

# Package ‘shrink’

June 25, 2018

**Type** Package

**Title** Shrinking characteristics of precision matrix estimators

**Version** 2.0

**Date** 2018-05-30

**Description** This package is an implementation of the methods described in “Shrinking Characteristics of Precision Matrix Estimators”. <https://arxiv.org/pdf/1704.04820.pdf>

**URL** <https://github.com/MGallow/shrink>

**BugReports** <https://github.com/MGallow/shrink/issues>

**License** GPL (>= 2)

**ByteCompile** TRUE

**NeedsCompilation** yes

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.0.1

**Imports** stats,  
parallel,  
foreach,  
ggplot2,  
dplyr

**Depends** Rcpp (>= 0.12.10),  
RcppProgress (>= 0.1),  
doParallel

**LinkingTo** Rcpp,  
RcppArmadillo,  
RcppProgress

**Suggests** testthat,  
knitr,  
rmarkdown,  
microbenchmark,  
glasso,  
pkgdown

**SystemRequirements** GNU make

**VignetteBuilder** knitr

## R topics documented:

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plot.shrink	<i>Plot shrink object</i>
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### Description

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

### Usage

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

### Arguments

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

### Examples

```
# NEED TO COMPLETE
```

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shrink	<i>Shrinking characteristics of precision matrix estimators</i>
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### Description

Shrinking characteristics of precision matrix estimators. Penalized precision matrix estimation using the ADMM 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:**  $\hat{\Omega}_\lambda = \arg \min_{\Omega \in S_+^p} \{Tr(S\Omega) - \log \det(\Omega) + \lambda \|A\Omega B - C\|_1\}$

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

### Usage

```
shrink(X = NULL, S = NULL, Y = NULL, A = diag(ncol(S)),
       B = diag(ncol(S)), C = diag(ncol(S)), nlam = 10, lam.min.ratio = 0.01,
       lam = NULL, path = FALSE, rho = 2, mu = 10, tau.inc = 2,
       tau.dec = 2, crit = c("ADMM", "loglik"), tol.abs = 1e-04,
       tol.rel = 1e-04, maxit = 10000, adjmaxit = NULL, K = 5,
       crit.cv = c("MSE", "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 NULL and <code>X</code> is provided instead then <code>S</code> will be computed automatically.
<code>Y</code>	option to provide $n \times r$ response matrix. Each row corresponds to a single response and each column contains $n$ response of a single feature/response.
<code>A</code>	option to provide user-specified matrix for penalty term. This matrix must have $p$ columns. Defaults to identity matrix.
<code>B</code>	option to provide user-specified matrix for penalty term. This matrix must have $p$ rows. Defaults to identity matrix.
<code>C</code>	option to provide user-specified matrix for penalty term. This matrix must have $nrow(A)$ rows and $ncol(B)$ columns. Defaults to identity matrix.
<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 log10 scale. <code>lam.max</code> is calculated to be the smallest <code>lam</code> such that all off-diagonal entries in $\Omega$ 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 NULL.
<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 TRUE, cores must be set to 1 and errors and optimal tuning parameters will be based on the full sample. Defaults to FALSE.
<code>rho</code>	initial step size for ADMM algorithm.
<code>mu</code>	factor for primal and residual norms in the ADMM algorithm. This will be used to adjust the step size <code>rho</code> after each iteration.
<code>tau.inc</code>	factor in which to increase step size <code>rho</code>
<code>tau.dec</code>	factor in which to decrease step size <code>rho</code>
<code>crit</code>	criterion for convergence (ADMM or loglik). If <code>crit = loglik</code> then iterations will stop when the relative change in log-likelihood is less than <code>tol.abs</code> . Default is ADMM and follows the procedure outlined in Boyd, et al.
<code>tol.abs</code>	absolute convergence tolerance. Defaults to $1e-4$ .
<code>tol.rel</code>	relative convergence tolerance. 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 (for each $\alpha$ ). This option is intended to be paired with warm starts and allows for 'one-step' estimators. Defaults to NULL.
<code>K</code>	specify the number of folds for cross validation.
<code>crit.cv</code>	cross validation criterion (MSE, loglik, AIC, or BIC). Defaults to MSE.
<code>start</code>	specify warm or cold start for cross validation. Default is warm.
<code>cores</code>	option to run CV in parallel. Defaults to <code>cores = 1</code> .

trace	option to display progress of CV. Choose one of progress to print a progress bar, print to print completed tuning parameters, or none.
alpha	elastic net mixing parameter contained in $[0, 1]$ . 0 = ridge, 1 = lasso. If a vector of parameters is provided, they should be in increasing order. Defaults to grid of values <code>seq(0, 1, 0.2)</code> .

### Details

For details on the implementation of 'shrink', see the vignette <https://mgallow.github.io/shrink/>.

### Value

returns class object `ADMMsigma` which includes:

Call	function call.
Iterations	number of iterations.
Tuning	optimal tuning parameter.
Lambdas	grid of lambda values for CV.
maxit	maximum number of iterations.
Omega	estimated penalized precision matrix.
Sigma	estimated covariance matrix from the penalized precision matrix (inverse of Omega).
Path	array containing the solution path. Solutions will be ordered in ascending alpha values for each lambda.
Z	final sparse update of estimated penalized precision matrix.
Y	final dual update.
rho	final step size.
Loglik	penalized log-likelihood for Omega
MIN.error	minimum average cross validation error (cv.crit) for optimal parameters.
AVG.error	average cross validation error (cv.crit) across all folds.
CV.error	cross validation errors (cv.crit).

### Author(s)

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### References

- Boyd, Stephen, Neal Parikh, Eric Chu, Borja Peleato, Jonathan Eckstein, and others. 2011. 'Distributed Optimization and Statistical Learning via the Alternating Direction Method of Multipliers.' *Foundations and Trends in Machine Learning* 3 (1). Now Publishers, Inc.: 1-122. [https://web.stanford.edu/~boyd/papers/pdf/admm\\_distr\\_stats.pdf](https://web.stanford.edu/~boyd/papers/pdf/admm_distr_stats.pdf)
- Hu, Yue, Chi, Eric C, and Allen, Genevera I. 2016. 'ADMM Algorithmic Regularization Paths for Sparse Statistical Machine Learning.' *Splitting Methods in Communication, Imaging, Science, and Engineering*. Springer: 433-459.
- Molstad, Aaron J., and Adam J. Rothman. (2017). 'Shrinking Characteristics of Precision Matrix Estimators.' *arXiv preprint arXiv: 1704.04820*. <https://arxiv.org/pdf/1704.04820.pdf>
- Rothman, Adam. 2017. 'STAT 8931 notes on an algorithm to compute the Lasso-penalized Gaussian likelihood precision matrix estimator.'

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### See Also

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### Examples

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# NEED TO COMPLETE
```

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