

Package: mispr (via r-universe)

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Type Package

Title Multiple Imputation with Sequential Penalized Regression

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Author Faisal Maqbool Zahid

Maintainer Faisal Maqbool Zahid <faisalmz99@yahoo.com>

Description Generates multivariate imputations using sequential regression with L2 penalty. For more details see Zahid and Heumann (2018) <[doi:10.1177/0962280218755574](https://doi.org/10.1177/0962280218755574)>.

Depends R (>= 3.3.0)

Imports e1071, MASS, penalized, stats

License GPL-2

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LazyData true

RoxygenNote 6.0.1

NeedsCompilation no

Repository CRAN

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`data1`*Simulated data with 50 covariates*

Description

`data1` artificially generated dataframe with $n=100$ and $p=50$. Missing values using MAR (missing at random) mechanism are artificially generated in 10 covariates.

Usage

```
data(data1)
```

Format

An object of class `data.frame` with 100 rows and 51 columns.

Examples

```
data(data1)
```

`data2`*Simulated data with 200 covariates*

Description

`data2` artificially generated dataframe with $n=100$ and $p=200$. Missing values using MAR (missing at random) mechanism are artificially generated in 10 covariates.

Usage

```
data(data2)
```

Format

An object of class `data.frame` with 100 rows and 201 columns.

Examples

```
data(data2)
```

Description

Generates Multivariate Imputations using sequential regression with L2 penalization.

Usage

```
mispr(x, x.select = FALSE, pen = FALSE, maxit = 5, m = 5,
      track = FALSE, init.method = "random", L2.fix = NULL, cv = TRUE,
      maxL2 = 2^10)
```

Arguments

<code>x</code>	A data frame or a matrix containing the incomplete data. Missing values are coded as NA.
<code>x.select</code>	A Boolean flag. If TRUE, linearly dependent columns will be removed before fitting of each imputation model. If FALSE, the linearly dependent columns will be removed only when number of predictors is greater than the sample size for fitting an imputation model. The default is FALSE.
<code>pen</code>	A Boolean flag. If TRUE, each imputation model will be fitted with L2 penalty. If FALSE, maximum likelihood estimation (MLE) will be used. However, if MLE fails, L2 penalty is used for fitting the imputation model. The default is FALSE.
<code>maxit</code>	A scalar giving the number of iterations. The default is 5.
<code>m</code>	Number of multiple imputations. The default is $m=5$.
<code>track</code>	A Boolean flag. If TRUE, <code>mispr</code> will print additional information about iterations on console. The default is FALSE for silent computation.
<code>init.method</code>	Method for initialization of missing values. <code>random</code> for filling NA in each column with a random sample from the observed values of that column. <code>median</code> for mean imputation.
<code>L2.fix</code>	Fixed value of ridge penalty (optional) to use for each imputation model. For default i.e., NULL, L2 penalty will be decided with k-fold cross-validation.
<code>cv</code>	A Boolean flag. If TRUE that is default, optimal value of L2 penalty will be decided independently for each imputation model using 5-fold cross-validation.
<code>maxL2</code>	The maximum value of the tuning parameter for L2 penalization to be used for optimizing the cross-validated likelihood. Default value is 2^{10} .

Details

Generates multiple imputations for incomplete multivariate data by fitting a sequence of regression models using L2 penalty iteratively. Missing data can occur in one or more variables of the data. In each step of the iteration, ridge regression is fitted according to the distributional form of the missing variable taken as a response. All other variables are taken as predictors. If some predictors are incomplete, the most #’recently generated imputations are used to complete the predictors before using them as a predictor.

Value

a list containing the number of imputed datasets, number of iterations used to obtain imputed data, list of multiply imputed datasets, and summary of missing values.

Author(s)

Faisal Maqbool Zahid <faisalmz99@yahoo.com>.

References

Zahid, F. M., and Heumann, C. (2018). Multiple imputation with sequential penalized regression. *Statistical Methods in Medical Research*, 0962280218755574.

Examples

```
data(data1)
# Select a subset of data1
x=data1[ , 1:10]
res1 = mispr(x)
# to get 3 multiply imputed datasets
res2 = mispr(x, m=3)
```

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* **datasets**

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