

Package ‘misaem’

October 13, 2022

Title Linear Regression and Logistic Regression with Missing Covariates

Version 1.0.1

Date 2021-04-07

Description Estimate parameters of linear regression and logistic regression with missing covariates with missing data, perform model selection and prediction, using EM-type algorithms. Jiang W., Josse J., Lavielle M., TraumaBase Group (2020) <[doi:10.1016/j.csda.2019.106907](https://doi.org/10.1016/j.csda.2019.106907)>.

Depends R (>= 3.4.0)

Encoding UTF-8

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URL <https://github.com/julierennes/misaem>

Imports mvtnorm, stats, MASS, norm, methods

Suggests knitr, rmarkdown, mice

LazyData false

VignetteBuilder knitr

RoxygenNote 7.1.1

NeedsCompilation no

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Repository CRAN

Date/Publication 2021-04-12 08:10:02 UTC

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combinations	<i>combinations</i>
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Description

Given all the possible patterns of missingness.

Usage

```
combinations(p)
```

Arguments

p Dimension of covariates.

Value

A matrix containing all the possible missing patterns. Each row indicates a pattern of missingness. "1" means "observed", 0 means "missing".

Examples

```
comb = combinations(5)
```

imputeEllP*Function for imputing single point for linear regression model*

Description

Function for imputing single point for linear regression model

Usage

```
imputeEllP(point, Sigma.inv)
```

Arguments

point	A single observation containing missing values.
Sigma.inv	Inverse of estimated Σ .

Value

Imputed observation.

likelihood_saem*likelihood_saem*

Description

Used in main function miss.saem. Calculate the observed log-likelihood for logistic regression model with missing data, using Monte Carlo version of Louis formula.

Usage

```
likelihood_saem(
  beta,
  mu,
  Sigma,
  Y,
  X.obs,
  rindic = as.matrix(is.na(X.obs)),
  whichcolXmissing = (1:ncol(rindic))[apply(rindic, 2, sum) > 0],
  mc.size = 2
)
```

Arguments

<code>beta</code>	Estimated parameter of logistic regression model.
<code>mu</code>	Estimated parameter μ .
<code>Sigma</code>	Estimated parameter Σ .
<code>Y</code>	Response vector $N \times 1$
<code>X.obs</code>	Design matrix with missingness $N \times p$
<code>rindic</code>	Missing pattern of <code>X.obs</code> . If a component in <code>X.obs</code> is missing, the corresponding position in <code>rindic</code> is 1; else 0.
<code>whichcolXmissing</code>	The column index in covariate containing at least one missing observation.
<code>mc.size</code>	Monte Carlo sampling size.

Value

Observed log-likelihood.

Examples

```
# Generate dataset
N <- 50 # number of subjects
p <- 3 # number of explanatory variables
mu.star <- rep(0,p) # mean of the explanatory variables
Sigma.star <- diag(rep(1,p)) # covariance
beta.star <- c(1, 1, 0) # coefficients
beta0.star <- 0 # intercept
beta.true = c(beta0.star,beta.star)
X.complete <- matrix(rnorm(N*p), nrow=N) %*% chol(Sigma.star) +
  matrix(rep(mu.star,N), nrow=N, byrow = TRUE)
p1 <- 1/(1+exp(-X.complete%*%beta.star-beta0.star))
y <- as.numeric(runif(N)<p1)
# Generate missingness
p.miss <- 0.10
patterns <- runif(N*p)<p.miss #missing completely at random
X.obs <- X.complete
X.obs[patterns] <- NA

# Observed log-likelihood
ll_obs = likelihood_saem(beta.true,mu.star,Sigma.star,y,X.obs)
```

Description

Calculate the likelihood or log-likelihood for one observation of logistic regression model .

Usage

```
log_reg(y, x, beta, iflog = TRUE)
```

Arguments

y	Response value (0 or 1).
x	Covariate vector of dimension $p \times 1$.
beta	Estimated parameter of logistic regression model.
iflog	If TRUE, log_reg calculate the log-likelihood; else likelihood.

Value

Likelihood or log-likelihood.

Examples

```
res = log_reg(1,c(1,2,3),c(1,-1,1))
```

louis_lr_saem

louis_lr_saem

Description

Used in main function miss.saem. Calculate the variance of estimated parameters for logistic regression model with missing data, using Monte Carlo version of Louis formula.

Usage

```
louis_lr_saem(
  beta,
  mu,
  Sigma,
  Y,
  X.obs,
  pos_var = 1:ncol(X.obs),
  rindic = as.matrix(is.na(X.obs)),
  whichcolXmissing = (1:ncol(rindic))[apply(rindic, 2, sum) > 0],
  mc.size = 2
)
```

Arguments

beta	Estimated parameter of logistic regression model.
mu	Estimated parameter μ .
Sigma	Estimated parameter Σ .
Y	Response vector $N \times 1$
X.obs	Design matrix with missingness $N \times p$
pos_var	Index of selected covariates.
rindic	Missing pattern of X.obs. If a component in X.obs is missing, the corresponding position in rindic is 1; else 0.
whichcolXmissing	The column index in covariate containing at least one missing observation.
mc.size	Monte Carlo sampling size.

Value

Variance of estimated β .

Examples

```
# Generate dataset
N <- 50 # number of subjects
p <- 3 # number of explanatory variables
mu.star <- rep(0,p) # mean of the explanatory variables
Sigma.star <- diag(rep(1,p)) # covariance
beta.star <- c(1, 1, 0) # coefficients
beta0.star <- 0 # intercept
beta.true = c(beta0.star,beta.star)
X.complete <- matrix(rnorm(N*p), nrow=N)%*%chol(Sigma.star) +
  matrix(rep(mu.star,N), nrow=N, byrow = TRUE)
p1 <- 1/(1+exp(-X.complete%*%beta.star-beta0.star))
y <- as.numeric(runif(N)<p1)
# Generate missingness
p.miss <- 0.10
patterns <- runif(N*p)<p.miss #missing completely at random
X.obs <- X.complete
X.obs[patterns] <- NA

# Louis formula to obtain variance of estimates
V_obs = louis_lr_saem(beta.true,mu.star,Sigma.star,y,X.obs)
```

Description

This function is used to perform statistical inference for logistic regression model with missing values, by algorithm SAEM.

Usage

```
miss.glm(formula, data, control = list(...), ...)
```

Arguments

formula	an object of class " formula " : a symbolic description of the logistic regression model to be fitted.
data	an optional data frame containing the variables in the model. If not found in data, the variables are taken from <code>environment(formula)</code> , typically the environment from which <code>miss.glm</code> is called.
control	a list of parameters for controlling the fitting process. For <code>miss.glm.fit</code> this is passed to miss.glm.control .
...	arguments to be used to form the default control argument if it is not supplied directly.

Value

An object of class "`miss.glm`": a list with following components:

coefficients	Estimated β .
ll	Observed log-likelihood.
var.covar	Variance-covariance matrix for estimated parameters.
s.err	Standard error for estimated parameters.
mu.X	Estimated μ .
Sig.X	Estimated Σ .
call	the matched call.
formula	the formula supplied.

Examples

```
# Generate dataset
N <- 100 # number of subjects
p <- 3 # number of explanatory variables
mu.star <- rep(0,p) # mean of the explanatory variables
Sigma.star <- diag(rep(1,p)) # covariance
beta.star <- c(1, 1, 0) # coefficients
beta0.star <- 0 # intercept
beta.true = c(beta0.star,beta.star)
X.complete <- matrix(rnorm(N*p), nrow=N) %*% chol(Sigma.star) +
  matrix(rep(mu.star,N), nrow=N, byrow = TRUE)
p1 <- 1/(1+exp(-X.complete%*%beta.star-beta0.star))
y <- as.numeric(runif(N)<p1)
```

```

# Generate missingness
p.miss <- 0.10
patterns <- runif(N*p)<=p.miss #missing completely at random
X.obs <- X.complete
X.obs[patterns] <- NA

df.obs = data.frame(y,X.obs)

# SAEM
miss.list = miss.glm(y~., data=df.obs, print_iter=FALSE, seed=100)
print(miss.list)
print(summary(miss.list))
summary(miss.list)$coef

```

miss.glm.control *Auxiliary for Controlling Fitting*

Description

Auxiliary function for [miss.glm](#) fitting. Typically only used internally by [miss.glm.fit](#).

Usage

```

miss.glm.control(
  maxruns = 500,
  tol_em = 1e-07,
  nmcmc = 2,
  tau = 1,
  k1 = 50,
  subsets = NA,
  seed = NA,
  print_iter = TRUE,
  var_cal = TRUE,
  ll_obs_cal = TRUE
)

```

Arguments

<code>maxruns</code>	maximum number of iterations. The default is <code>maxruns = 500</code> .
<code>tol_em</code>	the tolerance to stop SAEM. The default is <code>tol_em = 1e-7</code> .
<code>nmcmc</code>	the MCMC length. The default is <code>nmcmc = 2</code> .
<code>tau</code>	rate τ in the step size $(k - k_1)^{-\tau}$. The default is <code>tau = 1</code> .
<code>k1</code>	number of first iterations k_1 in the step size $(k - k_1)^{-\tau}$. The default is <code>k1=50</code> .
<code>subsets</code>	Index of selected covariates if any. The default is all the covariates.
<code>seed</code>	an integer as a seed set for the random generator.

<code>print_iter</code>	logical indicating if output should be produced for each iteration.
<code>var_cal</code>	logical indicating if the variance of estimated parameters should be calculated.
<code>ll_obs_cal</code>	logical indicating if the observed log-likelihood should be calculated.

Value

A list with components named as the arguments.

Examples

```
## For examples see example(miss.glm)
```

`miss.glm.fit`

Fitting Logistic Regression Models with Missing Values

Description

This function is used inside `miss.glm` to fit logistic regression model with missing values, by algorithm SAEM.

Usage

```
miss.glm.fit(x, y, control = list())
```

Arguments

<code>x</code>	design matrix with missingness $N \times p$.
<code>y</code>	response vector $N \times 1$.
<code>control</code>	a list of parameters for controlling the fitting process. For <code>miss.glm.fit</code> this is passed to <code>miss.glm.control</code> .

Value

a list with following components:

<code>coefficients</code>	Estimated β .
<code>ll</code>	Observed log-likelihood.
<code>var.covar</code>	Variance-covariance matrix for estimated parameters.
<code>s.err</code>	Standard error for estimated parameters.
<code>mu.X</code>	Estimated μ .
<code>Sig.X</code>	Estimated Σ .

Examples

```
## For examples see example(miss.glm)
```

miss.glm.model.select miss.glm.model.select

Description

Model selection for the logistic regression model with missing data.

Usage

```
miss.glm.model.select(Y, X, seed = NA)
```

Arguments

Y	Binary response vector $N \times 1$
X	Design matrix with missingness $N \times p$
seed	An integer as a seed set for the random generator. The default value is 200.

Value

An object of class "miss.glm".

Examples

```
# Generate dataset
N <- 40 # number of subjects
p <- 3 # number of explanatory variables
mu.star <- rep(0,p) # mean of the explanatory variables
Sigma.star <- diag(rep(1,p)) # covariance
beta.star <- c(1, 1, 0) # coefficients
beta0.star <- 0 # intercept
beta.true = c(beta0.star,beta.star)
X.complete <- matrix(rnorm(N*p), nrow=N) %*% chol(Sigma.star) +
  matrix(rep(mu.star,N), nrow=N, byrow = TRUE)
p1 <- 1/(1+exp(-X.complete%*%beta.star-beta0.star))
Y <- as.numeric(runif(N)<p1)
# Generate missingness
p.miss <- 0.10
patterns <- runif(N*p)<=p.miss #missing completely at random
X <- X.complete
X[patterns] <- NA
# model selection for SAEM
miss.model = miss.glm.model.select(Y,X,seed=100)
print(miss.model)
```

Description

This function is used to perform statistical inference for linear regression model with missing values, by algorithm EM.

Usage

```
miss.lm(formula, data, control = list(...), ...)
```

Arguments

- | | |
|---------|---|
| formula | an object of class " formula " : a symbolic description of the linear regression model to be fitted. |
| data | an optional data frame containing the variables in the model. If not found in <code>data</code> , the variables are taken from <code>environment(formula)</code> , typically the environment from which <code>miss.lm</code> is called. |
| control | a list of parameters for controlling the fitting process. For <code>miss.lm.fit</code> this is passed to miss.lm.control . |
| ... | arguments to be used to form the default control argument if it is not supplied directly. |

Value

An object of class "miss.lm": a list with following components:

- | | |
|--------------|--|
| coefficients | Estimated β . |
| ll | Observed log-likelihood. |
| s.resid | Estimated standard error for residuals. |
| s.err | Standard error for estimated parameters. |
| mu.X | Estimated μ . |
| Sig.X | Estimated Σ . |
| call | the matched call. |
| formula | the formula supplied. |

Examples

```
# Generate complete data
set.seed(1)
mu.X <- c(1, 1)
Sigma.X <- matrix(c(1, 1, 1, 4), nrow = 2)
n <- 50
p <- 2
```

```

X.complete <- matrix(rnorm(n*p), nrow=n) %*% chol(Sigma.X) +
  matrix(rep(mu.X, n), nrow=n, byrow = TRUE)
b <- c(2, 3, -1)
sigma.eps <- 0.25
y <- cbind(rep(1, n), X.complete) %*% b + rnorm(n, 0, sigma.eps)

# Add missing values
p.miss <- 0.10
patterns <- runif(n*p) < p.miss #missing completely at random
X.obs <- X.complete
X.obs[patterns] <- NA

# Estimate regression using EM
df.obs = data.frame(y, X.obs)
miss.list = miss.lm(y~, data=df.obs)
print(miss.list)
print(summary(miss.list))
summary(miss.list)$coef

```

miss.lm.control *Auxiliary for Controlling Fitting*

Description

Auxiliary function for **miss.lm** fitting. Typically only used internally by **miss.lm.fit**.

Usage

```
miss.lm.control(maxruns = 500, tol_em = 1e-07, print_iter = TRUE)
```

Arguments

maxruns	maximum number of iterations. The default is maxruns = 500.
tol_em	the tolerance to stop EM. The default is tol_em = 1e-4.
print_iter	logical indicating if output should be produced for each iteration.

Value

A list with components named as the arguments.

Examples

```
## For examples see example(miss.lm)
```

miss.lm.fit*Fitting Linear Regression Model with Missing Values*

Description

This function is used inside `miss.lm` to fit linear regression model with missing values, by EM algorithm.

Usage

```
miss.lm.fit(x, y, control = list())
```

Arguments

- | | |
|----------------------|---|
| <code>x</code> | design matrix with missingness $N \times p$. |
| <code>y</code> | response vector $N \times 1$. |
| <code>control</code> | a list of parameters for controlling the fitting process. For <code>miss.lm.fit</code> this is passed to <code>miss.lm.control</code> . |

Value

a list with following components:

- | | |
|---------------------------|--|
| <code>coefficients</code> | Estimated β . |
| <code>ll</code> | Observed log-likelihood. |
| <code>s.resid</code> | Estimated standard error for residuals. |
| <code>s.err</code> | Standard error for estimated parameters. |
| <code>mu.X</code> | Estimated μ . |
| <code>Sig.X</code> | Estimated Σ . |

Examples

```
## For examples see example(miss.lm)
```

miss.lm.model.select *miss.lm.model.select*

Description

Model selection for the linear regression model with missing data.

Usage

```
miss.lm.model.select(Y, X)
```

Arguments

Y	Response vector $N \times 1$
X	Design matrix with missingness $N \times p$

Value

An object of class "miss.lm".

Examples

```
# Generate complete data
set.seed(1)
mu.X <- c(1, 1)
Sigma.X <- matrix(c(1, 1, 1, 4), nrow = 2)
n <- 50
p <- 2
X.complete <- matrix(rnorm(n*p), nrow=n) %*% chol(Sigma.X) +
  matrix(rep(mu.X,n), nrow=n, byrow = TRUE)
b <- c(2, 0, -1)
sigma.eps <- 0.25
y <- cbind(rep(1, n), X.complete) %*% b + rnorm(n, 0, sigma.eps)

# Add missing values
p.miss <- 0.10
patterns <- runif(n*p)<=p.miss #missing completely at random
X.obs <- X.complete
X.obs[patterns] <- NA

# model selection
miss.model = miss.lm.model.select(y, X.obs)
print(miss.model)
```

predict.miss.glm	<i>Prediction on test with missing values for the logistic regression model.</i>
------------------	--

Description

Prediction on test with missing values for the logistic regression model.

Usage

```
## S3 method for class 'miss.glm'  
predict(object, newdata = NULL, seed = NA, method = "map", ...)
```

Arguments

- object a fitted object of class inheriting from "miss.glm".
 newdata a data frame in which to look for variables with which to predict. It can contain missing values.
 seed An integer as a seed set for the random generator.
 method The name of method to deal with missing values in test set. It can be 'map' (maximum a posteriori) or 'impute' (imputation by conditional expectation). Default is 'map'.
 ... Further arguments passed to or from other methods.

Value

- pr.saem The prediction result for logistic regression: the probability of response y=1.

Examples

```
# Generate dataset  

N <- 100 # number of subjects  

p <- 3 # number of explanatory variables  

mu.star <- rep(0,p) # mean of the explanatory variables  

Sigma.star <- diag(rep(1,p)) # covariance  

beta.star <- c(1, 1, 0) # coefficients  

beta0.star <- 0 # intercept  

beta.true = c(beta0.star,beta.star)  

X.complete <- matrix(rnorm(N*p), nrow=N)%*%chol(Sigma.star) +  

  matrix(rep(mu.star,N), nrow=N, byrow = TRUE)  

p1 <- 1/(1+exp(-X.complete%*%beta.star-beta0.star))  

y <- as.numeric(runif(N)<p1)  

# Generate missingness  

p.miss <- 0.10  

patterns <- runif(N*p)< p.miss #missing completely at random  

X.obs <- X.complete  

X.obs[patterns] <- NA
```

```

df.obs = data.frame(y,X.obs)

# SAEM
miss.list = miss.glm(y~, data=df.obs, print_iter=FALSE,seed=100)

# Generate new dataset for prediction
Nt <- 20
Xt <- matrix(rnorm(Nt*p), nrow=Nt) %*% chol(Sigma.star) +
  matrix(rep(mu.star,Nt), nrow=Nt, byrow = TRUE)
# Generate missingness in new dataset
patterns <- runif(Nt*p)< p.miss
Xt.obs <- Xt
Xt.obs[patterns] <- NA

# Prediction with missing values
miss.prob = predict(miss.list, data.frame(Xt.obs), method='map')
print(miss.prob)

```

`predict.miss.lm`*Prediction on test with missing values for the linear regression model.***Description**

Prediction on test with missing values for the linear regression model.

Usage

```
## S3 method for class 'miss.lm'
predict(object, newdata = NULL, seed = NA, ...)
```

Arguments

- `object` a fitted object of class inheriting from "miss.lm".
- `newdata` a data frame in which to look for variables with which to predict. It can contain missing values.
- `seed` An integer as a seed set for the random generator.
- `...` Further arguments passed to or from other methods.

Value

- | | |
|-------------------|--|
| <code>pr.y</code> | The prediction result for linear regression. |
|-------------------|--|

Examples

```

# Generate complete data
set.seed(1)
mu.X <- c(1, 1)
Sigma.X <- matrix(c(1, 1, 1, 4), nrow = 2)
n <- 50 # train set size
p <- 2 # number of covariates
X.complete <- matrix(rnorm(n*p), nrow=n) %*% chol(Sigma.X) +
  matrix(rep(mu.X,n), nrow=n, byrow = TRUE)
b <- c(2, 3, -1)
sigma.eps <- 0.25
y <- cbind(rep(1, n), X.complete) %*% b +
  rnorm(n, 0, sigma.eps)

# Add missing values
p.miss <- 0.10
patterns <- runif(n*p)< p.miss #missing completely at random
X.obs <- X.complete
X.obs[patterns] <- NA
# Estimate regression using EM
df.obs = data.frame(y ,X.obs)
miss.list = miss.lm(y~, data=df.obs)

# Generate new dataset for prediction
nt <- 20
Xt <- matrix(rnorm(nt*p), nrow=nt) %*% chol(Sigma.X) +
  matrix(rep(mu.X,nt), nrow=nt, byrow = TRUE)
# Generate missingness in new dataset
patterns <- runif(nt*p)< p.miss
Xt.obs <- Xt
Xt.obs[patterns] <- NA

# Prediction with missing values
miss.pred = predict(miss.list, data.frame(Xt.obs))
print(miss.pred)

```

`print.miss.glm` *Print miss.glm*

Description

Print results for class `miss.glm`.

Usage

```

## S3 method for class 'miss.glm'
print(x, digits = max(3L, getOption("digits") - 3L), ...)

```

Arguments

- x an object of class "miss.glm", usually, a result of a call to [miss.glm](#).
- digits minimal number of significant digits.
- ... further arguments passed to or from other methods.

Value

No return value, called for coefficient and standard error estimates print.

Examples

```
## For examples see example(miss.glm)
```

print.miss.lm

Print miss.lm

Description

Print results for class miss.lm.

Usage

```
## S3 method for class 'miss.lm'
print(x, digits = max(3L, getOption("digits")) - 3L), ...)
```

Arguments

- x an object of class "miss.lm", usually, a result of a call to [miss.lm](#).
- digits minimal number of significant digits.
- ... further arguments passed to or from other methods.

Value

No return value, called for coefficient and standard error estimates print.

Examples

```
## For examples see example(miss.lm)
```

```
print.summary.miss.glm
    Print Summary of miss.glm
```

Description

Print results for class `summary.miss.glm`.

Usage

```
## S3 method for class 'summary.miss.glm'
print(x, digits = max(3L, getOption("digits") - 3L), ...)
```

Arguments

<code>x</code>	an object of class "summary.miss.glm", usually, a result of a call to <code>summary.miss.glm</code> .
<code>digits</code>	minimal number of significant digits.
<code>...</code>	further arguments passed to or from other methods.

Value

No return value, called for summary print.

Examples

```
## For examples see example(miss.glm)
```

```
print.summary.miss.lm  Print Summary of miss.lm
```

Description

Print results for class `summary.miss.lm`.

Usage

```
## S3 method for class 'summary.miss.lm'
print(x, digits = max(3L, getOption("digits") - 3L), ...)
```

Arguments

<code>x</code>	an object of class "summary.miss.lm", usually, a result of a call to <code>summary.miss.lm</code> .
<code>digits</code>	minimal number of significant digits.
<code>...</code>	further arguments passed to or from other methods.

Value

No return value, called for summary print.

Examples

```
## For examples see example(miss.lm)
```

summary.miss.glm *Summarizing Fits for miss.glm*

Description

Summary for class `miss.glm`.

Usage

```
## S3 method for class 'miss.glm'
summary(object, ...)
```

Arguments

<code>object</code>	an object of class "miss.glm", usually, a result of a call to <code>miss.glm</code> .
<code>...</code>	Further arguments passed to or from other methods.

Value

An object of class "summary.miss.glm", a list with following components:

<code>coefficients</code>	The matrix of coefficients and standard errors
<code>loglikelihood</code>	Observed log-likelihood.
<code>call</code>	the component from <code>object</code> .
<code>formula</code>	the component from <code>object</code> .

Examples

```
## For examples see example(miss.glm)
```

summary.miss.lm *Summarizing Fits for miss.lm*

Description

Summary for class `miss.lm`.

Usage

```
## S3 method for class 'miss.lm'  
summary(object, ...)
```

Arguments

<code>object</code>	an object of class "miss.lm", usually, a result of a call to <code>miss.lm</code> .
<code>...</code>	Further arguments passed to or from other methods.

Value

An object of class "summary.miss.lm", a list with following components:

<code>coefficients</code>	The matrix of coefficients and standard errors.
<code>loglikelihood</code>	Observed log-likelihood.
<code>call</code>	the component from <code>object</code> .
<code>formula</code>	the component from <code>object</code> .

Examples

```
## For examples see example(miss.lm)
```

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