

Package ‘CVglasso’

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Type Package

Title Lasso Penalized Precision Matrix Estimation

Version 1.0

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Description Estimates a lasso penalized precision matrix via the blockwise coordinate descent (BCD). This package is a simple wrapper around the popular 'glasso' package that extends and enhances its capabilities. These enhancements include built-in cross validation and visualizations.
See Friedman et al (2008) <[doi:10.1093/biostatistics/kxm045](https://doi.org/10.1093/biostatistics/kxm045)> for details regarding the estimation method.

URL <https://github.com/MGallow/CVglasso>

BugReports <https://github.com/MGallow/CVglasso/issues>

License GPL (>= 2)

ByteCompile TRUE

Encoding UTF-8

LazyData true

RoxygenNote 6.0.1

Imports stats, parallel, foreach, ggplot2, dplyr, glasso

Depends doParallel

Suggests testthat

NeedsCompilation no

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CVglasso

*Penalized precision matrix estimation***Description**

Penalized precision matrix estimation using the graphical lasso (glasso) algorithm. Consider the case where X_1, \dots, X_n are iid $N_p(\mu, \Sigma)$ and we are tasked with estimating the precision matrix, denoted $\Omega \equiv \Sigma^{-1}$. This function solves the following optimization problem:

Objective: $\hat{\Omega}_\lambda = \arg \min_{\Omega \in S_+^p} \{Tr(S\Omega) - \log \det(\Omega) + \lambda \|\Omega\|_1\}$

where $\lambda > 0$ and we define $\|A\|_1 = \sum_{i,j} |A_{ij}|$.

Usage

```
CVglasso(X = NULL, S = NULL, nlam = 10, lam.min.ratio = 0.01,
         lam = NULL, diagonal = FALSE, path = FALSE, tol = 1e-04,
         maxit = 10000, adjmaxit = NULL, K = 5, crit.cv = c("loglik", "AIC",
         "BIC"), start = c("warm", "cold"), cores = 1, trace = c("progress",
         "print", "none"), ...)
```

Arguments

<code>X</code>	option to provide a $n \times p$ data matrix. Each row corresponds to a single observation and each column contains n observations of a single feature/variable.
<code>S</code>	option to provide a $p \times p$ sample covariance matrix (denominator n). If argument is <code>NULL</code> and <code>X</code> is provided instead then <code>S</code> will be computed automatically.
<code>nlam</code>	number of <code>lam</code> tuning parameters for penalty term generated from <code>lam.min.ratio</code> and <code>lam.max</code> (automatically generated). Defaults to 10.
<code>lam.min.ratio</code>	smallest <code>lam</code> value provided as a fraction of <code>lam.max</code> . The function will automatically generate <code>nlam</code> tuning parameters from <code>lam.min.ratio*lam.max</code> to <code>lam.max</code> in \log_{10} scale. <code>lam.max</code> is calculated to be the smallest <code>lam</code> such that all off-diagonal entries in Ω are equal to zero ($\alpha = 1$). Defaults to $1e-2$.
<code>lam</code>	option to provide positive tuning parameters for penalty term. This will cause <code>nlam</code> and <code>lam.min.ratio</code> to be disregarded. If a vector of parameters is provided, they should be in increasing order. Defaults to <code>NULL</code> .
<code>diagonal</code>	option to penalize the diagonal elements of the estimated precision matrix (Ω). Defaults to <code>FALSE</code> .
<code>path</code>	option to return the regularization path. This option should be used with extreme care if the dimension is large. If set to <code>TRUE</code> , <code>cores</code> must be set to 1 and errors and optimal tuning parameters will be based on the full sample. Defaults to <code>FALSE</code> .
<code>tol</code>	convergence tolerance. Iterations will stop when the average absolute difference in parameter estimates is less than <code>tol</code> times multiple. Defaults to $1e-4$.
<code>maxit</code>	maximum number of iterations. Defaults to $1e4$.

<code>adjmaxit</code>	adjusted maximum number of iterations. During cross validation this option allows the user to adjust the maximum number of iterations after the first <code>lam</code> tuning parameter has converged. This option is intended to be paired with <code>warm</code> starts and allows for 'one-step' estimators. Defaults to <code>NULL</code> .
<code>K</code>	specify the number of folds for cross validation.
<code>crit.cv</code>	cross validation criterion (<code>loglik</code> , <code>AIC</code> , or <code>BIC</code>). Defaults to <code>loglik</code> .
<code>start</code>	specify <code>warm</code> or <code>cold</code> start for cross validation. Default is <code>warm</code> .
<code>cores</code>	option to run CV in parallel. Defaults to <code>cores = 1</code> .
<code>trace</code>	option to display progress of CV. Choose one of <code>progress</code> to print a progress bar, <code>print</code> to print completed tuning parameters, or <code>none</code> .
<code>...</code>	additional arguments to pass to <code>glasso</code> .

Details

For details on the implementation of the 'glasso' function, see Tibshirani's website. <http://statweb.stanford.edu/~tibs/glasso/>.

Value

returns class object `CVglasso` which includes:

<code>Call</code>	function call.
<code>Iterations</code>	number of iterations
<code>Tuning</code>	optimal tuning parameters (<code>lam</code> and <code>alpha</code>).
<code>Lambdas</code>	grid of lambda values for CV.
<code>maxit</code>	maximum number of iterations for outer (blockwise) loop.
<code>Omega</code>	estimated penalized precision matrix.
<code>Sigma</code>	estimated covariance matrix from the penalized precision matrix (inverse of <code>Omega</code>).
<code>Path</code>	array containing the solution path. Solutions will be ordered by ascending lambda values.
<code>MIN.error</code>	minimum average cross validation error (<code>cv.crit</code>) for optimal parameters.
<code>AVG.error</code>	average cross validation error (<code>cv.crit</code>) across all folds.
<code>CV.error</code>	cross validation errors (<code>cv.crit</code>).

Author(s)

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References

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- Rothman, Adam. 2017. 'STAT 8931 notes on an algorithm to compute the Lasso-penalized Gaussssian likelihood precision matrix estimator.'

See Also

[plot.CVglasso](#)

Examples

```
# generate data from a sparse matrix
# first compute covariance matrix
S = matrix(0.7, nrow = 5, ncol = 5)
for (i in 1:5){
  for (j in 1:5){
    S[i, j] = S[i, j]^abs(i - j)
  }
}

# generate 100 x 5 matrix with rows drawn from iid N_p(0, S)
Z = matrix(rnorm(100*5), nrow = 100, ncol = 5)
out = eigen(S, symmetric = TRUE)
S.sqrt = out$vectors %*% diag(out$values^0.5)
S.sqrt = S.sqrt %*% t(out$vectors)
X = Z %*% S.sqrt

# lasso penalty CV
CVglasso(X)
```

plot.CVglasso	<i>Plot CVglasso object</i>
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Description

Produces a plot for the cross validation errors, if available.

Usage

```
## S3 method for class 'CVglasso'
plot(x, type = c("line", "heatmap"), footnote = TRUE,
     ...)
```

Arguments

x	class object CVglasso
type	produce either 'heatmap' or 'line' graph
footnote	option to print footnote of optimal values. Defaults to TRUE.
...	additional arguments.

Examples

```
# generate data from a sparse matrix
# first compute covariance matrix
S = matrix(0.7, nrow = 5, ncol = 5)
for (i in 1:5){
  for (j in 1:5){
    S[i, j] = S[i, j]^abs(i - j)
  }
}

# generate 100 x 5 matrix with rows drawn from iid N_p(0, S)
Z = matrix(rnorm(100*5), nrow = 100, ncol = 5)
out = eigen(S, symmetric = TRUE)
S.sqrt = out$vectors %*% diag(out$values^0.5)
S.sqrt = S.sqrt %*% t(out$vectors)
X = Z %*% S.sqrt

# produce line graph for CVglasso
plot(CVglasso(X))

# produce CV heat map for CVglasso
plot(CVglasso(X), type = 'heatmap')
```

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