

Package ‘matSPACE’

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Title Sparse Partial Correlation Estimation for Matrix-Variate Data

Version 0.1.0

Description Fits sparse partial correlation networks for matrix-variate data by extending the SPACE joint partial correlation estimation framework to a Kronecker-product covariance structure. All partial correlations are estimated simultaneously via an L1-penalized ('lasso') shooting algorithm within a single optimization framework, which preserves symmetry of the estimated network and avoids the tuning-parameter selection difficulties of separate node-wise regressions. Optional features include column reweighting, residual variance re-estimation across outer iterations, and automatic generation of a lasso penalty sequence for tuning.

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Encoding UTF-8

LinkingTo Rcpp

Imports Rcpp, stats

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URL <https://github.com/kimhyew1/matSPACE>

BugReports <https://github.com/kimhyew1/matSPACE/issues>

NeedsCompilation yes

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lambda.bound	Generate a log-spaced lasso penalty sequence for space()
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Description

Computes a decreasing, log-spaced sequence of K candidate lasso penalties (lam values) for tuning [space\(\)](#), analogous to the lambda_max-based grids used in lasso path algorithms. The largest value, lambda_max, is the largest off-diagonal entry of a variance-rescaled Gram matrix — the smallest penalty above which every off-diagonal partial correlation coefficient is driven to zero; the sequence descends geometrically to lambda_max * eps.

Usage

```
lambda.bound(dt, eps = 1e-06, K = 30)
```

Arguments

dt	list of n matrices, each p x q, in the same format expected by the data argument of space() .
eps	ratio of the smallest to the largest penalty in the returned sequence, i.e. lambda_min = lambda_max * eps.
K	number of penalty values to generate.

Value

A numeric vector of length K, decreasing geometrically from lambda_max to lambda_max * eps, suitable to pass one at a time as the lam argument of [space\(\)](#) (e.g. selecting among the fits with a BIC-type criterion).

Examples

```
set.seed(1)
p <- 5; q <- 4; n <- 3
data <- replicate(n, matrix(rnorm(p * q), p, q), simplify = FALSE)
lambda.bound(data, K = 10)
```

matSPACE	<i>Estimate row/column precision matrices for Kronecker-structured (matrix-variate) SPACE, with identifiability resolved</i>
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Description

Fits `space()` independently on the data (for the column precision V , $q \times q$) and on the transposed data (for the row precision U , $p \times p$), each over a lasso penalty path, selects the BIC-minimizing λ for U and for V separately at every scaling factor in `sf_vec`, and reconstructs the corresponding precision matrices (via `precision_from_parcor()`). Because the Kronecker product $\text{kronecker}(V, U)$ is invariant under $(V / c, U * c)$ for any $c > 0$, the pair is not separately identifiable from the data; the result is rescaled so that $V[1, 1] == 1$, using c equal to the raw fitted $V[1, 1]$ (U is multiplied by that same c), which preserves $\text{kronecker}(V, U)$ exactly.

Usage

```
matSPACE(
  data,
  lambda_V = NULL,
  lambda_U = NULL,
  K = 30,
  f_type_V = "equal",
  f_type_U = "equal",
  sf_vec = c(1, 1.5)
)
```

Arguments

<code>data</code>	list of n matrices, each $p \times q$; the matrix-variate observations, in the same format expected by the <code>data</code> argument of <code>space()</code> .
<code>lambda_V</code>	optional numeric vector of lasso penalties to use for the column (V , $q \times q$) fit. If <code>NULL</code> (default), generated via <code>lambda.bound()</code> on data with K values.
<code>lambda_U</code>	optional numeric vector of lasso penalties to use for the row (U , $p \times p$) fit. If <code>NULL</code> (default), generated via <code>lambda.bound()</code> on the transposed data with K values.
<code>K</code>	number of λ values to generate with <code>lambda.bound()</code> when <code>lambda_V</code> or <code>lambda_U</code> is <code>NULL</code> . Ignored for whichever of the two is supplied directly.
<code>f_type_V</code>	column weighting scheme forwarded to <code>space()</code> for the V fit; see <code>compute_weight()</code> .
<code>f_type_U</code>	column weighting scheme forwarded to <code>space()</code> for the U fit; see <code>compute_weight()</code> .
<code>sf_vec</code>	numeric vector of BIC scaling factors (the <code>sf</code> argument of <code>compute_BIC()</code>). A separate BIC-minimizing λ — and resulting identifiability-resolved (U , V) pair — is returned for each value.

Value

A named list, one element per value of `sf_vec` (named `sf_<value>`), each a list with components:

`V` `q x q` precision matrix at the BIC-minimizing `lambda_V`, rescaled so `V[1, 1] == 1`.

`U` `p x p` precision matrix at the BIC-minimizing `lambda_U`, rescaled by the same factor to preserve `kronecker(V, U)`.

`lambda_V, lambda_U` the BIC-minimizing lambda for `V` and `U` at this `sf`.

`BIC_V, BIC_U` the corresponding minimum BIC values.

`sf` the scaling factor this element was selected at.

The full lambda paths behind this selection are attached as a "path" attribute (`attr(fit, "path")`), a list with `V` and `U` components, each with `fits` (the raw `space()` fit at every lambda tried), `bic` (a `length(lambda) x length(sf_vec)` matrix, named `sf_<value>`), and `lambda`. Useful for plotting BIC (or the number of nonzero edges, from `fits[[i]]$ParCor`) against lambda without refitting.

Examples

```
set.seed(1)
p <- 4; q <- 3; n <- 3
data <- replicate(n, matrix(rnorm(p * q), p, q), simplify = FALSE)
fit <- matSPACE(data, K = 5, sf_vec = c(1, 1.5))
fit$sf_1$V
fit$sf_1$U
path <- attr(fit, "path")
plot(path$V$lambda, path$V$bic[, "sf_1"], type = "b")
```

space

Fit a matrix-variate SPACE sparse partial correlation model

Description

Estimates a sparse network of partial correlations among the `q` columns of matrix-variate data using an L1-penalized (lasso) SPACE-style shooting algorithm, with optional per-column reweighting and residual variance (`sig`) re-estimation across outer iterations.

Usage

```
space(
  data,
  lam,
  sig = NULL,
  f_type = "equal",
  iter = 2,
  beta_init = NULL,
  sig_init = NULL
)
```

Arguments

<code>data</code>	list of n matrices, each $p \times q$. All matrices share the same $p \times q$ shape; n is the number of independent replicates and the q columns are the variables whose pairwise partial correlations are estimated.
<code>lam</code>	lasso penalty applied to the off-diagonal partial correlation coefficients.
<code>sig</code>	optional length- q sigma vector; NULL = iterative estimate
<code>f_type</code>	"equal" "variance" "degree"
<code>iter</code>	number of outer iterations
<code>beta_init</code>	optional warm-start for beta: a $q \times q$ matrix/flat vector (row-major, $i*q+j$ layout) forwarded to the first internal <code>space_shooting()</code> call instead of the default univariate soft-threshold initialization. Only meaningful for the first outer iteration, since that is the only one whose <code>sigma_sr</code> ($\sqrt{\text{SIG}/\text{WEIGHT}}$) does not depend on <code>lam</code> — the first pass always starts from <code>SIG = rep(1, q)</code> , <code>WEIGHT = rep(1, q)</code> regardless of <code>lam</code> , so a beta fit at a neighboring lambda is a valid warm start there. Requires the compiled <code>space_shooting()</code> routine in <code>src/matSPACE.cpp</code> , which accepts a <code>beta_init</code> argument.
<code>sig_init</code>	optional length- q warm-start for the STARTING value of SIG when <code>SIG.update = TRUE</code> (i.e. <code>sig = NULL</code>). Unlike <code>sig</code> , this does not pin SIG constant — it is still re-estimated every outer iteration via <code>estimate_sigma()</code> ; it only replaces the <code>rep(1, q)</code> cold start at $i = 1$ with a neighboring lambda's converged SIG, so <code>f_type = "variance"/"degree"</code> don't force a uniform-weight first pass before ever seeing a realistic weighting.

Value

A list with components:

<code>ParCor</code>	$q \times q$ matrix of estimated partial correlations (diagonal = 1).
<code>beta</code>	$q \times q$ matrix of estimated regression coefficients.
<code>sig</code>	length- q vector of estimated column residual precisions (1/variance).
<code>f</code>	length- q vector of final per-column weights.
<code>total_iter</code>	total number of shooting iterations across all outer iterations.
<code>E</code>	list of n residual matrices ($p \times q$ each).

Examples

```
set.seed(1)
p <- 5; q <- 4; n <- 3
data <- replicate(n, matrix(rnorm(p * q), p, q), simplify = FALSE)
fit <- space(data, lam = 0.1, iter = 1)
fit$ParCor
```

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