

# Package: ADSIHT (via r-universe)

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**Type** Package

**Title** Adaptive Double Sparse Iterative Hard Thresholding

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**Description** Solving the high-dimensional double sparse linear regression via iterative hard thresholding algorithm. For more details, please see Zhang et al. (2024, <[DOI:10.48550/arXiv.2305.04182](https://doi.org/10.48550/arXiv.2305.04182)>).

**License** GPL (>= 3)

**Depends** R (>= 4.1.0)

**Imports** Matrix, mvnfast, Rcpp

**Suggests** knitr, rmarkdown, testthat

**LinkingTo** Rcpp, RcppEigen

**Encoding** UTF-8

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**RoxygenNote** 7.3.2

**NeedsCompilation** yes

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**Repository** CRAN

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## Contents

ADSIHT . . . . .	2
ADSIHT.ML . . . . .	4
gen.data . . . . .	6
<b>Index</b>	<b>8</b>

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ADSIHT	<i>Adaptive Double Sparse Iterative Hard Thresholding Algorithm (ADSIHT)</i>
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### Description

An implementation of the sparse group selection in linear regression model via ADSIHT.

### Usage

```
ADSIHT(
  x,
  y,
  group,
  s0,
  kappa = 0.9,
  ic.type = c("dsic", "loss"),
  ic.scale = 3,
  ic.coef = 3,
  L = 5,
  weight = rep(1, nrow(x)),
  coef1 = 1,
  coef2 = 1,
  eta = 0.8,
  max_iter = 20,
  method = "ols"
)
```

### Arguments

<code>x</code>	Input matrix, of dimension $n \times p$ ; each row is an observation vector and each column is a predictor.
<code>y</code>	The response variable of $n$ observations.
<code>group</code>	A vector indicating which group each variable belongs to. For variables in the same group, they should be located in adjacent columns of <code>x</code> and their corresponding index in <code>group</code> should be the same. Denote the first group as 1, the second 2, etc.
<code>s0</code>	A vector that controls the degrees with group. Default is $d^{(l-1)/(L-1)}$ : $1 \leq l \leq L$ , where $d$ is the maximum group size.
<code>kappa</code>	A parameter that controls the rapid of the decrease of threshold. Default is 0.9.
<code>ic.type</code>	The type of criterion for choosing the support size. Available options are "dsic", "loss". Default is "dsic".
<code>ic.scale</code>	A non-negative value used for multiplying the penalty term in information criterion. Default: <code>ic.scale</code> = 3.

ic.coef	A non-negative value used for multiplying the penalty term for choosing the optimal stopping time. Default: ic.coef = 3.
L	The length of the sequence of $s_0$ . Default: L = 5.
weight	The weight of the samples, with the default value set to 1 for each sample.
coef1	A positive value to control the sub-optimal stopping time.
coef2	A positive value to control the overall stopping time. A small value leads to larger search range.
eta	A parameter controls the step size in the gradient descent step. Default: eta = 0.8.
max_iter	A parameter that controls the maximum number of line search, ignored if OLS is employed.
method	Whether ols (default) or linesearch method should be employed.

### Value

A list object comprising:

beta	A $p$ -by-length( $s_0$ ) matrix of coefficients, stored in column format.
intercept	A length( $s_0$ ) vector of intercepts
.	
lambda	A length( $s_0$ ) vector of threshold values
A_out	The selected variables given threshold value in lambda.
ic	The values of the specified criterion for each fitted model given threshold lambda.

### Author(s)

Yanhang Zhang, Zhifan Li, Shixiang Liu, Jianxin Yin.

### Examples

```
n <- 200
m <- 100
d <- 10
s <- 5
s0 <- 5
data <- gen.data(n, m, d, s, s0)
fit <- ADSIHT(data$x, data$y, data$group)
fit$A_out[which.min(fit$ic)]
```

**Description**

An implementation of the sparse group selection in linear regression model via ADSIHT.

**Usage**

```
ADSIHT.ML(
  x_list,
  y_list,
  group_list,
  s0,
  kappa = 0.9,
  ic.type = c("dsic", "loss"),
  ic.scale = 3,
  ic.coef = 3,
  L = 5,
  weight,
  coef1 = 1,
  coef2 = 1,
  eta = 0.8,
  max_iter = 20,
  method = "ols"
)
```

**Arguments**

<code>x_list</code>	The list of input matrix.
<code>y_list</code>	The list of response variable.
<code>group_list</code>	A vector indicating which group each variable belongs to. For variables in the same group, they should be located in adjacent columns of <code>x</code> and their corresponding index in <code>group</code> should be the same. Denote the first group as 1, the second 2, etc.
<code>s0</code>	A vector that controls the degrees with group. Default is $d^{(l-1)/(L-1)}$ : $1 \leq l \leq L$ , where <code>d</code> is the maximum group size.
<code>kappa</code>	A parameter that controls the rapid of the decrease of threshold. Default is 0.9.
<code>ic.type</code>	The type of criterion for choosing the support size. Available options are "dsic", "loss". Default is "dsic".
<code>ic.scale</code>	A non-negative value used for multiplying the penalty term in information criterion. Default: <code>ic.scale</code> = 3.
<code>ic.coef</code>	A non-negative value used for multiplying the penalty term for choosing the optimal stopping time. Default: <code>ic.coef</code> = 3.
<code>L</code>	The length of the sequence of <code>s0</code> . Default: <code>L</code> = 5.

weight	The weight of the samples, with the default value set to 1 for each sample.
coef1	A positive value to control the sub-optimal stopping time.
coef2	A positive value to control the overall stopping time. A small value leads to larger search range.
eta	A parameter controls the step size in the gradient descent step. Default: eta = 0.8.
max_iter	A parameter that controls the maximum number of line search, ignored if OLS is employed.
method	Whether ols (default) or linesearch method should be employed.

### Value

A list object comprising:

beta	A $p$ -by- $\text{length}(s_0)$ matrix of coefficients, stored in column format.
intercept	A $\text{length}(s_0)$ vector of intercepts
.	.
lambda	A $\text{length}(s_0)$ vector of threshold values
A_out	The selected variables given threshold value in lambda.
ic	The values of the specified criterion for each fitted model given threshold lambda.

### Author(s)

Yanhang Zhang, Zhifan Li, Shixiang Liu, Jianxin Yin.

### Examples

```

set.seed(1)
n <- 200
p <- 100
K <- 4
s <- 5
s0 <- 2
x_list <- lapply(1:K, function(x) matrix(rnorm(n*p, 0, 1), nrow = n))
vec <- rep(0, K * p)
non_sparse_groups <- sample(1:p, size = s, replace = FALSE)
for (group in non_sparse_groups) {
  group_indices <- seq(group, K * p, by = p)
  non_zero_indices <- sample(group_indices, size = s0, replace = FALSE)
  vec[non_zero_indices] <- rep(2, s0)
}
y_list <- lapply(1:K, function(i) return(
  y = x_list[[i]] %*% vec[((i-1)*p+1):(i*p)]+rnorm(n, 0, 0.5))
)
fit <- ADSIHT.ML(x_list, y_list)
fit$A_out[, which.min(fit$ic)]

```

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gen.data                      *Generate simulated data*

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### Description

Generate simulated data for sparse group linear model.

### Usage

```
gen.data(
  n,
  m,
  d,
  s,
  s0,
  cor.type = 1,
  beta.type = 1,
  rho = 0.5,
  sigma1 = 1,
  sigma2 = 1,
  seed = 1
)
```

### Arguments

n	The number of observations.
m	The number of groups of interest.
d	The group size of each group. Only even group structure is allowed here.
s	The number of important groups in the underlying regression model.
s0	The number of important variables in each important group.
cor.type	The structure of correlation. cor.type = 1 denotes the independence structure, where the covariance matrix has $(i, j)$ entry equals $I(i \neq j)$ . cor.type = 2 denotes the exponential structure, where the covariance matrix has $(i, j)$ entry equals $\rho^{ i-j }$ . cor.type = 3 denotes the constant structure, where the non-diagonal entries of covariance matrix are $\rho$ and diagonal entries are 1.
beta.type	The structure of coefficients. beta.type = 1 denotes the homogenous setup, where each entry has the same magnitude. beta.type = 2 denotes the heterogeneous structure, where the coefficients are drawn from a normal distribution.
rho	A parameter used to characterize the pairwise correlation in predictors. Default is 0.5.
sigma1	The value controlling the strength of the gaussian noise. A large value implies strong noise. Default sigma1 = 1.
sigma2	The value controlling the strength of the coefficients. A large value implies large coefficients. Default sigma2 = 1.
seed	random seed. Default: seed = 1.

**Value**

A list object comprising:

<code>x</code>	Design matrix of predictors.
<code>y</code>	Response variable.
<code>beta</code>	The coefficients used in the underlying regression model.
<code>group</code>	The group index of each variable.
<code>true.group</code>	The important groups in the sparse group linear model.
<code>true.variable</code>	The important variables in the sparse group linear model.

**Author(s)**

Yanhang Zhang, Zhifan Li, Jianxin Yin.

**Examples**

```
# Generate simulated data
n <- 200
m <- 100
d <- 10
s <- 5
s0 <- 5
data <- gen.data(n, m, d, s, s0)
str(data)
```

# Index

ADSIHT, [2](#)  
ADSIHT.ML, [4](#)  
gen.data, [6](#)