Package ‘LassoGEE’

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**Type**  Package

**Title**  High-Dimensional Lasso Generalized Estimating Equations

**Version**  1.0

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**Description**  
Fits generalized estimating equations with L1 regularization to longitudinal data with high dimensional covariates. Use a efficient iterative composite gradient descent algorithm.

**License**  GPL (>= 2)

**URL**  <https://github.com/liygCR/LassoGEE>

**Depends**  R (>= 3.6.0)

**Encoding**  UTF-8

**LazyData**  true

**Imports**  Rcpp (>= 1.0.4), PGEE, MASS, mvtnorm, caret, SimCorMultRes

**LinkingTo**  Rcpp, RcppArmadillo

**RoxygenNote**  7.1.1

**NeedsCompilation**  yes

**Repository**  CRAN

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Cross-validation for LassoGEE.

Description

Does k-fold cross-validation for LassoGEE to select tuning parameter value for longitudinal data with working independence structure.

Usage

```r
cv.LassoGEE(
  X,
  y,
  id,
  family,
  method = c("CGD", "RWL"),
  scale.fix,
  scale.value,
  fold,
  lambda.vec,
  maxiter,
  tol
)
```

Arguments

- `X`: A design matrix of dimension \((nm) \times p\).
- `y`: A response vector of length \(m \times n\).
- `id`: A vector for identifying subjects/clusters.
- `family`: A family object: a list of functions and expressions for defining link and variance functions. Families supported here is same as in `PGEE` which are binomial, gaussian, gamma and poisson.
- `method`: The algorithms that are available. "CGD" represents the I-CGD algorithm, and "RWL" represents re-weighted least square algorithm.
- `scale.fix`: A logical variable; if true, the scale parameter is fixed at the value of `scale.value`. The default value is TRUE.
- `scale.value`: If `scale.fix = TRUE`, this assigns a numeric value to which the scale parameter should be fixed. The default value is 1.
- `fold`: The number of folds used in cross-validation.
- `lambda.vec`: A vector of tuning parameters that will be used in the cross-validation.
- `maxiter`: The number of iterations that is used in the estimation algorithm. The default value is 50.
- `tol`: The tolerance level that is used in the estimation algorithm. The default value is \(1e^{-3}\).
Value

An object class of cv.LassoGEE.

References


See Also

LassoGEE

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IC

Information Criterion for selecting the tuning parameter.

Description

Information Criterion for a fitted LassoGEE object with the AIC, BIC, or GCV criteria.

Usage

IC(obj, criterion = c("BIC", "AIC", "GCV", "AICc", "EBIC"))

Arguments

obj A fitted LassoGEE object.

criterion The criterion by which to select the regularization parameter. One of "AIC", "BIC", "GCV", "AICc", or "EBIC"; default is "BIC".

Value

IC The calculated model selection criteria

References

LassoGEE

Function to fit penalized GEE by I-CGD algorithm.

Description

This function fits a $L_1$ penalized GEE model to longitudinal data by I-CGD algorithm or re-weighted least square algorithm.

Usage

LassoGEE(
  X,
  y,
  id,
  family = binomial("probit"),
  lambda,
  corstr = "independence",
  method = c("CGD", "RWL"),
  beta.ini = NULL,
  R = NULL,
  scale.fix = TRUE,
  scale.value = 1,
  maxiter = 50,
  tol = 0.001,
  silent = TRUE,
  Mv = NULL,
  verbose = TRUE
)

Arguments

X
  A design matrix of dimension (nm) * p.

y
  A response vector of length m * n.

id
  A vector for identifying subjects/clusters.

family
  A family object representing one of the built-in families. Families supported here are the same as in PGEE, e.g., binomial, gaussian, gamma and poisson, and the corresponding link functions are supported, e.g., identity, and probit.

lambda
  A user supplied value for the penalization parameter.

corstr
  A character string that indicates the correlation structure among the repeated measurements of a subject. Structures supported in LassoGEE are "AR1", "exchangeable", "unstructured", and "independence". The default corstr type is "independence".

method
  The algorithms that are available. "CGD" represents the I-CGD algorithm, and "RWL" represents re-weighted least square algorithm.

beta.ini
  User specified initial values for regression parameters. The default value is NULL.
R

User specified correlation matrix. The default value is NULL.

scale.fix

A logical variable. The default value is TRUE, then the value of the scale parameter is fixed to scale.value.

scale.value

If scale.fix = TRUE, a numeric value will be assigned to the fixed scale parameter. The default value is 1.

maxiter

The maximum number of iterations used in the algorithm. The default value is 50.

tol

The tolerance level used in the algorithm. The default value is 1e-3.

silent

A logical variable; if false, the iteration counts at each iteration of CGD are printed. The default value is TRUE.

Mv

If either "stat_M_dep", or "non_stat_M_dep" is specified in corstr, then this assigns a numeric value for Mv. Otherwise, the default value is NULL.

verbose

A logical variable; Print the out loop iteration counts. The default value is TRUE.

Value

A list containing the following components:

betaest

return final estimation

beta_all_step

return estimate in each iteration

inner.count

iterative count in each stage

outer.iter

iterate number of outer loop

References


See Also

cv.LassoGEE

Examples

# required R package
library(mvtnorm)
laboratory(SimCorMultRes)
#
set.seed(123)
p <- 200
s <- ceiling(p^{1/3})
n <- ceiling(10 * s * log(p))
m <- 4
# covariance matrix of p number of continuous covariates
X.sigma <- matrix(0, p, p)
{
for (i in 1:p)
X.sigma[i,] <- 0.5^(abs(1:p)-i))
}
# generate matrix of covariates
X <- as.matrix(rmvnorm(n*m, mean = rep(0,p), X.sigma))

# true regression parameter associated with the covariate
bt <- runif(s, 0.05, 0.5) # = rep(1/s,s)
beta.true <- c(bt,rep(0,p-s))
# intercept
beta_intercepts <- 0
# unstructure
tt <- runif(m*m,-1,1)
Rtmp <- t(matrix(tt, m,m))%*%matrix(tt, m,m)+diag(1,4)
R_tr <- diag(diag(Rtmp)^{-1/2})%*%Rtmp%*%diag(diag(Rtmp)^{-1/2})
diag(R_tr) = round(diag(R_tr))

# library(SimCorMultRes)
# simulation of clustered binary responses
simulated_binary_dataset <- rbin(clsize = m, intercepts = beta_intercepts,
betas = beta.true, xformula = ~X, cor.matrix = R_tr,
link = "probit")
lambda <- 0.2* s *sqrt(log(p)/n)
data = simulated_binary_dataset$simdata
y = data$y
X = data$X
id = data$id

ptm <- proc.time()
nCGDfit = LassoGEE(X = X, y = y, id = id, family = binomial("probit"),
lambda = lambda, corstr = "unstructured")
proc.time() - ptm
betaest <- nCGDfit$betaest

print.cv.LassoGEE

print a cross-validated LassoGEE object

Description

Print a summary of the results of cross-validation for a LassoGEE model.

Usage

## S3 method for class 'cv.LassoGEE'
print(x, digits = NULL, ...)
print.LassoGEE

Arguments

x fitted 'cv.LassoGEE' object
digits significant digits in printout
... additional print arguments

Details

A summary of the cross-validated fit is produced. `print.cv.LassoGEE(object)` will print the summary for a sequence of lambda.

References


See Also

LassoGEE, and cv.LassoGEE methods.

print.LassoGEE  print a LassoGEE object

Description

Print a summary of the results of a LassoGEE model.

Usage

```r
## S3 method for class 'LassoGEE'
print(x, digits = NULL, ...)
```

Arguments

x fitted 'LassoGEE' object
digits significant digits in printout
... additional print arguments

Details

A summary of the cross-validated fit is produced. `print.cv.LassoGEE(object)` will print the summary includes Working Correlation and Returned Error Value.

References

See Also

LassoGEE, and cv.LassoGEE methods.
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