***Description**

Calculate the estimator on the DISR method

Usage

```
DISR(data = 0, data0, real = TRUE, example = FALSE, D)
```

Arguments

<code>data</code>	is the original data set
<code>data0</code>	is the missing data set
<code>real</code>	is to judge whether the data set is a real missing data set
<code>example</code>	is to judge whether the data set is a simulation example
<code>D</code>	is the number of nodes

Value

<code>XDISR</code>	is the estimator on the DISR method
<code>MSEDISR</code>	is the MSE value of the DISR method
<code>MAEDISR</code>	is the MAE value of the DISR method
<code>REDISR</code>	is the RE value of the DISR method
<code>GCVDISR</code>	is the GCV value of the DISR method
<code>timeDISR</code>	is the time cost of the DISR method

Examples

```
library(MASS)
n=100;p=10;per=0.1
X0=data=matrix(mvrnorm(n*p,0,1),n,p)
m=round(per*n*p,digits=0)
mr=sample(1:(n*p),m,replace=FALSE)
X0[mr]=NA;data0=X0
DISR(data=data,data0=data0,real=FALSE,example=FALSE,D=2)
```

HCV

HCV

Description

Hepatitis C virus

Usage

```
data("HCV")
```

Format

The format is: num [1:615, 1:13] 1 1 1 1 1 1 1 1 1 ... - attr(*, "dimnames")=List of 2 ..\$: chr [1:615] "1" "2" "3" "4"\$: chr [1:13] "Category" "Age" "Sex" "ALB" ...

Details

There are 31 missing values in the data set, accounting for 0.39 percent.

Source

UCI repository

References

Lichtinghagen, R., Pietsch, D., Bantel, H., Manns, M., Brand, K. and Bahr, Matthias. (2013). The Enhanced Liver Fibrosis (ELF) Score: Normal Values, Influence Factors and Proposed Cut-Off Values.. *Journal of hepatology*. 59. 236-242.

Examples

```
data(HCV)
## maybe str(HCV) ; plot(HCV) ...
```

ISR

Iterated score regression

Description

Calculate the estimator on the ISR method

Usage

```
ISR(data = 0, data0, real = TRUE, example = FALSE)
```

Arguments

data	is the original data set
data0	is the missing data set
real	is to judge whether the data set is a real missing data set
example	is to judge whether the data set is a simulation example

Value

XISR	is the estimator on the ISR method
MSEISR	is the MSE value of the ISR method
MAEISR	is the MAE value of the ISR method
REISR	is the RE value of the ISR method
GCVISR	is the GCV value of the ISR method
timeISR	is the time cost of the ISR method

Examples

```
library(MASS)
n=100;p=10;per=0.1
X0=data=matrix(mvrnorm(n*p,0,1),n,p)
m=round(per*n*p,digits=0)
mr=sample(1:(n*p),m,replace=FALSE)
X0[mr]=NA;data0=X0
ISR(data=data,data0=data0,real=FALSE,example=FALSE)
```

Mean

Mean method

Description

Calculate the estimator on the Mean method

Usage

```
Mean(data = 0, data0, real = TRUE, example = FALSE)
```

Arguments

data	is the original data set
data0	is the missing data set
real	is to judge whether the data set is a real missing data set
example	is to judge whether the data set is a simulation example

Value

XMean	is the estimator on the Mean method
MSEMean	is the MSE value of the Mean method
MAEMean	is the MAE value of the Mean method
REMean	is the RE value of the Mean method
GCVMean	is the GCV value of the Mean method
timeMean	is the time cost of the Mean method

Examples

```
library(MASS)
n=100;p=10;per=0.1
X0=data=matrix(mvrnorm(n*p,0,1),n,p)
m=round(per*n*p,digits=0)
mr=sample(1:(n*p),m,replace=FALSE)
X0[mr]=NA;data0=X0
Mean(data=data,data0=data0,real=FALSE,example=FALSE)
```

MMLPCA

*Modified maximum likelihood principal component analysis***Description**

Calculate the estimator on the ISR method

Usage

```
MMLPCA(data = 0, data0, real = TRUE, example = FALSE)
```

Arguments

data	is the original data set
data0	is the missing data set
real	is to judge whether the data set is a real missing data set
example	is to judge whether the data set is a simulation example

Value

XMMLPCA	is the estimator on the MMLPCA method
MSEMMLPCA	is the MSE value of the MMLPCA method
MAEMMLPCA	is the MAE value of the MMLPCA method
REMMLPCA	is the RE value of the MMLPCA method
GCVMMLPCA	is the GCV value of the MMLPCA method
timeMMLPCA	is the time cost of the MMLPCA method

Examples

```

library(MASS)
n=100;p=10;per=0.1
X0=data=matrix(mvrnorm(n*p,0,1),n,p)
m=round(per*n*p,digits=0)
mr=sample(1:(n*p),m,replace=FALSE)
X0[mr]=NA;data0=X0
MMLPCA(data=data,data0=data0,real=FALSE,example=FALSE)

```

MNIPALS

*Modified nonlinear iterative partial least squares method***Description**

Calculate the estimator on the MNIPALS method

Usage

```
MNIPALS(data = 0, data0, real = TRUE, example = FALSE)
```

Arguments

data	is the original data set
data0	is the missing data set
real	is to judge whether the data set is a real missing data set
example	is to judge whether the data set is a simulation example

Value

XMNIPALS	is the estimator on the MNIPALS method
MSEMNIPALS	is the MSE value of the MNIPALS method
MAEMNIPALS	is the MAE value of the MNIPALS method
REMNIPALS	is the RE value of the MNIPALS method
GVMNIPALS	is the GCV value of the MNIPALS method
timeMNIPALS	is the time cost of the MNIPALS method

Examples

```

library(MASS)
n=100;p=10;per=0.1
X0=data=matrix(mvrnorm(n*p,0,1),n,p)
m=round(per*n*p,digits=0)
mr=sample(1:(n*p),m,replace=FALSE)
X0[mr]=NA;data0=X0
MNIPALS(data=data,data0=data0,real=FALSE,example=FALSE)

```

 MRPCA

Modified regularized PCA

Description

Calculate the estimator on the MRPCA method

Usage

```
MRPCA(data = 0, data0, real = TRUE, example = FALSE)
```

Arguments

data	is the original data set
data0	is the missing data set
real	is to judge whether the data set is a real missing data set
example	is to judge whether the data set is a simulation example

Value

XMRPCA	is the estimator on the MRPCA method
MSEMRPCA	is the MSE value of the MRPCA method
MAEMRPCA	is the MAE value of the MRPCA method
REMRPCA	is the RE value of the MRPCA method
GCVMRPCA	is the GCV value of the MRPCA method
timeMRPCA	is the time cost of the MRPCA method

Examples

```
library(MASS)
n=100;p=10;per=0.1
X0=data=matrix(mvrnorm(n*p,0,1),n,p)
m=round(per*n*p,digits=0)
mr=sample(1:(n*p),m,replace=FALSE)
X0[mr]=NA;data0=X0
MRPCA(data=data,data0=data0,real=FALSE,example=FALSE)
```

orange	<i>orange</i>
--------	---------------

Description

orange

Usage

```
data("orange")
```

Format

The format is: num [1:12, 1:8] 4.79 4.58 4.71 6.58 NA ... - attr(*, "dimnames")=List of 2 ..\$: chr [1:12] "1" "2" "3" "4"\$: chr [1:8] "Color.intensity" "Odor.intensity" "Attack.intensity" "Sweet" ...

Details

There are 19 missing values in the data set, accounting for 19.79 percent.

Source

<http://factominer.free.fr/missMDA/index.html>

References

Josse J, Husson F (2016). missMDA: A Package for Handling Missing Values in Multivariate Data Analysis. *Journal of Statistical Software*, 70(1), 1–31.

Examples

```
data(orange)
## maybe str(orange) ; plot(orange) ...
```

ozone	<i>ozone</i>
-------	--------------

Description

ozone

Usage

```
data("ozone")
```

Format

A data frame with 112 observations on the following 11 variables.

max03 a numeric vector

T9 a numeric vector

T12 a numeric vector

T15 a numeric vector

Ne9 a numeric vector

Ne12 a numeric vector

Ne15 a numeric vector

Vx9 a numeric vector

Vx12 a numeric vector

Vx15 a numeric vector

max03v a numeric vector

Details

There are 115 missing values in it, accounting for 9.96 percent.

Source

<http://factominer.free.fr/missMDA/index.html>

References

Audigier, V., Husson, F., and Josse, J. (2014). A principal components method to impute missing values for mixed data. *Advances in Data Analysis and Classification*, 10(1), 5-26.

Examples

```
data(ozone)
## maybe str(ozone) ; plot(ozone) ...
```

PM2.5

PM2.5

Description

Beijing PM2.5

Usage

```
data("PM2.5")
```

Format

The format is: num [1:43824, 1:12] 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 ... - attr(*, "dimnames")=List of 2 ..\$: chr [1:43824] "1" "2" "3" "4"\$: chr [1:12] "year" "month" "day" "hour" ...

Details

It records 43824 daily measurements on 12 variables and there are 2067 missing values on 2067 measurements, accounting for 0.00393.

Source

UCI repository

References

X. Liang, T. Zou, B. Guo, S. Li, H. Zhang, S. Zhang, H. Huang, and S. Chen. Assessing Beijing's PM2.5 pollution: severity, weather impact, APEC and winter heating. *Proceedings of the Royal Society A*, 471(2182):1–20, 2015.

Examples

```
data(PM2.5)
## maybe str(PM2.5) ; plot(PM2.5) ...
```

review

review

Description

Travel reviews

Usage

```
data("review")
```

Format

The format is: num [1:980, 1:10] 0.93 1.02 1.22 0.45 0.51 0.99 0.9 0.74 1.12 0.7 ... - attr(*, "dimnames")=List of 2 ..\$: chr [1:980] "User_1" "User_2" "User_3" "User_4"\$: chr [1:10] "Category_1" "Category_2" "Category_3" "Category_4" ...

Details

980 travelers' reviews of 10 different types of travel facilities in East Asia

Source

UCI repository

References

Renjith, S., Sreekumar, A., and Jathavedan, M. (2018). Evaluation of partitioning clustering algorithms for processing social media data in tourism domain. 2018 IEEE Recent Advances in Intelligent Computational Systems (RAICS), 127-131.

Examples

```
data(review)
## maybe str(review) ; plot(review) ...
```

 SR

Score regression

Description

Calculate the estimator on the SR method

Usage

```
SR(data = 0, data0, real = TRUE, example = FALSE)
```

Arguments

data	is the original data set
data0	is the missing data set
real	is to judge whether the data set is a real missing data set
example	is to judge whether the data set is a simulation example

Value

XSR	is the estimator on the SR method
MSESR	is the MSE value of the SR method
MAESR	is the MAE value of the SR method
RESR	is the RE value of the SR method
GCVSR	is the GCV value of the SR method
timeSR	is the time cost of the SR method

Examples

```
library(MASS)
n=100;p=10;per=0.1
X0=data=matrix(mvrnorm(n*p,0,1),n,p)
m=round(per*n*p,digits=0)
mr=sample(1:(n*p),m,replace=FALSE)
X0[mr]=NA;data0=X0
SR(data=data,data0=data0,real=FALSE,example=FALSE)
```

Index

* datasets

- CKD, [2](#)
 - HCV, [4](#)
 - orange, [9](#)
 - ozone, [9](#)
 - PM2.5, [10](#)
 - review, [11](#)
-
- CKD, [2](#)
 - DISR, [3](#)
 - HCV, [4](#)
 - ISR, [4](#)
 - Mean, [5](#)
 - MMLPCA, [6](#)
 - MNIPALS, [7](#)
 - MRPCA, [8](#)
 - orange, [9](#)
 - ozone, [9](#)
 - PM2.5, [10](#)
 - review, [11](#)
 - SR, [12](#)