

Package ‘ISR’

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Title The Iterated Score Regression-Based Estimation

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Description We use the ISR to handle with PCA-based missing data with high correlation, and the DISR to handle with distributed PCA-based missing data. The philosophy of the package is described in Guo G. (2024) <[doi:10.1080/03610918.2022.2091779](https://doi.org/10.1080/03610918.2022.2091779)>.

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NeedsCompilation no

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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

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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*Caculate the estimator with the DISR method*

Description

Caculate the estimator with the DISR method

Usage

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

Arguments

| | |
|---------|---|
| data | is the orignal 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 |
| D | is the number of nodes |

Value

| | |
|----------|-------------------------------------|
| XDISR | is the estimator on the DISR method |
| MSEDISR | is the MSE value of the DISR method |
| MAEDISR | is the MAE value of the DISR method |
| REDISR | is the RE value of the DISR method |
| GCVDISR | is the GCV value of the DISR method |
| timeDISR | 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 | <i>HCV</i> |
|-----|------------|

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

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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 | <i>Caculate the estimator with the ISR method</i> |
|-----|---|

Description

Caculate the estimator with the ISR method

Usage

```
ISR(data, 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

*Caculate the estimator on the Mean method***Description**

Caculate the estimator on the Mean method

Usage

Mean(data, 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

*Caculate the estimator on the MMLPCA method***Description**

Caculate the estimator on the MMLPCA method

Usage

```
MMLPCA(data, 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

Caculate the estimator on the MNIPALS method

Description

Caculate the estimator on the MNIPALS method

Usage

```
MNIPALS(data, 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
- GCVMNIPALS 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

*Caculate the estimator on the MRPCA method***Description**

Caculate the estimator on the MRPCA method

Usage

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

Arguments

| | |
|---------|---|
| data | is the orignal 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)
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**orange*

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**ozone*

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) ...
```

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

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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

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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

Caculate the estimator on the SR method

Description

Caculate the estimator on the SR method

Usage

```
SR(data, 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 |

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)
```

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