

Package ‘rrpack’

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Title Reduced-Rank Regression

Version 0.1-13

Description Multivariate regression methodologies including classical reduced-rank regression (RRR) studied by Anderson (1951) <[doi:10.1214/aoms/1177729580](https://doi.org/10.1214/aoms/1177729580)> and Reinsel and Velu (1998) <[doi:10.1007/978-1-4757-2853-8](https://doi.org/10.1007/978-1-4757-2853-8)>, reduced-rank regression via adaptive nuclear norm penalization proposed by Chen et al. (2013) <[doi:10.1093/biomet/ast036](https://doi.org/10.1093/biomet/ast036)> and Mukherjee et al. (2015) <[doi:10.1093/biomet/asx080](https://doi.org/10.1093/biomet/asx080)>, robust reduced-rank regression (R4) proposed by She and Chen (2017) <[doi:10.1093/biomet/asx032](https://doi.org/10.1093/biomet/asx032)>, generalized/mixed-response reduced-rank regression (mRRR) proposed by Luo et al. (2018) <[doi:10.1016/j.jmva.2018.04.011](https://doi.org/10.1016/j.jmva.2018.04.011)>, row-sparse reduced-rank regression (SRRR) proposed by Chen and Huang (2012) <[doi:10.1080/01621459.2012.734178](https://doi.org/10.1080/01621459.2012.734178)>, reduced-rank regression with a sparse singular value decomposition (RSSVD) proposed by Chen et al. (2012) <[doi:10.1111/j.1467-9868.2011.01002.x](https://doi.org/10.1111/j.1467-9868.2011.01002.x)> and sparse and orthogonal factor regression (SOFAR) proposed by Uematsu et al. (2019) <[doi:10.1109/TIT.2019.2909889](https://doi.org/10.1109/TIT.2019.2909889)>.

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Author Kun Chen [aut, cre] (<<https://orcid.org/0000-0003-3579-5467>>),
Wenjie Wang [aut] (<<https://orcid.org/0000-0003-0363-3180>>),
Jun Yan [ctb] (<<https://orcid.org/0000-0003-4401-7296>>)

Maintainer Kun Chen <kun.chen@uconn.edu>

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R topics documented:

| | |
|------------------------|----|
| cv.mrrr | 2 |
| cv.rrr | 4 |
| cv.sofar | 5 |
| cv.srrr | 7 |
| mrrr | 8 |
| plot | 10 |
| r4 | 12 |
| rrpack-coef | 14 |
| rrr | 15 |
| rrr.cookD | 17 |
| rrr.fit | 18 |
| rrr.leverage | 19 |
| rrr.sim1 | 20 |
| rrr.sim2 | 21 |
| rrr.sim3 | 22 |
| rrr.sim4 | 23 |
| rrr.sim5 | 24 |
| rrs.fit | 25 |
| rssvd | 26 |
| sofar | 28 |
| srrr | 30 |
| summary | 32 |

| | |
|--------------|-----------|
| Index | 34 |
|--------------|-----------|

| | |
|---------|--|
| cv.mrrr | <i>Mixed-response reduced-rank regression with rank selected by cross validation</i> |
|---------|--|

Description

Mixed-response reduced-rank regression with rank selected by cross validation

Usage

```
cv.mrrr(
  Y,
  X,
  is.pca = NULL,
  offset = NULL,
  ctrl.id = c(),
  family = list(gaussian(), binomial(), poisson()),
  familygroup = NULL,
  maxrank = min(ncol(Y), ncol(X)),
  penstr = list(),
  init = list(),
```

```

control = list(),
nfold = 5,
foldid = NULL,
nlam = 20,
warm = FALSE
)

```

Arguments

| | |
|-------------|---|
| Y | response matrix |
| X | covariate matrix |
| is.pca | If TRUE, mixed principal component analysis with X=I |
| offset | matrix of the same dimension as Y for offset |
| ctrl.id | indices of unpenalized predictors |
| family | a list of family functions as used in glm |
| familygroup | a list of family indices of the responses |
| maxrank | integer giving the maximum rank allowed. |
| penstr | a list of penalty structure of SVD. |
| init | a list of initial values of kappaC0, kappaS0, C0, and S0 |
| control | a list of controlling parameters for the fitting |
| nfold | number of folds in cross validation |
| foldid | to specify the folds if desired |
| nlam | number of tuning parameters; not effective when using rank constrained estimation |
| warm | if TRUE, use warm start in fitting the solution paths |

Value

S3 mrrr object, a list containing

| | |
|-----|------------------------------------|
| fit | the output from the selected model |
| dev | deviance measures |

Examples

```

## Not run:
library(rrpack)
simdata <- rrr.sim3(n = 100, p = 30, q.mix = c(5, 20, 5),
                  nrank = 2, mis.prop = 0.2)

Y <- simdata$Y
Y_mis <- simdata$Y.mis
X <- simdata$X
X0 <- cbind(1,X)
C <- simdata$C
family <- simdata$family
familygroup <- simdata$familygroup

```

```

svdX0d1 <- svd(X0)$d[1]
init1 = list(kappaC0 = svdX0d1 * 5)
offset = NULL
control = list(epsilon = 1e-4, sv.tol = 1e-2, maxit = 2000,
              trace = FALSE, gammaC0 = 1.1, plot.cv = TRUE,
              conv.obj = TRUE)
fit.cv.mrrr <- cv.mrrr(Y_mis, X, family = family,
                    familygroup = familygroup,
                    maxrank = 20,
                    penstr = list(penaltySVD = "rankCon",
                                  lambdaSVD = c(1 : 6)),
                    control = control, init = init1,
                    nfold = 10, nlam = 50)

summary(fit.cv.mrrr)
coef(fit.cv.mrrr)
fit.mrrr <- fit.cv.mrrr$fit

## plot(svd(fit.mrrr$coef[- 1,])$d)
plot(C ~ fit.mrrr$coef[- 1, ])
abline(a = 0, b = 1)

## End(Not run)

```

cv.rrr

Reduced-rank regression with rank selected by cross validation

Description

Reduced-rank regression with rank selected by cross validation

Usage

```

cv.rrr(
  Y,
  X,
  nfold = 10,
  maxrank = min(dim(Y), dim(X)),
  norder = NULL,
  coefSVD = FALSE
)

```

Arguments

| | |
|---------|---|
| Y | response matrix |
| X | covariate matrix |
| nfold | number of folds |
| maxrank | maximum rank allowed |
| norder | for constructing the folds |
| coefSVD | If TRUE, svd of the coefficient is returned |

Value

a list containing rr estimates from cross validation

References

Chen, K., Dong, H. and Chan, K.-S. (2013) Reduced rank regression via adaptive nuclear norm penalization. *Biometrika*, 100, 901–920.

Examples

```
library(rrpack)
p <- 50; q <- 50; n <- 100; nrank <- 3
mydata <- rrr.sim1(n, p, q, nrank, s2n = 1, sigma = NULL,
                 rho_X = 0.5, rho_E = 0.3)
rfit_cv <- with(mydata, cv.rrr(Y, X, nfold = 10, maxrank = 10))
summary(rfit_cv)
coef(rfit_cv)
```

cv.sofar

Sparse orthogonal factor regression tuned by cross validation

Description

Sparse orthogonal factor regression tuned by cross validation

Usage

```
cv.sofar(
  Y,
  X,
  nrank = 1,
  su = NULL,
  sv = NULL,
  nfold = 5,
  norder = NULL,
  modstr = list(),
  control = list(),
  screening = FALSE
)
```

Arguments

| | |
|-------|---|
| Y | response matrix |
| X | covariate matrix |
| nrank | an integer specifying the desired rank/number of factors |
| su | a scaling vector for U such that $U^T U = \text{diag}(s_u)$ |

| | |
|-----------|--|
| sv | a scaling vector for V such that $V^T V = \text{diag}(s_v)$ |
| nfold | number of fold; used for cv.sofar |
| norder | observation orders to constrect data folds; used for cv.sofar |
| modstr | a list of internal model parameters controlling the model fitting |
| control | a list of internal computation parameters controlling optimization |
| screening | If TRUE, marginal screening via lasso is performed before sofar fitting. |

Details

The model parameters can be specified through argument `modstr`. The available elements include

- `mu`: parameter in the augmented Lagrangian function.
- `mugamma`: increment of `mu` along iterations to speed up computation.
- `WA`: weight matrix for A .
- `WB`: weight matrix for B .
- `Wd`: weight matrix for d .
- `wgamma`: power parameter in constructing adaptive weights.

The model fitting can be controled through argument `control`. The avilable elements include

- `nlam`: number of lambda triplets to be used.
- `lam.min.factor`: set the smallest lambda triplets as a fraction of the estimation `lambda.max` triplets.
- `lam.max.factor`: set the largest lambda triplets as a multiple of the estimation `lambda.max` triplets.
- `lam.AB.factor`: set the relative penalty level between A/B and D .
- `penA,penB,penD`: if TRUE, penalty is applied.
- `lamA`: sequence of tuning parameters for A .
- `lamB`: sequence of tuning parameters for B .
- `lamD`: sequence of tuning parameters for d .
- `methodA`: penalty for penalizing A .
- `methodB`: penalty for penalizing B .
- `epsilon`: convergence tolerance.
- `maxit`: maximum number of iterations.
- `innerEpsilon`: convergence tolerance for inner subroutines.
- `innerMaxit`: maximum number of iterations for inner subroutines.
- `sv.tol`: tolerance for singular values.

 cv.srrr

 Row-sparse reduced-rank regression tuned by cross validation

Description

Row-sparse reduced-rank regression tuned by cross validation

Usage

```
cv.srrr(
  Y,
  X,
  nrank = 1,
  method = c("glasso", "adglasso"),
  nfold = 5,
  norder = NULL,
  A0 = NULL,
  V0 = NULL,
  modstr = list(),
  control = list()
)
```

Arguments

| | |
|---------|--|
| Y | response matrix |
| X | covariate matrix |
| nrank | prespecified rank |
| method | group lasso or adaptive group lasso |
| nfold | fold number |
| norder | for constructing the folds |
| A0 | initial value |
| V0 | initial value |
| modstr | a list of model parameters controlling the model fitting |
| control | a list of parameters for controlling the fitting process |

Details

Model parameters controlling the model fitting can be specified through argument `modstr`. The available elements include

- `lamA`: tuning parameter sequence.
- `nlam`: number of tuning parameters; no effect if `lamA` is specified.
- `minLambda`: minimum lambda value, no effect if `lamA` is specified.
- `maxLambda`: maximum lambda value, no effect if `lamA` is specified.

- WA: adaptive weights. If NULL, the weights are constructed from RRR.
- wgamma: power parameter for constructing adaptive weights.

Similarly, the computational parameters controlling optimization can be specified through argument control. The available elements include

- epsilon: epsilon convergence tolerance.
- maxit: maximum number of iterations.
- inner.eps: used in inner loop.
- inner.maxit: used in inner loop.

Value

A list of fitting results

References

Chen, L. and Huang, J.Z. (2012) Sparse reduced-rank regression for simultaneous dimension reduction and variable selection. *Journal of the American Statistical Association*. 107:500, 1533–1545.

mrrr

Generalized or mixed-response reduced-rank regression

Description

Performs either rank constrained maximum likelihood estimation or singular value penalized estimation.

Usage

```
mrrr(
  Y,
  X,
  is.pca = NULL,
  offset = NULL,
  ctrl.id = c(),
  family = list(gaussian(), binomial()),
  familygroup = NULL,
  maxrank = min(ncol(Y), ncol(X)),
  penstr = list(),
  init = list(),
  control = list()
)
```


Arguments

| | |
|-------------|--|
| Y | response matrix |
| X | covariate matrix |
| is.pca | If TRUE, mixed principal component analysis with $X=I$ |
| offset | matrix of the same dimension as Y for offset |
| ctrl.id | indices of unpenalized predictors |
| family | a list of family functions as used in glm |
| familygroup | a list of family indices of the responses |
| maxrank | integer giving the maximum rank allowed. Usually this can be set to $\min(n,p,q)$ |
| penstr | a list of penalty structure of SVD, contains <code>penstr\$penaltySVD</code> is the penalty of SVD, <code>penstr\$lambdaSVD</code> is the regularization parameter |
| init | a list of initial values of <code>kappaC0</code> , <code>kappaS0</code> , <code>C0</code> , and <code>S0</code> |
| control | a list of controlling parameters for the fitting |

Details

The model fitting process can be fine tuned through argument `control`. The available elements for `control` include

- `epsilon`: positive convergence tolerance `epsilon`; the iterations converge when $|new - old| / (old + 0.1) < epsilon$. treated as zero.
- `sv.tol`: tolerance for singular values.
- `maxit`: integer giving the maximal number of iterations.
- `trace`: logical indicating if tracing the objective is needed.
- `conv.obj`: if TRUE, track objective function.
- `equal.phi`: if TRUE, use a single dispersion parameter for Gaussian responses.
- `plot.obj`: if TRUE, plot obj values along iterations; for checking only
- `plot.cv`: if TRUE, plot cross validation error.
- `gammaC0`: adaptive scaling to speed up computation.

Similarly, the available elements for arguments `penstr` specifying penalty structure of SVD include

- `penaltySVD`: penalty for reducing rank
- `lambdaSVD`: tuning parameter. For `penaltySVD = rankCon`, this is the specified rank.

Value

S3 `mrrr` object, a list containing

| | |
|------------------------|----------------------------------|
| <code>obj</code> | the objective function tracking |
| <code>converged</code> | TRUE/FALSE for convergence |
| <code>coef</code> | the estimated coefficient matrix |
| <code>outlier</code> | the estimated outlier matrix |
| <code>nrnk</code> | the rank of the fitted model |

Examples

```

library(rrpack)
simdata <- rrr.sim3(n = 100, p = 30, q.mix = c(5, 20, 5),
                  nrank = 2, mis.prop = 0.2)

Y <- simdata$Y
Y_mis <- simdata$Y.mis
X <- simdata$X
X0 <- cbind(1, X)
C <- simdata$C
family <- simdata$family
familygroup <- simdata$familygroup
svdX0d1 <- svd(X0)$d[1]
init1 = list(kappaC0 = svdX0d1 * 5)
offset = NULL
control = list(epsilon = 1e-4, sv.tol = 1e-2, maxit = 2000,
              trace = FALSE, gammaC0 = 1.1, plot.cv = TRUE,
              conv.obj = TRUE)
fit.mrrr <- mrrr(Y_mis, X, family = family, familygroup = familygroup,
               penstr = list(penaltySVD = "rankCon", lambdaSVD = 2),
               control = control, init = init1)
summary(fit.mrrr)
coef(fit.mrrr)
par(mfrow = c(1, 2))
plot(fit.mrrr$obj)
plot(C ~ fit.mrrr$coef[- 1 ,])
abline(a = 0, b = 1)

```

plot

Scatter Plot

Description

S3 methods generating scatter plot for some objects generated by rrpck using ggplot2. An ggplot2 object is returned so that users are allowed to easily further customize the plot.

Usage

```

## S3 method for class 'rrr'
plot(
  x,
  y = NULL,
  layer = 1L,
  xlab = paste("latent predictor ", layer, sep = ""),
  ylab = paste("latent response ", layer, sep = ""),
  ...
)

## S3 method for class 'sofar'
plot(

```

```
x,  
y = NULL,  
layer = 1L,  
xlab = paste("latent predictor ", layer, sep = ""),  
ylab = paste("latent response ", layer, sep = ""),  
...  
)  
  
## S3 method for class 'cv.sofar'  
plot(  
  x,  
  y = NULL,  
  layer = 1L,  
  xlab = paste("latent predictor ", layer, sep = ""),  
  ylab = paste("latent response ", layer, sep = ""),  
  ...  
)  
  
## S3 method for class 'srrr'  
plot(  
  x,  
  y = NULL,  
  layer = 1L,  
  xlab = paste("latent predictor ", layer, sep = ""),  
  ylab = paste("latent response ", layer, sep = ""),  
  ...  
)  
  
## S3 method for class 'cv.srrr'  
plot(  
  x,  
  y = NULL,  
  layer = 1L,  
  xlab = paste("latent predictor ", layer, sep = ""),  
  ylab = paste("latent response ", layer, sep = ""),  
  ...  
)  
  
## S3 method for class 'rssvd'  
plot(  
  x,  
  y = NULL,  
  layer = 1L,  
  xlab = paste("latent predictor ", layer, sep = ""),  
  ylab = paste("latent response ", layer, sep = ""),  
  ...  
)
```

Arguments

| | |
|-------|---|
| x | Some object generated by rrpck. |
| y | NULL. Do not need to specify. |
| layer | The unit-rank layer to plot; cannot be larger than the estimated rank |
| xlab | Label of X axis. |
| ylab | Label of Y axis. |
| ... | Other arguments for future usage. |

Value

ggplot2 object.

 r4

Robust reduced-rank regression

Description

Perform robust reduced-rank regression.

Usage

```
r4(
  Y,
  X,
  maxrank = min(dim(Y), dim(X)),
  method = c("rowl0", "rowl1", "entrywise"),
  Gamma = NULL,
  ic.type = c("AIC", "BIC", "PIC"),
  modstr = list(),
  control = list()
)
```

Arguments

| | |
|---------|--|
| Y | a matrix of response (n by q) |
| X | a matrix of covariate (n by p) |
| maxrank | maximum rank for fitting |
| method | outlier detection method, either entrywise or rowwise |
| Gamma | weighting matrix in the loss function |
| ic.type | information criterion, AIC, BIC or PIC |
| modstr | a list of model parameters controlling the model fitting |
| control | a list of parameters for controlling the fitting process |

Details

The model parameters can be controlled through argument `modstr`. The available elements include

- `nlam`: parameter in the augmented Lagrangian function.
- `adaptive`: if TRUE, use leverage values for adaptive penalization. The default value is FALSE.
- `weights`: user supplied weights for adaptive penalization.
- `minlam`: maximum proportion of outliers.
- `maxlam`: maximum proportion of good observations.
- `delid`: discarded observation indices for initial estimation.

The model fitting can be controlled through argument `control`. The available elements include

- `epsilon`: convergence tolerance.
- `maxit`: maximum number of iterations.
- `qr.tol`: tolerance for qr decomposition.
- `tol`: tolerance.

Value

a list consisting of

| | |
|-----------------------------|--|
| <code>coef.path</code> | solutuon path of regression coefficients |
| <code>s.path</code> | solutuon path of sparse mean shifts |
| <code>s.norm.path</code> | solutuon path of the norms of sparse mean shifts |
| <code>ic.path</code> | paths of information criteria |
| <code>ic.smooth.path</code> | smoothed paths of information criteria |
| <code>lambda.path</code> | paths of the tuning parameter |
| <code>id.solution</code> | ids of the selected solutions on the path |
| <code>ic.best</code> | lowest values of the information criteria |
| <code>rank.best</code> | rank values of selected solutions |
| <code>coef</code> | estimated regression coefficients |
| <code>s</code> | estimated sparse mean shifts |
| <code>rank</code> | rank estimate |

References

She, Y. and Chen, K. (2017) Robust reduced-rank regression. *Biometrika*, 104 (3), 633–647.

Examples

```
## Not run:
library(rrpack)
n <- 100; p <- 500; q <- 50
xrank <- 10; nrank <- 3; rmax <- min(n, p, q, xrank)
nlam <- 100; gamma <- 2
rho_E <- 0.3
rho_X <- 0.5
nlev <- 0
vlev <- 0
vout <- NULL
vlevsd <- NULL
nout <- 0.1 * n
s2n <- 1
voutsd <- 2
simdata <- rrr.sim5(n, p, q, nrank, rx = xrank, s2n = s2n,
                   rho_X = rho_X, rho_E = rho_E, nout = nout, vout = vout,
                   voutsd = voutsd, nlev = nlev, vlev = vlev, vlevsd = vlevsd)

Y <- simdata$Y
X <- simdata$X
fit <- r4(Y, X, maxrank = rmax,
          method = "rowl0", ic.type = "PIC")

summary(fit)
coef(fit)
which(apply(fit$s, 1, function(a) sum(a^2)) != 0)

## End(Not run)
```

rrpack-coef

Estimated coefficients

Description

S3 methods extracting estimated coefficients for objects generated by rrpack.

Usage

```
## S3 method for class 'mrrr'
coef(object, ...)

## S3 method for class 'cv.mrrr'
coef(object, ...)

## S3 method for class 'r4'
coef(object, ...)

## S3 method for class 'rrr'
coef(object, ...)
```

```
## S3 method for class 'rrr.fit'
coef(object, ...)

## S3 method for class 'cv.rrr'
coef(object, ...)

## S3 method for class 'srrr'
coef(object, ...)

## S3 method for class 'sofar'
coef(object, ...)

## S3 method for class 'rssvd'
coef(object, ...)
```

Arguments

| | |
|--------|-----------------------------------|
| object | Object generated by rrpck. |
| ... | Other arguments for future usage. |

Value

A numeric matrix.

 rrr

Multivariate reduced-rank linear regression

Description

Produce solution paths of reduced-rank estimators and adaptive nuclear norm penalized estimators; compute the degrees of freedom of the RRR estimators and select a solution via certain information criterion.

Usage

```
rrr(
  Y,
  X,
  penaltySVD = c("rank", "ann"),
  ic.type = c("GIC", "AIC", "BIC", "BICP", "GCV"),
  df.type = c("exact", "naive"),
  maxrank = min(dim(Y), dim(X)),
  modstr = list(),
  control = list()
)
```

Arguments

| | |
|-------------------------|--|
| <code>Y</code> | a matrix of response (n by q) |
| <code>X</code> | a matrix of covariate (n by p) |
| <code>penaltySVD</code> | 'rank': rank-constrained estimation; 'ann': adaptive nuclear norm estimation. |
| <code>ic.type</code> | the information criterion to be used; currently supporting 'AIC', 'BIC', 'BICP', 'GCV', and 'GIC'. |
| <code>df.type</code> | 'exact': the exact degrees of freedoms based on SURE theory; 'naive': the naive degree of freedoms based on counting number of free parameters |
| <code>maxrank</code> | an integer of maximum desired rank. |
| <code>modstr</code> | a list of model parameters controlling the model fitting |
| <code>control</code> | a list of parameters for controlling the fitting process: 'sv.tol' controls the tolerance of singular values; 'qr.tol' controls the tolerance of QR decomposition for the LS fit |

Details

Model parameters can be specified through argument `modstr`. The available include

- `gamma`: A scalar power parameter of the adaptive weights in `penalty == "ann"`.
- `nlambda`: The number of lambda values; no effect if `penalty == "count"`.
- `lambda`: A vector of user-specified rank values if `penalty == "count"` or a vector of penalty values if `penalty == "ann"`.

The available elements for argument `control` include

- `sv.tol`: singular value tolerance.
- `qr.tol`: QR decomposition tolerance.

Value

S3 `rrr` object, a list consisting of

| | |
|-------------------------|---|
| <code>call</code> | original function call |
| <code>Y</code> | input matrix of response |
| <code>X</code> | input matrix of covariate |
| <code>A</code> | right singular matrix of the least square fitted matrix |
| <code>Ad</code> | a vector of squared singular values of the least square fitted matrix |
| <code>coef.ls</code> | coefficient estimate from LS |
| <code>Spath</code> | a matrix, each column containing shrinkage factors of the singular values of a solution; the first four objects can be used to recover all reduced-rank solutions |
| <code>df.exact</code> | the exact degrees of freedom |
| <code>df.naive</code> | the naive degrees of freedom |
| <code>penaltySVD</code> | the method of low-rank estimation |
| <code>sse</code> | a vector of sum of squared errors |

| | |
|------|--|
| ic | a vector of information criterion |
| coef | estimated coefficient matrix |
| U | estimated left singular matrix such that XU/\sqrt{n} is orthogonal |
| V | estimated right singular matrix that is orthogonal |
| D | estimated singular value matrix such that $C = UDV^t$ |
| rank | estimated rank |

References

Chen, K., Dong, H. and Chan, K.-S. (2013) Reduced rank regression via adaptive nuclear norm penalization. *Biometrika*, 100, 901–920.

Examples

```
library(rrpack)
p <- 50; q <- 50; n <- 100; nrank <- 3
mydata <- rrr.sim1(n, p, q, nrank, s2n = 1, sigma = NULL,
                 rho_X = 0.5, rho_E = 0.3)
rfit <- with(mydata, rrr(Y, X, maxrank = 10))
summary(rfit)
coef(rfit)
plot(rfit)
```

rrr.cookD

Cook's distance in reduced-rank regression for model diagnostics

Description

Compute Cook's distance for model diagnostics in rrr estimation.

Usage

```
rrr.cookD(Y, X = NULL, nrank = 1, qr.tol = 1e-07)
```

Arguments

| | |
|--------|------------------|
| Y | response matrix |
| X | covariate matrix |
| nrank | model rank |
| qr.tol | tolerance |

Value

a list containing diagnostics measures

References

Chen, K. Model diagnostics in reduced-rank estimation. *Statistics and Its interface*, 9, 469–484.

rrr.fit

*Fitting reduced-rank regression with a specific rank***Description**

Given a response matrix and a covariate matrix, this function fits reduced rank regression for a specified rank. It reduces to singular value decomposition if the covariate matrix is the identity matrix.

Usage

```
rrr.fit(Y, X, nrank = 1, weight = NULL, coefSVD = FALSE)
```

Arguments

| | |
|---------|---|
| Y | a matrix of response (n by q) |
| X | a matrix of covariate (n by p) |
| nrank | an integer specifying the desired rank |
| weight | a square matrix of weight (q by q); The default is the identity matrix |
| coefSVD | logical indicating the need for SVD for the coefficient matrix in the output; used in ssvd estimation |

Value

S3 rrr object, a list consisting of

| | |
|-----------|------------------------------|
| coef | coefficient of rrr |
| coef.ls | coefficient of least square |
| fitted | fitted value of rrr |
| fitted.ls | fitted value of least square |
| A | right singular matrix |
| Ad | a vector of singular values |
| rank | rank of the fitted rrr |

Examples

```
Y <- matrix(rnorm(400), 100, 4)
X <- matrix(rnorm(800), 100, 8)
rfit <- rrr.fit(Y, X, nrank = 2)
coef(rfit)
```

| | |
|--------------|---|
| rrr.leverage | <i>Leverage scores and Cook's distance in reduced-rank regression for model diagnostics</i> |
|--------------|---|

Description

Compute leverage scores and Cook's distance for model diagnostics in rrr estimation.

Usage

```
rrr.leverage(Y, X = NULL, nrank = 1, qr.tol = 1e-07)
```

Arguments

| | |
|--------|--|
| Y | a matrix of response (n by q) |
| X | a matrix of covariate (n by p) |
| nrank | an integer specifying the desired rank |
| qr.tol | tolerance to be passed to 'qr' |

Value

'rrr.leverage' returns a list containing a vector of leverages and a scalar of the degrees of freedom (sum of leverages). 'rrr.cooks' returns a list containing

| | |
|-----------|--------------------|
| residuals | residuals matrix |
| mse | mean squared error |
| leverage | leverage |
| cookD | Cook's distance |
| df | degrees of freedom |

References

Chen, K. Model diagnostics in reduced-rank estimation. *Statistics and Its interface*, 9, 469–484.

`rrr.sim1`*Simulation model 1*

Description

Similar to the the RSSVD simulation model in Chen, Chan, Stenseth (2012), JRSSB.

Usage

```
rrr.sim1(  
  n = 50,  
  p = 25,  
  q = 25,  
  nrank = 3,  
  s2n = 1,  
  sigma = NULL,  
  rho_X = 0.5,  
  rho_E = 0  
)
```

Arguments

| | |
|----------------------|---|
| <code>n, p, q</code> | model dimensions |
| <code>nrank</code> | model rank |
| <code>s2n</code> | signal to noise ratio |
| <code>sigma</code> | error variance. If specified, then <code>s2n</code> has no effect |
| <code>rho_X</code> | correlation parameter in the generation of predictors |
| <code>rho_E</code> | correlation parameter in the generation of random errors |

Value

simulated model and data

References

Chen, K., Chan, K.-S. and Stenseth, N. C. (2012) Reduced rank stochastic regression with a sparse singular value decomposition. *Journal of the Royal Statistical Society: Series B*, 74, 203–221.

`rrr.sim2`*Simulation model 2*

Description

Similar to the the SRRR simulation model in Chen and Huang (2012), JASA

Usage

```
rrr.sim2(  
  n = 100,  
  p = 50,  
  p0 = 10,  
  q = 50,  
  q0 = 10,  
  nrank = 3,  
  s2n = 1,  
  sigma = NULL,  
  rho_X = 0.5,  
  rho_E = 0  
)
```

Arguments

| | |
|--------------------|---|
| <code>n</code> | sample size |
| <code>p</code> | number of predictors |
| <code>p0</code> | number of relevant predictors |
| <code>q</code> | number of responses |
| <code>q0</code> | number of relevant responses |
| <code>nrank</code> | model rank |
| <code>s2n</code> | signal to noise ratio |
| <code>sigma</code> | error variance. If specified, then <code>s2n</code> has no effect |
| <code>rho_X</code> | correlation parameter in the generation of predictors |
| <code>rho_E</code> | correlation parameter in the generation of random errors |

Value

simulated model and data

References

Chen, L. and Huang, J.Z. (2012) Sparse reduced-rank regression for simultaneous dimension reduction and variable selection. *Journal of the American Statistical Association*, 107:500, 1533–1545.

`rrr.sim3`*Simulation model 3*

Description

Generate data from a mixed-response reduced-rank regression model

Usage

```
rrr.sim3(  
  n = 100,  
  p = 30,  
  q.mix = c(5, 20, 5),  
  nrank = 2,  
  intercept = rep(0.5, 30),  
  mis.prop = 0.2  
)
```

Arguments

| | |
|------------------------|---|
| <code>n</code> | sample size |
| <code>p</code> | number of predictors |
| <code>q.mix</code> | numbers of Gaussian, Bernolli and Poisson responses |
| <code>nrank</code> | model rank |
| <code>intercept</code> | a vector of intercept |
| <code>mis.prop</code> | missing proportion |

Value

simulated model and data

References

Chen, K., Luo, C., and Liang, J. (2017) Leveraging mixed and incomplete outcomes through a mixed-response reduced-rank regression. *Technical report*.

rrr.sim4

*Simulation model 4***Description**

Generate data from a mean-shifted reduced-rank regression model

Usage

```
rrr.sim4(
  n = 100,
  p = 12,
  q = 8,
  nrank = 3,
  s2n = 1,
  rho_X = 0,
  rho_E = 0,
  nout = 10,
  vout = NULL,
  voutsd = 2,
  nlev = 10,
  vlev = 10,
  vlevsd = NULL,
  SigmaX = "CorrCS",
  SigmaE = "CorrCS"
)
```

Arguments

| | |
|--------|--|
| n | sample size |
| p | number of predictors |
| q | numbers of responses |
| nrank | model rank |
| s2n | signal to noise ratio |
| rho_X | correlation parameter for predictors |
| rho_E | correlation parameter for errors |
| nout | number of outliers; should be smaller than n |
| vout | control mean-shifted value of outliers |
| voutsd | control mean-shifted magnitude of outliers |
| nlev | number of high-leverage outliers |
| vlev | control value of leverage |
| vlevsd | control magnitude of leverage |
| SigmaX | correlation structure of predictors |
| SigmaE | correlation structure of errors |

Value

simulated model and data

References

She, Y. and Chen, K. (2017) Robust reduced-rank regression. *Biometrika*, 104 (3), 633–647.

rrr.sim5

Simulation model 5

Description

Generate data from a mean-shifted reduced-rank regression model

Usage

```
rrr.sim5(  
  n = 40,  
  p = 100,  
  q = 50,  
  nrank = 5,  
  rx = 10,  
  s2n = 1,  
  rho_X = 0,  
  rho_E = 0,  
  nout = 10,  
  vout = NULL,  
  voutsd = 2,  
  nlev = 10,  
  vlev = 10,  
  vlevsd = NULL,  
  SigmaX = "CorrCS",  
  SigmaE = "CorrCS"  
)
```

Arguments

| | |
|-------|--------------------------------------|
| n | sample size |
| p | number of predictors |
| q | numbers of responses |
| nrank | model rank |
| rx | rank of the design matrix |
| s2n | signal to noise ratio |
| rho_X | correlation parameter for predictors |
| rho_E | correlation parameter for errors |

| | |
|--------|--|
| nout | number of outliers; should be smaller than n |
| vout | control mean-shifted value of outliers |
| voutsd | control mean-shifted magnitude of outliers |
| nlev | number of high-leverage outliers |
| vlev | control value of leverage |
| vlevsd | control magnitude of leverage |
| SigmaX | correlation structure of predictors |
| SigmaE | correlation structure of errors |

Value

simulated model and data

References

She, Y. and Chen, K. (2017) Robust reduced-rank regression. *Biometrika*, 104 (3), 633–647.

| | |
|---------|--|
| rrs.fit | <i>Fitting reduced-rank ridge regression with given rank and shrinkage penalty</i> |
|---------|--|

Description

Fitting reduced-rank ridge regression with given rank and shrinkage penalty

Usage

```
rrs.fit(Y, X, nrank = min(ncol(Y), ncol(X)), lambda = 1, coefSVD = FALSE)
```

Arguments

| | |
|---------|--|
| Y | a matrix of response (n by q) |
| X | a matrix of covariate (n by p) |
| nrank | an integer specifying the desired rank |
| lambda | tuning parameter for the ridge penalty |
| coefSVD | logical indicating the need for SVD for the coefficient matrix in the output |

Value

S3 rrr object, a list consisting of

| | |
|-----------|------------------------------|
| coef | coefficient of rrs |
| coef.ls | coefficient of least square |
| fitted | fitted value of rrs |
| fitted.ls | fitted value of least square |
| A | right singular matrix |
| Ad | sigular value vector |
| nrank | rank of the fitted rrr |

References

Mukherjee, A. and Zhu, J. (2011) Reduced rank ridge regression and its kernal extensions.

Mukherjee, A., Chen, K., Wang, N. and Zhu, J. (2015) On the degrees of freedom of reduced-rank estimators in multivariate regression. *Biometrika*, 102, 457–477.

Examples

```
library(rrpack)
Y <- matrix(rnorm(400), 100, 4)
X <- matrix(rnorm(800), 100, 8)
rfit <- rrs.fit(Y, X)
```

 rssvd

Reduced-rank regression with a sparse singular value decomposition

Description

Reduced-rank regression with a sparse singular value decomposition using the iterative exclusive extraction algorithm.

Usage

```
rssvd(
  Y,
  X,
  nrank,
  ic.type = c("BIC", "BICP", "AIC"),
  orthX = FALSE,
  control = list(),
  screening = FALSE
)
```

Arguments

| | |
|-----------|--|
| Y | response matrix |
| X | covariate matrix |
| nrank | integer specification of the desired rank |
| ic.type | character specifying which information criterion to use to select the best: 'BIC', 'BICP', and 'AIC' |
| orthX | logical indicating if X is orthogonal, in which case a faster algorithm is used |
| control | a list of parameters controlling the fitting process |
| screening | If TRUE, marginal screening via glm is performed before srrr fitting. |

Details

The model fitting can be controlled through argument `control`. The available elements include

- `maxit`: maximum number of iterations.
- `epsilon`: convergence tolerance.
- `innerMaxit`: maximum number of iterations for inner steps.
- `innerEpsilon`: convergence tolerance for inner steps.
- `nlambda`: number of tuning parameters.
- `adaptive`: if TRUE, use adaptive penalization.
- `gamma0`: power parameter for constructing adaptive weights.
- `minLambda`: multiply factor to determine the minimum lambda.
- `niter.eea`: the number of iterations in the iterative exclusive extraction algorithm.
- `df.tol`: tolerance.

Value

S3 `rssvd.path` object, a list consisting of

| | |
|--------------------|--|
| <code>Upath</code> | solution path of U |
| <code>Vpath</code> | solution path of V |
| <code>Dpath</code> | solution path of D |
| <code>U</code> | estimated left singular matrix that is orthogonal |
| <code>V</code> | estimated right singular matrix that is orthogonal |
| <code>D</code> | estimated singular values such that $C=UDV^t$ |
| <code>rank</code> | estimated rank |

References

Chen, K., Chan, K.-S. and Stenseth, N. C. (2012) Reduced rank stochastic regression with a sparse singular value decomposition. *Journal of the Royal Statistical Society: Series B*, 74, 203–221.

Examples

```

library(rrpack)
## Simulate data from a sparse factor regression model
p <- 50; q <- 50; n <- 100; nrank <- 3
mydata <- rrr.sim1(n, p, q, nrank, s2n = 1, sigma = NULL,
                 rho_X = 0.5, rho_E = 0.3)
fit1 <- with(mydata, rssid(Y, X, nrank = nrank + 1))
summary(fit1)
plot(fit1)

```

sofar

*Sparse orthogonal factor regression***Description**

Compute solution paths of sparse orthogonal factor regression

Usage

```

sofar(
  Y,
  X,
  nrank = 1,
  su = NULL,
  sv = NULL,
  ic.type = c("GIC", "BIC", "AIC", "GCV"),
  modstr = list(),
  control = list(),
  screening = FALSE
)

```

Arguments

| | |
|-----------|--|
| Y | response matrix |
| X | covariate matrix |
| nrank | an integer specifying the desired rank/number of factors |
| su | a scaling vector for U such that $U^T U = \text{diag}(s_u)$. |
| sv | a scaling vector for V such that $V^T V = \text{diag}(s_v)$. |
| ic.type | select tuning method; the default is GIC |
| modstr | a list of internal model parameters controlling the model fitting |
| control | a list of internal computation parameters controlling optimization |
| screening | If TRUE, marginal screening via lasso is performed before sofar fitting. |

Details

The model parameters can be specified through argument `modstr`. The available elements include

- `mu`: parameter in the augmented Lagrangian function.
- `mugamma`: increment of `mu` along iterations to speed up computation.
- `WA`: weight matrix for A.
- `WB`: weight matrix for B.
- `Wd`: weight matrix for d.
- `wgamma`: power parameter in constructing adaptive weights.

The model fitting can be controlled through argument `control`. The available elements include

- `nlam`: number of lambda triplets to be used.
- `lam.min.factor`: set the smallest lambda triplets as a fraction of the estimation `lambda.max` triplets.
- `lam.max.factor`: set the largest lambda triplets as a multiple of the estimation `lambda.max` triplets.
- `lam.AB.factor`: set the relative penalty level between A/B and D.
- `penA,penB,penD`: if TRUE, penalty is applied.
- `lamA`: sequence of tuning parameters for A.
- `lamB`: sequence of tuning parameters for B.
- `lamD`: sequence of tuning parameters for d.
- `methodA`: penalty for penalizing A.
- `methodB`: penalty for penalizing B.
- `epsilon`: convergence tolerance.
- `maxit`: maximum number of iterations.
- `innerEpsilon`: convergence tolerance for inner subroutines.
- `innerMaxit`: maximum number of iterations for inner subroutines.
- `sv.tol`: tolerance for singular values.

Value

A `sofar` object containing

| | |
|---------------------|------------------------------|
| <code>call</code> | original function call |
| <code>Y</code> | input response matrix |
| <code>X</code> | input predictor matrix |
| <code>Upath</code> | solution path of U |
| <code>Dpath</code> | solution path of D |
| <code>Vpath</code> | solution path of D |
| <code>Rpath</code> | path of estimated rank |
| <code>icpath</code> | path of information criteria |

| | |
|---------|--|
| lam.id | ids of selected lambda for GIC, BIC, AIC and GCV |
| p.index | ids of predictors which passed screening |
| q.index | ids of responses which passed screening |
| lamA | tuning sequence for A |
| lamB | tuning sequence for B |
| lamD | tuning sequence for D |
| U | estimated left singular matrix that is orthogonal (factor weights) |
| V | estimated right singular matrix that is orthogonal (factor loadings) |
| D | estimated singular values |
| rank | estimated rank |

References

Uematsu, Y., Fan, Y., Chen, K., Lv, J., & Lin, W. (2019). SOFAR: large-scale association network learning. *IEEE Transactions on Information Theory*, 65(8), 4924–4939.

Examples

```
## Not run:
library(rrpack)
## Simulate data from a sparse factor regression model
p <- 100; q <- 50; n <- 100; nrank <- 3
mydata <- rrr.sim1(n, p, q, nrank, s2n = 1,
                  sigma = NULL, rho_X = 0.5, rho_E = 0.3)

Y <- mydata$Y
X <- mydata$X

fit1 <- sofar(Y, X, ic.type = "GIC", nrank = nrank + 2,
             control = list(methodA = "adlasso", methodB = "adlasso"))
summary(fit1)
plot(fit1)

fit1$U
crossprod(fit1$U) #check orthogonality
fit1$V
crossprod(fit1$V) #check orthogonality

## End(Not run)
```

srrr

Row-sparse reduced-rank regression

Description

Row-sparse reduced-rank regression for a prespecified rank; produce a solution path for selecting predictors

Usage

```
srrr(
  Y,
  X,
  nrank = 2,
  method = c("glasso", "adglasso"),
  ic.type = c("BIC", "BICP", "AIC", "GCV", "GIC"),
  A0 = NULL,
  V0 = NULL,
  modstr = list(),
  control = list(),
  screening = FALSE
)
```

Arguments

| | |
|-----------|---|
| Y | response matrix |
| X | covariate matrix |
| nrank | prespecified rank |
| method | group lasso or adaptive group lasso |
| ic.type | information criterion |
| A0 | initial value |
| V0 | initial value |
| modstr | a list of model parameters controlling the model fitting |
| control | a list of parameters for controlling the fitting process |
| screening | If TRUE, marginal screening via glm is performed before srrr fitting. |

Details

Model parameters controlling the model fitting can be specified through argument `modstr`. The available elements include

- `lamA`: tuning parameter sequence.
- `nlam`: number of tuning parameters; no effect if `lamA` is specified.
- `minLambda`: minimum lambda value, no effect if `lamA` is specified.
- `maxLambda`: maximum lambda value, no effect if `lamA` is specified.
- `WA`: adaptive weights. If `NULL`, the weights are constructed from `RRR`.
- `wgamma`: power parameter for constructing adaptive weights.

Similarly, the computational parameters controlling optimization can be specified through argument `control`. The available elements include

- `epsilon`: epsilon convergence tolerance.
- `maxit`: maximum number of iterations.
- `inner.eps`: used in inner loop.
- `inner.maxit`: used in inner loop.

Value

A list of fitting results

References

Chen, L. and Huang, J. Z. (2012) Sparse reduced-rank regression for simultaneous dimension reduction and variable selection. *Journal of the American Statistical Association*. 107:500, 1533–1545.

Examples

```
library(rrpack)
p <- 100; n <- 100; nrank <- 3
mydata <- rrr.sim2(n, p, p0 = 10, q = 50, q0 = 10, nrank = 3,
                 s2n = 1, sigma = NULL, rho_X = 0.5, rho_E = 0)
fit1 <- with(mydata, srrr(Y, X, nrank = 3))
summary(fit1)
coef(fit1)
plot(fit1)
```

summary

Summarize rrpacK Objects

Description

S3 methods summarizing objects generated by rrpacK.

Usage

```
## S3 method for class 'mrrr'
summary(object, ...)

## S3 method for class 'cv.mrrr'
summary(object, ...)

## S3 method for class 'r4'
summary(object, ...)

## S3 method for class 'rrr'
summary(object, ...)

## S3 method for class 'cv.rrr'
summary(object, ...)

## S3 method for class 'sofar'
summary(object, ...)

## S3 method for class 'cv.sofar'
```



```
summary(object, ...)  
  
## S3 method for class 'srrr'  
summary(object, ...)  
  
## S3 method for class 'cv.srrr'  
summary(object, ...)  
  
## S3 method for class 'rssvd'  
summary(object, ...)
```

Arguments

| | |
|--------|-----------------------------------|
| object | Object generated from rrpck. |
| ... | Other arguments for future usage. |

Index

coef.cv.mrrr (rrpack-coef), 14
coef.cv.rrr (rrpack-coef), 14
coef.mrrr (rrpack-coef), 14
coef.r4 (rrpack-coef), 14
coef.rrr (rrpack-coef), 14
coef.rssvd (rrpack-coef), 14
coef.sofar (rrpack-coef), 14
coef.srrr (rrpack-coef), 14
cv.mrrr, 2
cv.rrr, 4
cv.sofar, 5
cv.srrr, 7

mrrr, 8

plot, 10

r4, 12
rrpack-coef, 14
rrr, 15
rrr.cookD, 17
rrr.fit, 18
rrr.leverage, 19
rrr.sim1, 20
rrr.sim2, 21
rrr.sim3, 22
rrr.sim4, 23
rrr.sim5, 24
rrs.fit, 25
rssvd, 26

sofar, 28
srrr, 30
summary, 32