

# Package ‘miselect’

July 22, 2025

**Title** Variable Selection for Multiply Imputed Data

**Version** 0.9.2

**Description** Penalized regression methods, such as lasso and elastic net, are used in many biomedical applications when simultaneous regression coefficient estimation and variable selection is desired. However, missing data complicates the implementation of these methods, particularly when missingness is handled using multiple imputation. Applying a variable selection algorithm on each imputed dataset will likely lead to different sets of selected predictors, making it difficult to ascertain a final active set without resorting to ad hoc combination rules. 'miselect' presents Stacked Adaptive Elastic Net (saenet) and Grouped Adaptive LASSO (galasso) for continuous and binary outcomes, developed by Du et al (2022) <doi:10.1080/10618600.2022.2035739>. They, by construction, force selection of the same variables across multiply imputed data. 'miselect' also provides cross validated variants of these methods.

**License** GPL-3

**Encoding** UTF-8

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**Suggests** mice, knitr, rmarkdown, testthat

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coef.cv.galasso	<i>Extract Coefficients From a "cv.galasso" Object</i>
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Description

Extract Coefficients From a "cv.galasso" Object

Usage

```
## S3 method for class 'cv.galasso'  
coef(object, lambda = object$lambda.min, ...)
```

Arguments

object	A "cv.galasso" fit
lambda	Chosen value of lambda. Must be between "min(lambda)" and "max(lambda)". Default is "lambda.min"
...	Additional unused arguments

Value

A list of numeric vectors containing the coefficients from running galasso on lambda for each imputation.

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coef.cv.saenet	<i>Extract Coefficients From a "cv.saenet" Object</i>
----------------	---

---

**Description**

Extract Coefficients From a "cv.saenet" Object

**Usage**

```
## S3 method for class 'cv.saenet'
coef(object, lambda = object$lambda.min, alpha = object$alpha.min, ...)
```

**Arguments**

object	A "cv.saenet" fit
lambda	Chosen value of lambda. Must be between "min(lambda)" and "max(lambda)". Default is "lambda.min"
alpha	Chosen value of alpha. Must be between "min(alpha)" and "max(alpha)". Default is "alpha.min"
...	Additional unused arguments

**Value**

A numeric vector containing the coefficients from running saenet on lambda and alpha.

---

coef.galasso	<i>Extract Coefficients From a "galasso" Object</i>
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---

**Description**

Extract Coefficients From a "galasso" Object

**Usage**

```
## S3 method for class 'galasso'
coef(object, lambda, ...)
```

**Arguments**

object	A "galasso" fit
lambda	Chosen value of lambda. Must be between "min(lambda)" and "max(lambda)". Default is "lambda.min"
...	Additional unused arguments

**Value**

A list of length D containing the coefficient estimates from running galasso on lambda.

---

coef.saenet	<i>Extract Coefficients From a "saenet" Object</i>
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### Description

coef.galasso averages the estimates across imputations to return a single vector instead of a matrix.

### Usage

```
## S3 method for class 'saenet'
coef(object, lambda, alpha, ...)
```

### Arguments

object	A "cv.saenet" fit
lambda	Chosen value of lambda. Must be between "min(lambda)" and "max(lambda)". Default is "lambda.min"
alpha	Chosen value of alpha. Must be between "min(alpha)" and "max(alpha)". Default is "alpha.min"
...	Additional unused arguments

### Value

A numeric vector containing the coefficients from running saenet on lambda and alpha.

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cv.galasso	<i>Cross Validated Multiple Imputation Grouped Adaptive LASSO</i>
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### Description

Does k-fold cross-validation for galasso, and returns an optimal value for lambda.

### Usage

```
cv.galasso(
  x,
  y,
  pf,
  adWeight,
  family = c("gaussian", "binomial"),
  nlambda = 100,
  lambda.min.ratio = ifelse(isTRUE(all.equal(adWeight, rep(1, p))), 0.001, 1e-06),
  lambda = NULL,
  nfolds = 5,
```

```

    foldid = NULL,
    maxit = 1000,
    eps = 1e-05
)

```

### Arguments

<code>x</code>	A length <code>m</code> list of <code>n * p</code> numeric matrices. No matrix should contain an intercept, or any missing values
<code>y</code>	A length <code>m</code> list of length <code>n</code> numeric response vectors. No vector should contain missing values
<code>pf</code>	Penalty factor. Can be used to differentially penalize certain variables
<code>adWeight</code>	Numeric vector of length <code>p</code> representing the adaptive weights for the L1 penalty
<code>family</code>	The type of response. "gaussian" implies a continuous response and "binomial" implies a binary response. Default is "gaussian".
<code>nlambda</code>	Length of automatically generated "lambda" sequence. If "lambda" is non NULL, "nlambda" is ignored. Default is 100
<code>lambda.min.ratio</code>	Ratio that determines the minimum value of "lambda" when automatically generating a "lambda" sequence. If "lambda" is not NULL, "lambda.min.ratio" is ignored. Default is 1e-4
<code>lambda</code>	Optional numeric vector of lambdas to fit. If NULL, galasso will automatically generate a lambda sequence based off of <code>nlambda</code> and <code>lambda.min.ratio</code> . Default is NULL
<code>nfolds</code>	Number of foldid to use for cross validation. Default is 5, minimum is 3
<code>foldid</code>	an optional length <code>n</code> vector of values between 1 and <code>cv.galasso</code> will automatically generate folds
<code>maxit</code>	Maximum number of iterations to run. Default is 10000
<code>eps</code>	Tolerance for convergence. Default is 1e-5

### Details

`cv.galasso` works by adding a group penalty to the aggregated objective function to ensure selection consistency across imputations. Simulations suggest that the "stacked" objective function approaches (i.e., `saenet`) tend to be more computationally efficient and have better estimation and selection properties.

### Value

An object of type "cv.galasso" with 7 elements:

**call** The call that generated the output.

**lambda** The sequence of lambdas fit.

**cvm** Average cross validation error for each "lambda". For family = "gaussian", "cvm" corresponds to mean squared error, and for binomial "cvm" corresponds to deviance.

**cvse** Standard error of "cvm".

**galasso.fit** A "galasso" object fit to the full data.

**lambda.min** The lambda value for the model with the minimum cross validation error.

**lambda.1se** The lambda value for the sparsest model within one standard error of the minimum cross validation error.

**df** The number of nonzero coefficients for each value of lambda.

## References

Du, J., Boss, J., Han, P., Beesley, L. J., Kleinsasser, M., Goutman, S. A., ... & Mukherjee, B. (2022). Variable selection with multiply-imputed datasets: choosing between stacked and grouped methods. *Journal of Computational and Graphical Statistics*, 31(4), 1063-1075. <doi:10.1080/10618600.2022.2035739>

## Examples

```
library(miselect)
library(mice)

set.seed(48109)

# Using the mice defaults for sake of example only.
mids <- mice(miselect.df, m = 5, printFlag = FALSE)
dfs <- lapply(1:5, function(i) complete(mids, action = i))

# Generate list of imputed design matrices and imputed responses
x <- list()
y <- list()
for (i in 1:5) {
  x[[i]] <- as.matrix(dfs[[i]][, paste0("X", 1:20)])
  y[[i]] <- dfs[[i]]$Y
}

pf <- rep(1, 20)
adWeight <- rep(1, 20)

fit <- cv.galasso(x, y, pf, adWeight)

# By default 'coef' returns the betas for lambda.min.
coef(fit)
```

## Description

Does k-fold cross-validation for saenet, and returns optimal values for lambda and alpha.

**Usage**

```

cv.saenet(
  x,
  y,
  pf,
  adWeight,
  weights,
  family = c("gaussian", "binomial"),
  alpha = 1,
  nlambdas = 100,
  lambda.min.ratio = ifelse(isTRUE(all.equal(adWeight, rep(1, p))), 0.001, 1e-06),
  lambda = NULL,
  nfolds = 5,
  foldid = NULL,
  maxit = 1000,
  eps = 1e-05
)

```

**Arguments**

<code>x</code>	A length <code>m</code> list of <code>n * p</code> numeric matrices. No matrix should contain an intercept, or any missing values
<code>y</code>	A length <code>m</code> list of length <code>n</code> numeric response vectors. No vector should contain missing values
<code>pf</code>	Penalty factor of length <code>p</code> . Can be used to differentially penalize certain variables. 0 indicates to not penalize the covariate
<code>adWeight</code>	Numeric vector of length <code>p</code> representing the adaptive weights for the L1 penalty
<code>weights</code>	Numeric vector of length <code>n</code> containing the proportion observed (non-missing) for each row in the un-imputed data.
<code>family</code>	The type of response. "gaussian" implies a continuous response and "binomial" implies a binary response. Default is "gaussian".
<code>alpha</code>	Elastic net parameter. Can be a vector to cross validate over. Default is 1
<code>nlambdas</code>	Length of automatically generated "lambda" sequence. If "lambda" is non NULL, "nlambdas" is ignored. Default is 100
<code>lambda.min.ratio</code>	Ratio that determines the minimum value of "lambda" when automatically generating a "lambda" sequence. If "lambda" is not NULL, "lambda.min.ratio" is ignored. Default is 1e-3
<code>lambda</code>	Optional numeric vector of lambdas to fit. If NULL, galasso will automatically generate a lambda sequence based off of <code>nlambdas</code> and <code>lambda.min.ratio</code> . Default is NULL
<code>nfolds</code>	Number of foldid to use for cross validation. Default is 5, minimum is 3
<code>foldid</code>	an optional length <code>n</code> vector of values between 1 and <code>cv.galasso</code> will automatically generate folds
<code>maxit</code>	Maximum number of iterations to run. Default is 1000
<code>eps</code>	Tolerance for convergence. Default is 1e-5

## Details

cv.saenet works by stacking the multiply imputed data into a single matrix and running a weighted adaptive elastic net on it. Simulations suggest that the "stacked" objective function approaches tend to be more computationally efficient and have better estimation and selection properties.

Due to stacking, the automatically generated lambda sequence cv.saenet generates may end up underestimating lambda.max, and thus the degrees of freedom may be nonzero at the first lambda value.

## Value

An object of type "cv.saenet" with 9 elements:

**call** The call that generated the output.

**lambda** Sequence of lambdas fit.

**cvm** Average cross validation error for each lambda and alpha. For family = "gaussian", "cvm" corresponds to mean squared error, and for binomial "cvm" corresponds to deviance.

**cvse** Standard error of "cvm".

**saenet.fit** A "saenet" object fit to the full data.

**lambda.min** The lambda value for the model with the minimum cross validation error.

**lambda.1se** The lambda value for the sparsest model within one standard error of the minimum cross validation error.

**alpha.min** The alpha value for the model with the minimum cross validation error.

**alpha.1se** The alpha value for the sparsest model within one standard error of the minimum cross validation error.

**df** The number of nonzero coefficients for each value of lambda and alpha.

## References

Du, J., Boss, J., Han, P., Beesley, L. J., Kleinsasser, M., Goutman, S. A., ... & Mukherjee, B. (2022). Variable selection with multiply-imputed datasets: choosing between stacked and grouped methods. *Journal of Computational and Graphical Statistics*, 31(4), 1063-1075. <doi:10.1080/10618600.2022.2035739>

## Examples

```
library(miselect)
library(mice)

set.seed(48109)

# Using the mice defaults for sake of example only.
mids <- mice(miselect.df, m = 5, printFlag = FALSE)
dfs <- lapply(1:5, function(i) complete(mids, action = i))

# Generate list of imputed design matrices and imputed responses
x <- list()
y <- list()
for (i in 1:5) {
```



```

      x[[i]] <- as.matrix(dfs[[i]][, paste0("X", 1:20)])
      y[[i]] <- dfs[[i]]$Y
    }

    # Calculate observational weights
    weights <- 1 - rowMeans(is.na(miselect.df))
    pf      <- rep(1, 20)
    adWeight <- rep(1, 20)

    # Since 'Y' is a binary variable, we use 'family = "binomial"'
    fit <- cv.saenet(x, y, pf, adWeight, weights, family = "binomial")

    # By default 'coef' returns the betas for (lambda.min , alpha.min)
    coef(fit)

    # You can also cross validate over alpha

    fit <- cv.saenet(x, y, pf, adWeight, weights, family = "binomial",
                     alpha = c(.5, 1))
    # Get selected variables from the 1 standard error rule
    coef(fit, lambda = fit$lambda.1se, alpha = fit$alpha.1se)

```

---

galasso

---

*Multiple Imputation Grouped Adaptive LASSO*


---

## Description

galasso fits an adaptive LASSO for multiply imputed data. "galasso" supports both continuous and binary responses.

## Usage

```

galasso(
  x,
  y,
  pf,
  adWeight,
  family = c("gaussian", "binomial"),
  nlambdas = 100,
  lambda.min.ratio = ifelse(isTRUE(all.equal(adWeight, rep(1, p))), 0.001, 1e-06),
  lambda = NULL,
  maxit = 10000,
  eps = 1e-05
)

```

**Arguments**

<code>x</code>	A length <code>m</code> list of <code>n * p</code> numeric matrices. No matrix should contain an intercept, or any missing values
<code>y</code>	A length <code>m</code> list of length <code>n</code> numeric response vectors. No vector should contain missing values
<code>pf</code>	Penalty factor. Can be used to differentially penalize certain variables
<code>adWeight</code>	Numeric vector of length <code>p</code> representing the adaptive weights for the L1 penalty
<code>family</code>	The type of response. "gaussian" implies a continuous response and "binomial" implies a binary response. Default is "gaussian".
<code>nlambda</code>	Length of automatically generated "lambda" sequence. If "lambda" is non NULL, "nlambda" is ignored. Default is 100
<code>lambda.min.ratio</code>	Ratio that determines the minimum value of "lambda" when automatically generating a "lambda" sequence. If "lambda" is not NULL, "lambda.min.ratio" is ignored. Default is 1e-4
<code>lambda</code>	Optional numeric vector of lambdas to fit. If NULL, galasso will automatically generate a lambda sequence based off of <code>nlambda</code> and <code>lambda.min.ratio</code> . Default is NULL
<code>maxit</code>	Maximum number of iterations to run. Default is 10000
<code>eps</code>	Tolerance for convergence. Default is 1e-5

**Details**

galasso works by adding a group penalty to the aggregated objective function to ensure selection consistency across imputations. The objective function is:

$$\begin{aligned} & \operatorname{argmin}_{\beta_{jk}} -L(\beta_{jk}|X_{ijk}, Y_{ik}) \\ & + \lambda * \sum_{j=1}^p \hat{a}_j * pf_j * \sqrt{\sum_{k=1}^m \beta_{jk}^2} \end{aligned}$$

Where  $L$  is the log likelihood,  $a$  is the adaptive weights, and  $pf$  is the penalty factor. Simulations suggest that the "stacked" objective function approach (i.e., saenet) tends to be more computationally efficient and have better estimation and selection properties. However, the advantage of galasso is that it allows one to look at the differences between coefficient estimates across imputations.

**Value**

An object with type `galasso` and subtype `galasso.gaussian` or `galasso.binomial`, depending on which family was used. Both subtypes have 4 elements:

**lambda** Sequence of lambda fit.

**coef** a list of length `D` containing the coefficient estimates from running galasso at each value of lambda. Each element in the list is a `nlambda x (p+1)` matrix.

**df** Number of nonzero betas at each value of lambda.

## References

Du, J., Boss, J., Han, P., Beesley, L. J., Kleinsasser, M., Goutman, S. A., ... & Mukherjee, B. (2022). Variable selection with multiply-imputed datasets: choosing between stacked and grouped methods. *Journal of Computational and Graphical Statistics*, 31(4), 1063-1075. <doi:10.1080/10618600.2022.2035739>

## Examples

```
library(miselect)
library(mice)

mids <- mice(miselect.df, m = 5, printFlag = FALSE)
dfs <- lapply(1:5, function(i) complete(mids, action = i))

# Generate list of imputed design matrices and imputed responses
x <- list()
y <- list()
for (i in 1:5) {
  x[[i]] <- as.matrix(dfs[[i]][, paste0("X", 1:20)])
  y[[i]] <- dfs[[i]]$Y
}

pf <- rep(1, 20)
adWeight <- rep(1, 20)

fit <- galasso(x, y, pf, adWeight)
```

---

miselect.df	<i>Synthetic Example Data For "miselect"</i>
-------------	--

---

## Description

This synthetic data is taken from the first simulation case from the miselect paper

## Usage

```
miselect.df
```

## Format

A data.frame with 500 observations on 21 variables:

**Y** Binary response.

**X1-X20** Covariates with missing data.

---

print.cv.galasso	<i>Print cv.galasso Objects</i>
------------------	---------------------------------

---

**Description**

print.cv.galasso print the fit and returns it invisibly.

**Usage**

```
## S3 method for class 'cv.galasso'  
print(x, ...)
```

**Arguments**

x	An object of type "cv.galasso" to print
...	Further arguments passed to or from other methods

---

print.cv.saenet	<i>Print cv.saenet Objects</i>
-----------------	--------------------------------

---

**Description**

print.cv.saenet print the fit and returns it invisibly.

**Usage**

```
## S3 method for class 'cv.saenet'  
print(x, ...)
```

**Arguments**

x	An object of type "cv.saenet" to print
...	Further arguments passed to or from other methods

saenet

*Multiple Imputation Stacked Adaptive Elastic Net***Description**

Fits an adaptive elastic net for multiply imputed data. The data is stacked and is penalized that each imputation selects the same betas at each value of lambda. "saenet" supports both continuous and binary responses.

**Usage**

```
saenet(
  x,
  y,
  pf,
  adWeight,
  weights,
  family = c("gaussian", "binomial"),
  alpha = 1,
  nlambdas = 100,
  lambda.min.ratio = ifelse(isTRUE(all.equal(adWeight, rep(1, p))), 0.001, 1e-06),
  lambda = NULL,
  maxit = 1000,
  eps = 1e-05
)
```

**Arguments**

x	A length m list of n * p numeric matrices. No matrix should contain an intercept, or any missing values
y	A length m list of length n numeric response vectors. No vector should contain missing values
pf	Penalty factor. Can be used to differentially penalize certain variables
adWeight	Numeric vector of length p representing the adaptive weights for the L1 penalty
weights	Numeric vector of length n containing the proportion observed (non-missing) for each row in the un-imputed data.
family	The type of response. "gaussian" implies a continuous response and "binomial" implies a binary response. Default is "gaussian".
alpha	Elastic net parameter. Can be a vector to cross validate over. Default is 1
nlambdas	Length of automatically generated "lambda" sequence. If "lambda" is non NULL, "nlambdas" is ignored. Default is 100
lambda.min.ratio	Ratio that determines the minimum value of "lambda" when automatically generating a "lambda" sequence. If "lambda" is not NULL, "lambda.min.ratio" is ignored. Default is 1e-3

<code>lambda</code>	Optional numeric vector of lambdas to fit. If NULL, galasso will automatically generate a lambda sequence based off of <code>nlambda</code> and <code>lambda.min.ratio</code> . Default is NULL
<code>maxit</code>	Maximum number of iterations to run. Default is 1000
<code>eps</code>	Tolerance for convergence. Default is 1e-5

## Details

saenet works by stacking the multiply imputed data into a single matrix and running a weighted adaptive elastic net on it. The objective function is:

$$\begin{aligned} \operatorname{argmin}_{\beta_j} & -\frac{1}{n} \sum_{k=1}^m \sum_{i=1}^n o_i * L(\beta_j | Y_{ik}, X_{ijk}) \\ & + \lambda \left( \alpha \sum_{j=1}^p \hat{a}_j * pf_j |\beta_j| \right. \\ & \left. + (1 - \alpha) \sum_{j=1}^p pf_j * \beta_j^2 \right) \end{aligned}$$

Where L is the log likelihood,  $o = w / m$ ,  $a$  is the adaptive weights, and  $pf$  is the penalty factor. Simulations suggest that the "stacked" objective function approach (i.e., saenet) tends to be more computationally efficient and have better estimation and selection properties. However, the advantage of galasso is that it allows one to look at the differences between coefficient estimates across imputations.

## Value

An object with type `saenet` and subtype `saenet.gaussian` or `saenet.binomial`, depending on which family was used. Both subtypes have 4 elements:

**lambda** Sequence of lambda fit.

**coef**  $n\lambda \times p + 1$  tensor representing the estimated betas at each value of lambda and alpha.

**df** Number of nonzero betas at each value of lambda and alpha.

## References

Du, J., Boss, J., Han, P., Beesley, L. J., Kleinsasser, M., Goutman, S. A., ... & Mukherjee, B. (2022). Variable selection with multiply-imputed datasets: choosing between stacked and grouped methods. *Journal of Computational and Graphical Statistics*, 31(4), 1063-1075. <doi:10.1080/10618600.2022.2035739>

## Examples

```
library(miselect)
library(mice)

mids <- mice(miselect.df, m = 5, printFlag = FALSE)
dfs <- lapply(1:5, function(i) complete(mids, action = i))
```

```
# Generate list of imputed design matrices and imputed responses
x <- list()
y <- list()
for (i in 1:5) {
  x[[i]] <- as.matrix(dfs[[i]][, paste0("X", 1:20)])
  y[[i]] <- dfs[[i]]$Y
}

# Calculate observational weights
weights <- 1 - rowMeans(is.na(miselect.df))
pf      <- rep(1, 20)
adWeight <- rep(1, 20)

# Since 'Y' is a binary variable, we use 'family = "binomial"'
fit <- saenet(x, y, pf, adWeight, weights, family = "binomial")
```

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