Package 'gicf'

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Description

Penalised likelihood estimation of a covariance matrix via the ridge-regularised covglasso estimator described in Cibinel et al. (2024) <doi:10.48550/arXiv.2410.02403>.

Details

This package optimises the penalised loglikelihood function of a Gaussian covariance graphical model via an iterative coordinate descent algorithm.

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References

Cibinel, L., A. Roverato, and V. Vinciotti (2024). A unified approach to penalized likelihood estimation of covariance matrices in high dimensions. arXiv, arXiv:2410.02403.

gcgmloglik	Gaussian Covariance Graphical Model Loglikelihood function

Description

Computes the penalised loglikelihood function of a Gaussian covariance graph model.

Usage

```
gcgmloglik(Sigma, S, n, lambda = 0, kappa = 0)
```

Arguments

Sigma	The covariance matrix.
S	The sample covariance matrix.
n	The size of the observed dataset.
lambda	A non-negative lasso parameter.
kappa	A non-negative ridge regularisation parameter.

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Details

When imposing sparsity on the covariance matrix of a multivariate Gaussian distribution, the resulting model can be interpreted as a covariance graphical model, i.e., the independence structure of the components of the random vector can be encoded by a graph in which the nodes are identified with the variables and a missing edge between two nodes implies that the corresponding variables are marginally independent.

In particular, this model admits both a ridge and a lasso penalty, resulting in the loglikelihood function

$$-\mathrm{log}|\Sigma|-\mathrm{trace}(\Sigma^{-1}S)-\lambda\|\Sigma-\mathrm{diag}(\Sigma)\|_1-\kappa\|\Sigma^{-1}\|_1,$$
 where $\lambda,\kappa\geq 0.$

Value

The value of the penalised loglikelihood function.

Examples

```
# An example with a banded covariance matrix
library(mvtnorm)
set.seed(1234)
p <- 10
n <- 500
# Create banded covariance matrix with three bands
band1 <- cbind(1:(p - 1), 2:p)
band2 <- cbind(1:(p - 2), 3:p)
band3 <- cbind(1:(p - 3), 4:p)
idxs <- rbind(band1, band2, band3)</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[idxs] <- 0.5
Sigma <- Sigma + t(Sigma)</pre>
diag(Sigma) <- 2
# Generate data
data <- rmvnorm(n, sigma = Sigma)</pre>
S \leftarrow cov(data) * (n - 1)/n
# Fix a value of lambda and kappa
lambda <- 0.07
kappa <- 0.5
# Gaussian loglikelihood
print("Gaussian loglikelihood:")
print(gcgmloglik(Sigma, S, n))
# Penalised Gaussian loglikelihood
print("Penalised Gaussian loglikelihood:")
print(gcgmloglik(Sigma, S, n, lambda, kappa))
```

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Penalised maximum likelihood covariance matrix estimation

Description

Estimation of a sparse covariance matrix via the ridge-regularised covglasso estimator described in Cibinel et al. (2024).

Usage

```
gicf(
  data = NULL,
  S = NULL,
  n = NULL,
  lambda = 0,
  kappa = 0,
  max.iter = 2500,
  tol = 1e-04,
  Sigma.init = NULL,
  adj = NULL
)
```

Arguments

data	A numerical matrix whose rows contain the observations of multivariate normal random vector. If NULL, the sample covariance matrix S and the dataset size n must be provided.
S	The sample covariance matrix. Must be provided if data is NULL.
n	The dataset size. Must be provided if data is NULL.
lambda	A vector of non-negative lasso parameters. For efficency purposes, should be sorted from largest to smallest.
kappa	A non-negative ridge regularisation parameter.
max.iter	The maximum number of iterations allowed for the coordinate descent algorithm.
tol	A numerical tolerance below which quantities are treated as zero.
Sigma.init	The initial guess for the coordinate descent algorithm. Defaults to the diagonal of the sample covariance matrix.
adj	An optional matrix whose pattern of zeroes is enforced onto the final output of the algorithm.

Details

This function computes the ridge-regularised covglasso estimator of the covariance matrix of a multivariate normal distribution, that is it computes the maximum of the penalised log-likelihood

$$-\log|\Sigma| - \operatorname{trace}(\Sigma^{-1}S) - \lambda \|\Sigma - \operatorname{diag}(\Sigma)\|_1 - \kappa \|\Sigma^{-1}\|_1,$$

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where $\lambda, \kappa \geq 0$. The optimum is computed via a coordinate descent algorithm, resulting in an approach which unifies and extends the methods of Chaudhuri et. al (2007), Warton (2008), Bien and Tibshirani (2011) and Wang (2014).

Value

If a scalar value for lambda is provided, a list containing the following elements.

sigma	The estimate of the covariance matrix.
omega	The inverse of the estimated covariance matrix.
loglik	The (unpenalised) log-likelihood at the optimum.
loglikpen	The (penalised) log-likelihood at the optimum.
it	The number of iterations needed to reach convergence.

If a vector of values of lambda is provided, the output is a list in which each entry is itself a list, structured as above, associated with the corresponding value of lambda.

References

Chaudhuri, S., M. Drton, and T. S. Richardson (2007). Estimation of a covariance matrix with zeros. Biometrika 94 (1), 199–216.

Cibinel, L., A. Roverato, and V. Vinciotti (2024). A unified approach to penalized likelihood estimation of covariance matrices in high dimensions. arXiv, arXiv:2410.02403.

Bien, J. and R. J. Tibshirani (2011). Sparse estimation of a covariance matrix. Biometrika 98 (4), 807–820.

Wang, H. (2014). Coordinate descent algorithm for covariance graphical lasso. Statistics and Computing 24, 521–529.

Warton, D. I. (2008). Penalized normal likelihood and ridge regularization of correlation and covariance matrices. Journal of the American Statistical Association 103 (481), 340–349.

Examples

```
# An example with a banded covariance matrix
library(mvtnorm)

set.seed(1234)

p <- 10
n <- 500

# Create banded covariance matrix with three bands
band1 <- cbind(1:(p - 1), 2:p)
band2 <- cbind(1:(p - 2), 3:p)</pre>
```

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```
band3 <- cbind(1:(p - 3), 4:p)
idxs <- rbind(band1, band2, band3)</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[idxs] <- 0.5
Sigma <- Sigma + t(Sigma)
diag(Sigma) <- 2</pre>
# Generate data
data <- rmvnorm(n, sigma = Sigma)</pre>
# Fit a path of estimates
lambdas <- seq(0, 0.15, 0.01)
fit <- gicf(data, lambda = lambdas, kappa = 0.1)</pre>
# Explore one particular estimate
onefit <- fit[[5]]</pre>
image(onefit$sigma != 0)
# Redo the fit, but this time fix the correct sparsity pattern
fit2 <- gicf(data, lambda = lambdas, kappa = 0.1, adj = Sigma)</pre>
onefit2 <- fit2[[5]]</pre>
image(onefit2$sigma != 0)
```

Hyperparameters

Maximum effective range of regularisation parameters

Description

Compute the effective range of the regularisation parameters κ and λ .

Usage

```
lambdamax(S, kappa = 0, adj = 1 - diag(1, nrow(S)))
kappamax(S, lambda, adj = 1 - diag(1, nrow(S)))
```

Arguments

S The sample covariance matrix.

kappa, lambda The non-negative ridge regularisation/lasso shrinkage parameters.

An optional matrix whose pattern of zeroes is to be enforced onto the final output

of the Generalised Iterative Conditional Fitting algorithm.

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Details

These utility functions describe the boundary of the region

$$\mathcal{H} = \{(\kappa, \lambda) \in \mathbb{R}^2_{>0} : \lambda \le \lambda_{MAX}(\kappa)\},\$$

with

$$\lambda \leq \lambda_{MAX}(\kappa) \iff \kappa \leq \kappa_{MAX}(\lambda),$$

$$\lambda_{MAX}(\kappa) = \max_{i,j \in \mathcal{G}} \frac{|s_{ij}|}{(s_{ii} + \kappa)(s_{jj} + \kappa)},$$

$$\kappa_{MAX}(\lambda) = \max_{\substack{i,j \in \mathcal{G} \\ g_{ij}(\lambda) \geq 0}} \left\{ \sqrt{\frac{1}{4}(s_{ii} + s_{jj}) + g_{ij}(\lambda)} - \frac{1}{2}(s_{ii} + s_{jj}) \right\},$$

$$g_{ij}(\lambda) = \frac{|s_{ij}|}{\lambda} - s_{ii}s_{jj}.$$

Here S is the sample covariance matrix and \mathcal{G} is a graph whose adjacency matrix has the same sparsity pattern as adj. If the parameters (κ, λ) lay outside of \mathcal{H} , and the starting point of the Generalised Iterative Conditional Fitting algorithm is $\operatorname{diag}(S + \kappa I)$, then the output will also be $\operatorname{diag}(S + \kappa I)$.

Value

lambdamax returns a scalar value representing $\lambda_{MAX}(\kappa)$. kappamax returns a scalar value representing $\kappa_{MAX}(\lambda)$

Examples

```
# An example with a banded covariance matrix
library(mvtnorm)
set.seed(1234)
p <- 10
n <- 500
# Create banded covariance matrix with three bands
band1 <- cbind(1:(p - 1), 2:p)
band2 <- cbind(1:(p - 2), 3:p)
band3 <- cbind(1:(p - 3), 4:p)
idxs <- rbind(band1, band2, band3)
Sigma <- matrix(0, p, p)</pre>
Sigma[idxs] <- 0.5
Sigma <- Sigma + t(Sigma)</pre>
diag(Sigma) <- 2
# Generate data
data <- rmvnorm(n, sigma = Sigma)</pre>
S \leftarrow cov(data) * (n - 1)/n
```

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```
# Fix a value of lambda and compute k_max
lambda <- 0.07
k.max <- kappamax(S, lambda = lambda)
# Check that fit is diagonal
fit \leftarrow gicf(S = S, n = n, lambda = lambda, kappa = k.max)
image(fit$sigma != 0)
# Fix a value of kappa and compute l_max
kappa <- 1.15
1.max <- lambdamax(S, kappa = kappa)</pre>
# Check that fit is diagonal
fit <- gicf(S = S, n = n, lambda = l.max, kappa = kappa)
image(fit$sigma != 0)
# Repeat steps above, but with correct sparsity pattern specified
lambda <- 0.07
k.max <- kappamax(S, lambda = lambda, adj = Sigma)</pre>
fit <- gicf(S = S, n = n, lambda = lambda, kappa = k.max, adj = Sigma)
image(fit$sigma != 0)
kappa <- 1.15
1.max <- lambdamax(S, kappa = kappa, adj = Sigma)</pre>
fit <- gicf(S = S, n = n, lambda = l.max, kappa = kappa, adj = Sigma)</pre>
image(fit$sigma != 0)
```

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