

Package ‘cops’

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Title Cluster Optimized Proximity Scaling

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Description Cluster optimized proximity scaling (COPS) refers to multidimensional scaling (MDS) methods that aim at pronouncing the clustered appearance of the configuration (Rusch, Mair & Hornik, 2021, <doi:10.1080/10618600.2020.1869027>). They achieve this by transforming proximities/distances with power functions and augment the fitting criterion with a clusteredness index, the OP-TICS Cordillera (Rusch, Hornik & Mair, 2018, <doi:10.1080/10618600.2017.1349664>). There are two variants: One for finding the configuration directly (COPS-C) for ratio, power, interval and non-metric MDS (Borg & Groenen, 2005, ISBN:978-0-387-28981-6), and one for using the augmented fitting criterion to find optimal parameters (P-COPS). The package contains various functions, wrappers, methods and classes for fitting, plotting and displaying different MDS models in a COPS framework like ratio, interval and non-metric MDS for COPS-C and P-COPS with Torgerson scaling (Torgerson, 1958, ISBN:978-0471879459), scaling by majorizing a complex function (SMA-COF; de Leeuw, 1977, <<https://escholarship.org/uc/item/4ps3b5mj>>), Sammon mapping (Sammon, 1969, <doi:10.1109/T-C.1969.222678>), elastic scaling (McGee, 1966, <doi:10.1111/j.2044-8317.1966.tb00367.x>), s-stress (Takane, Young & de Leeuw, 1977, <doi:10.1007/BF02293745>), r-stress (de Leeuw, Groenen & Mair, 2016, <<https://rpubs.com/deleeuw/142619>>), power-stress (Buja & Swayne, 2002 <doi:10.1007/s00357-001-0031-0>), restricted power stress, approximated power stress, power elastic scaling, power Sammon mapping (Rusch, Mair & Hornik, 2021, <doi:10.1080/10618600.2020.1869027>). All of these models can also solely be fit as MDS with power transformations. The package further contains a function for pattern search optimization, the “Adaptive Luus-Jakola Algorithm” (Rusch, Mair & Hornik, 2021, <doi:10.1080/10618600.2020.1869027>).

Depends R (>= 3.1.2), cordillera (>= 0.7-2), smacof (>= 1.10-4)

Imports MASS, minqa, pso, scatterplot3d, NlcOptim, Rsolnp, dfoptim, subplex, cmaes, crs, nloptr, rgenoud, GenSA

Enhances stats

Suggests testthat, R.rsp, rmarkdown

VignetteBuilder R.rsp

License GPL-2 | GPL-3

LazyData true

URL <http://r-forge.r-project.org/projects/stops/>

RoxygenNote 7.1.1

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R topics documented:

BankingCrisesDistances	3
cmdscale	3
conf_adjust	4
cops	5
copstress	8
copstressMin	9
cop_apstress	12
cop_cmdscale	14
cop_elastic	15
cop_powerelastic	17
cop_powermds	19
cop_powersammon	20
cop_powerstress	22
cop_rpowerstress	23
cop_rstress	25
cop_sammon	27
cop_sammon2	28
cop_smacofSphere	30
cop_smacofSym	31
cop_sstress	33
doubleCenter	35
enorm	35
ljoptim	36
mkBmat	37
mkPower	38
pcops	38
pdist	41
plot.cops	42
plot.pcops	43
plot.smacofP	44
plot3dstatic	46

BankingCrisesDistances 3

plot3dstatic.cmdscale	46
powerStressFast	47
powerStressMin	49
procruster	50
sammon	51
scale_adjust	51
secularEq	52
sqdist	52
torgerson	53

Index 54

BankingCrisesDistances
Banking Crises Distances

Description

Matrix of Jaccard distances between 70 countries (Hungary and Greece were combined to be the same observation) based on their binary time series of having had a banking crises in a year from 1800 to 2010 or not. See `data(bankingCrises)` in package `Ecdat` for more info. The last column is Reinhart & Rogoffs classification as a low (3), middle- (2) or high-income country (1).

Format

A 69 x 70 matrix.

Source

`data(bankingCrises)` in library(`Ecdat`)

`cmdscale` *Wrapper to cmdscale for S3 class*

Description

Wrapper to `cmdscale` for S3 class

Usage

`cmdscale(d, k = 2, eig = TRUE, ...)`

Arguments

d	a distance structure such as that returned by 'dist' or a full symmetric matrix containing the dissimilarities
k	the maximum dimension of the space which the data are to be represented in
eig	indicates whether eigenvalues should be returned.
...	additional parameters passed to cmdscale. See cmdscale

Value

See [cmdscale](#). This wrapper only adds an extra slot to the list with the call, adds column labels to the \$points and assigns S3 class 'cmdscale'.

Examples

```
dis<-as.matrix(smacof::kinshipdelta)
res<-cmdscale(dis)
```

conf_adjust	<i>conf_adjust: a function to procrustes adjust two matrices</i>
-------------	--

Description

conf_adjust: a function to procrustes adjust two matrices

Usage

```
conf_adjust(conf1, conf2, verbose = FALSE, eps = 1e-12, itmax = 100)
```

Arguments

conf1	reference configuration, a numeric matrix
conf2	another configuration, a numeric matrix
verbose	should adjustment be output; default to FALSE
eps	numerical accuracy
itmax	maximum number of iterations

Value

a list with ref.conf being the reference configuration, other.conf the adjusted configuration and comparison.conf the comparison configuration

Description

About the package `cops`: Cluster optimized proximity scaling (COPS) refers to multidimensional scaling methods that aim at pronouncing the clustered appearance of the configuration. They achieve this by transforming proximities/distances with power functions and augment the fitting criterion with a clusteredness index, the OPTICS Cordillera (Rusch, Hornik & Mair 2018). There are two variants: One for finding the configuration directly for given parameters (COPS-C), and one for using the augmented fitting criterion to find optimal parameters for the power transformations (P-COPS). The package contains various functions, wrappers, methods and classes for fitting, plotting and displaying different MDS models in a COPS framework like Torgerson scaling, SMA-COF, Sammon mapping, elastic scaling, symmetric SMACOF, spherical SMACOF, `sstress`, `rstress`, `powermds`, power elastic scaling, power sammon mapping, `powerstress`. All of these models can also solely be fit as MDS with power transformations. The package further contains functions for optimization (Adaptive LJ Algorithmus).

About the function `cops`: The high level function allows for minimizing `copstress` for a clustered MDS configuration. Allows to choose COPS-C (finding a configuration from `copstress` with cordillera penalty) and profile COPS (finding hyperparameters for MDS models with power transformations). It is a wrapper for `copstressMin` and `pcops`.

Usage

```
cops(
  dis,
  variant = c("1", "2", "Variant1", "Variant2", "v1", "v2", "COPS-C", "P-COPS",
    "configuration-c", "profile", "copstress-c", "p-copstress", "COPS-P", "copstress-p",
    "cops-c", "p-cops", "copsc", "pcops"),
  ...
)
```

Arguments

<code>dis</code>	a dissimilarity matrix or a <code>dist</code> object
<code>variant</code>	a character string specifying which variant of COPS to fit. Allowed is any of the following "1","2","Variant1","Variant2","v1","v2","COPS-C","P-COPS","configuration-c","profile","copstress-c","p-copstress". Defaults to "COPS-C".
<code>...</code>	arguments to be passed to <code>copstressMin</code> (for Variant 1) or <code>pcops</code> (for Variant 2).

Details

The `cops` package provides five categories of important functions:

Models & Algorithms:

- `cops()` ... high level interface to fit COPS models as described in Rusch et al. (2021). By setting `cordweight` to zero they can also be used to fit metric MDS for many different models, see below.
- `copstressMin()`... The workhorse for fitting a COPS-C model. Can also be called directly.
- `pcops()`... The workhorse for fitting a P-COPS model. Can also be called directly.
- `powerStressMin()`... a workhorse for fitting s-stress, r-stress (de Leeuw, 2014), p-stress (e.g., Rusch et al., 2021), Sammon mapping with power transformations (`powersammon`) and elastic scaling with power transformation (`powerelastic`). They can conveniently also be fitted via the `cops` functions and setting `stressweight=1` and `cordweight` or by the dedicated functions starting with `cops_XXX` where `XXX` is the method and setting `stressweight=1` and `cordweight=0`. It uses the nested majorization algorithm for r-stress of De Leeuw (2014).

Optimization functions:

- `ljoptim()` ... An (adaptive) version of the Luus-Jakola random search

Wrappers and convenience functions:

- `conf_adjust()`: procrustes adjustment of configurations
- `cmdscale()`, `sammon()`: wrappers that return S3 objects to be used with `cops`
- `copstress()` ... a function to calculate `copstress` (Rusch et al., 2021)
- `cop_smacofSym()`, `cop_sammon()`, `cop_cmdscale()`, `cop_rstress()`, `cop_powerstress()`, `cop_smacofSphere()`, `cop_sammon2()`, `cop_elastic()`, `cop_sstress()`, `cop_powerelastic()`, `cop_powersammon()`: cop versions of these MDS models.

Methods: For most of the objects returned by the high-level functions S3 classes and methods for standard generics were implemented, including `print`, `summary`, `plot`, `plot3dstatic`.

References:

- Rusch, T., Hornik, K. & Mair, P. (2018) Assessing and quantifying clusteredness: The OPTICS Cordillera. *Journal of Computational and Graphical Statistics*, 27 (1), 220-233. doi: [10.1080/10618600.2017.1349664](https://doi.org/10.1080/10618600.2017.1349664)
- Rusch, T., Mair, P. & Hornik, K. (2021) Cluster optimized proximity scaling. *Journal of Computational and Graphical Statistics*. doi: [10.1080/10618600.2020.1869027](https://doi.org/10.1080/10618600.2020.1869027)

Authors: Thomas Rusch, Jan de Leeuw, Patrick Mair

Maintainer: Thomas Rusch

Value

For COPS-C Variant 1 see [copstressMin](#), for P-COPS Variant 2 see [pcops](#)

Examples

```
data(BankingCrisesDistances)
```

```
# shorthand function for COPS-C (finding configuration with copstress)
```

```

res<-cops(BankingCrisesDistances[,1:69],variant="COPS-C",
          stressweight=0.98,cordweight=0.02,itmax=1000)
# Note: itmax is very small here for illustration; will give a non-convergence
# warning of the optimizer which disappears at itmax=275000

res
summary(res)
plot(res)
plot(res,"reachplot")
plot(res,"transplot")
plot(res,"Shepard")
#shorthand function for P-COPS (hyperparameter search for powerstress)
res<-cops(BankingCrisesDistances[,1:69],variant="P-COPS")
res
summary(res)
plot(res)
plot(res,"reachplot")
plot(res,"transplot")
plot(res,"Shepard")

dis<-as.matrix(smacof::kinshipdelta)

#COPS-C with equal weight to stress and cordillera
res1<-cops(dis,variant="COPS-C",stressweight=0.5,cordweight=0.5,
           minpts=2,itmax=1000) #use higher itmax in real
res1
summary(res1)
plot(res1)
plot(res1,"reachplot")

#s-stress type copstress (i.e. kappa=2, lambda=2)
res3<-cops(dis,variant="COPS-C",kappa=2,lambda=2,stressweight=0.5,cordweight=0.5)
res3
summary(res3)
plot(res3)

# power-stress type profile copstress
# search for optimal kappa and lambda between
# kappa=0.5,lambda=0.5 and kappa=2,lambda=5
# nu is fixed on -1
ws<-1/dis
diag(ws)<-1
res5<-cops(dis,variant="P-COPS",loss="powerstress",
           theta=c(1.4,3,-1), lower=c(1,0.5,-1),upper=c(3,5,-1),
           weightmat=ws, stressweight=0.9,cordweight=0.1)
res5
summary(res5)
plot(res5)

```

 copstress

Calculates copstress for given MDS object

Description

Calculates copstress for given MDS object

Usage

```
copstress(
  obj,
  stressweight = 1,
  cordweight = 5,
  q = 1,
  minpts = 2,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  normed = TRUE,
  scale = c("std", "sd", "proc", "none"),
  init,
  ...
)
```

Arguments

obj	MDS object (supported are sammon, cmdscale, smacof, rstress, powermds)
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to 2
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value that prints information on the fitting process; >2 is very verbose (copstress level), >3 is extremely (up to MDS optimization level)
normed	should the cordillera be normed; defaults to TRUE
scale	should the configuration be scale adjusted.
init	a reference configuration when doing procrustes adjustment
...	additional arguments to be passed to the cordillera function

Value

A list with the components

- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- cordillera: the cordillera object

copstressMin

Fitting a COPS-C Model (COPS Variant 1).

Description

Minimizing Copstress to obtain a clustered MDS configuration with given hyperparameters theta.

Usage

```
copstressMin(
  delta,
  kappa = 1,
  lambda = 1,
  nu = 1,
  theta = c(kappa, lambda, nu),
  type = c("ratio", "interval", "ordinal"),
  ties = "primary",
  weightmat = 1 - diag(nrow(delta)),
  ndim = 2,
  init = NULL,
  stressweight = 0.975,
  cordweight = 0.025,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  dmax = NULL,
  rang,
  optimmethod = c("NelderMead", "Newuoa", "BFGS", "SANN", "hjk", "solnl", "solnp",
    "subplex", "snomadr", "hjk-Newuoa", "hjk-BFGS", "BFGS-hjk", "Newuoa-hjk", "cmaes",
    "direct", "direct-Newuoa", "direct-BFGS", "genoud", "gensa"),
  verbose = 0,
  scale = c("sd", "rmsq", "std", "proc", "none"),
  normed = TRUE,
  accuracy = 1e-07,
  itmax = 5000,
  stresstype = c("stress-1", "stress"),
  ...
)
```

Arguments

delta	numeric matrix or dist object of a matrix of proximities
kappa	power transformation for fitted distances
lambda	power transformation for proximities
nu	power transformation for weights
theta	the theta vector of powers; the first is kappa (for the fitted distances if it exists), the second lambda (for the observed proximities if it exist), the third is nu (for the weights if it exists) . If less than three elements are is given as argument, it will be recycled. Defaults to 1 1 1. Will override any kappa, lmbda, nu parameters if they are given and do not match
type	what type of MDS to fit. Currently one of "ratio", "interval" or "ordinal". Default is "ratio".
ties	the handling of ties for ordinal (nonmetric) MDS. Possible are "primary" (default), "secondary" or "tertiary".
weightmat	(optional) a matrix of nonnegative weights; defaults to 1 for all off diagonals
ndim	number of dimensions of the target space
init	(optional) initial configuration
stressweight	weight to be used for the fit measure; defaults to 0.975
cordweight	weight to be used for the cordillera; defaults to 0.025
q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
dmax	The winsorization limit of reachability distances in the OPTICS Cordillera. If supplied, it should be either a numeric value that matches max(rang) or NULL; if NULL it is found as 1.5 times (for kappa >1) or 1 times (for kappa <=1) the maximum reachbilty value of the power torgerson model with the same lambda. If dmax and rang are supplied and dmax is not max(rang), a warning is given and rang takes precedence.
rang	range of the reachabilities to be considered. If missing it is found from the initial configuration by taking 0 as the lower boundary and dmax (see above) as upper boundary. See also cordillera
optimmethod	What optimizer to use? Choose one string of 'Newuoa' (from package minqa), 'NelderMead', 'hjk' (Hooke-Jeeves algorithm from dfoptim), 'solnl' (from nl-cOptim), 'solnp' (from Rsolnp), 'subplex' (from subplex), 'SANN' (simulated annealing), 'BFGS', 'snomadr' (from crs), 'genoud' (from rgenoud), 'gensa' (from GenSA), 'cmaes' (from cmaes) and 'direct' (from nloptr). See the according R packages for details on these solvers. There are also combinations that proved to work well good, like 'hjk-Newuoa', 'hjk-BFGS', 'BFGS-hjk', 'Newuoa-hjk', 'direct-Newuoa' and 'direct-BFGS'. Usually hjk, BFGS, newuoa, subplex and solnl work rather well in an acceptable time frame (depending on the smoothness of copstress). Default is 'hjk-Newuoa'.
verbose	numeric value hat prints information on the fitting process; >2 is very verbose

scale	Allows to scale the configuration for the OC (the scaled configuration is also returned as \$conf). One of "none" (so no scaling), "sd" (configuration divided by the highest standard deviation of the columns), "std" (standardize all columns !NOTE: This does not preserve the relative distances of the optimal config), "proc" (procrustes adjustment to the initial fit) and "rmsq" (configuration divided by the maximum root mean square of the columns). Default is "sd".
normed	should the cordillera be normed; defaults to TRUE
accuracy	numerical accuracy, defaults to 1e-7
itmax	maximum number of iterations. Defaults to 5000. If itmax is (too) small, some optimizers will print warnings. For example, for optimizers using NEWUOA, an iteration number of $10 * \text{length}(\text{par})^2$ is recommended. The number of parameters to optimize over for the COPS problem is number of objects * target space dimensions and can grow large very quickly, so being able to live with these warnings is probably a good idea.
stresstype	which stress to use in the copstress. Defaults to stress-1. If anything else is set, explicitly normed stress which is $(\text{stress}-1)^2$. Using stress-1 puts more weight on MDS fit.
...	additional arguments to be passed to the optimization procedure

Value

A list with the components

- delta: the original transformed dissimilarities
- obsdiss: the explicitly normed transformed dissimilarities (which are approximated by the fit)
- confdist: the fitted distances
- conf: the configuration to which the scaling of argument scale was applied
- confo: the unscaled but explicitly normed configuration returned from the fitting procedure. Scaling applied to confo gives conf.
- par, pars : the theta vector of powers transformations (κ, λ, ν)
- niter: number of iterations of the optimizer.
- stress: the square root of explicitly normalized stress (calculated for confo).
- spp: stress per point
- ndim: number of dimensions
- model: Fitted model name with optimizer
- call: the call
- nobj: the number of objects
- type, loss, losstype: stresstype
- stress.m: The stress used for copstress. If stresstype="stress-1" this is like \$stress else it is stress^2
- stress.en: another ways to calculate the stress
- deltaorig: the original untransformed dissimilarities

- copstress: the copstress loss value
- resmat: the matrix of residuals
- weightmat: the matrix of untransformed weights
- OC: the (normed) OPTICS Cordillera object (calculated for scaled conf)
- OCv: the (normed) OPTICS Cordillera value alone (calculated for scaled conf)
- optim: the object returned from the optimization procedure
- stressweight, cordweight: the weights of the stress and OC respectively (v_1 and v_2)
- optimmethod: The solver used
- type: the type of MDS fitted

Examples

```
dis<-as.matrix(smacof::kinshipdelta)

#Copstress with equal weight to stress and cordillera
res1<-copstressMin(dis, stressweight=0.5, cordweight=0.5,
                  itmax=1000) #use higher itmax about 10000
res1
summary(res1)
plot(res1) #super clustered
```

cop_apstress

PCOPS versions of approximated power stress models. This uses an approximation and makes use of smacof.

Description

PCOPS versions of approximated power stress models. This uses an approximation and makes use of smacof.

Usage

```
cop_apstress(
  dis,
  theta = c(1, 1, 1),
  ndim = 2,
  weightmat = NULL,
  init = NULL,
  itmaxi = 1000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
```

```

    epsilon = 10,
    rang = NULL,
    verbose = 0,
    normed = TRUE,
    scale = "sd",
    stresstype = "default"
)

```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; this is either a scalar of the tau and epsilon transformation for the observed proximities, or a vector where the first is the kappa argument for the fitted distances (here internally fixed to 1) and the second the tau argument and the third the epsilon argument. Defaults to 1 1 1
ndim	number of dimensions of the target space
weightmat	(optional) a binary matrix of nonnegative weights
init	(optional) initial configuration
itmaxi	number of iterations. default is 1000.
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value that prints information on the fitting process; >2 is extremely verbose
normed	should the cordillera be normed; defaults to TRUE
scale	should the configuration be scale adjusted
stresstype	which stress to report. Only takes smacofs default stress currently.

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value

- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure (which has all smacofB elements and some more)
- cordillera: the cordillera object

 cop_cmdscale

PCOPS version of strain

Description

PCOPS version of strain

Usage

```
cop_cmdscale(
  dis,
  theta = c(1, 1, 1),
  weightmat = NULL,
  ndim = 2,
  init = NULL,
  itmaxi = 1000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  scale = "sd",
  normed = TRUE,
  stresstype = "default"
)
```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; this is either a scalar of the lambda transformation for the observed proximities, or a vector where the first is the kappa argument for the fitted distances (here internally fixed to 1) and the second and third the lambda and the nu argument (the latter is fixed to 1). Defaults to 1 1 1
weightmat	(optional) a matrix of nonnegative weights
ndim	number of dimensions of the target space
init	(optional) initial configuration
itmaxi	number of iterations. No effect here.

...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the corrdillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value hat prints information on the fitting process; >2 is extremely verbose
scale	should the configuration be scale adjusted
normed	should the cordillera be normed; defaults to TRUE
stresstype	which stress to report. Only takes cmdscases default stress currently.

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_elastic

PCOPS versions of elastic scaling models (via smacofSym)

Description

PCOPS versions of elastic scaling models (via smacofSym)

Usage

```

cop_elastic(
  dis,
  theta = c(1, 1, -2),
  ndim = 2,
  weightmat = 1,
  init = NULL,
  itmaxi = 1000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  normed = TRUE,
  scale = "sd",
  stresstype = "default"
)

```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; this is either a scalar of the lambda transformation for the observed proximities, or a vector where the first is the kappa argument for the fitted distances (here internally fixed to 1) and the second the lambda argument and the third the nu argument (here internally fixed to -2). Defaults to 1 1 -2
ndim	number of dimensions of the target space
weightmat	(optional) a matrix of nonnegative weights (NOT the elscal weights)
init	(optional) initial configuration
itmaxi	number of iterations. default is 1000.
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the corrdillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value hat prints information on the fitting process; >2 is extremely verbose

normed	should the cordillera be normed; defaults to TRUE
scale	should the configuration be scale adjusted
stresstype	which stress to report. Only takes smacofs default stress currently.

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (κ , λ)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_powerelastic *PCOPS version of elastic scaling with powers*

Description

PCOPS version of elastic scaling with powers

Usage

```
cop_powerelastic(
  dis,
  theta = c(1, 1, -2),
  weightmat = 1 - diag(nrow(dis)),
  init = NULL,
  ndim = 2,
  itmaxi = 10000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  scale = "sd",
  normed = TRUE,
  stresstype = c("default", "stress1", "rawstress", "normstress", "enormstress",
    "enormstress1")
)
```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; the first is kappa (for the fitted distances), the second lambda (for the observed proximities) and nu as the third (fixed to -2). If a scalar for the free parameters is given it is recycled. Defaults to 1 1 -2.
weightmat	(optional) a matrix of nonnegative weights
init	(optional) initial configuration
ndim	number of dimensions of the target space
itmaxi	number of iterations. default is 10000.
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value that prints information on the fitting process; >2 is extremely verbose
scale	should the configuration be scale adjusted
normed	should the cordillera be normed; defaults to TRUE
stresstype	which stress to report? Defaults to explicitly normed stress

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_powermds

PCOPS version of powermds

Description

PCOPS version of powermds

Usage

```
cop_powermds(
  dis,
  theta = c(1, 1, 1),
  weightmat = 1 - diag(nrow(dis)),
  init = NULL,
  ndim = 2,
  itmaxi = itmaxi,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  scale = "sd",
  normed = TRUE,
  stresstype = c("default", "stress1", "rawstress", "normstress", "enormstress",
    "enormstress1")
)
```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; the first is kappa (for the fitted distances), the second lambda (for the observed proximities), nu is fixed to 1. If a scalar is given it is recycled. Defaults to 1 1 1.
weightmat	(optional) a matrix of nonnegative weights
init	(optional) initial configuration
ndim	number of dimensions of the target space
itmaxi	number of iterations. default is 10000.
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the cordillera; defaults to 1

minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value that prints information on the fitting process; >2 is extremely verbose
scale	should the configuration be scale adjusted
normed	should the cordillera be normed; defaults to TRUE
stresstype	which stress to report? Defaults to whatever whim is my default (currently explicitly normed stress)

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_powersammon

PCOPS version of sammon with powers

Description

PCOPS version of sammon with powers

Usage

```
cop_powersammon(
  dis,
  theta = c(1, 1, -1),
  weightmat = 1 - diag(nrow(dis)),
  init = NULL,
  ndim = 2,
  itmaxi = 10000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
```

```

q = 1,
minpts = ndim + 1,
epsilon = 10,
rang = NULL,
verbose = 0,
scale = "sd",
normed = TRUE,
stresstype = c("default", "stress1", "rawstress", "normstress", "enormstress",
  "enormstress1")
)

```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; the first is kappa (for the fitted distances), the second lambda (for the observed proximities), the third nu (fixed to -1). If a scalar is given it is recycled for the free parameters. Defaults to 1 1 -1.
weightmat	(optional) a matrix of nonnegative weights
init	(optional) initial configuration
ndim	number of dimensions of the target space
itmaxi	number of iterations. default is 10000.
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value hat prints information on the fitting process; >2 is extremely verbose
scale	should the configuration be scale adjusted
normed	should the cordillera be normed; defaults to TRUE
stresstype	which stress to report? Defaults to explicitly normed stress

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value

- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

 cop_powerstress

COPS version of powerstress

Description

COPS version of powerstress

Usage

```
cop_powerstress(
  dis,
  theta = c(1, 1, 1),
  weightmat = 1 - diag(nrow(dis)),
  init = NULL,
  ndim = 2,
  itmaxi = 10000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  scale = "sd",
  normed = TRUE,
  stresstype = c("default", "stress1", "rawstress", "normstress", "enormstress",
    "enormstress1")
)
```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; the first is kappa (for the fitted distances), the second lambda (for the observed proximities), the third nu (for the weights). If a scalar is given it is recycled. Defaults to 1 1 1.
weightmat	(optional) a matrix of nonnegative weights
init	(optional) initial configuration
ndim	number of dimensions of the target space
itmaxi	number of iterations. default is 10000.

...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value that prints information on the fitting process; >2 is extremely verbose
scale	should the configuration be scale adjusted
normed	should the cordillera be normed; defaults to TRUE
stresstype	which stress to report? Defaults to explicitly normed stress

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_rpowerstress *COPS version of restricted powerstress*

Description

COPS version of restricted powerstress

Usage

```

cop_rpowerstress(
  dis,
  theta = c(1, 1, 1),
  weightmat = 1 - diag(nrow(dis)),
  init = NULL,
  ndim = 2,
  itmaxi = 10000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  scale = "sd",
  normed = TRUE,
  stresstype = c("default", "stress1", "rawstress", "normstress", "enormstress",
    "enormstress1")
)

```

Arguments

<code>dis</code>	numeric matrix or dist object of a matrix of proximities
<code>theta</code>	the theta vector of powers; the first two arguments are for kappa and lambda and must be equal (for the dissimilarities and observed proximities), the third nu (for the weights). If a scalar is given it is recycled, if a vector of length is given, it gets expanded to <code>c(theta[1],theta[1],theta[2])</code> . Defaults to 1 1 1.
<code>weightmat</code>	(optional) a matrix of nonnegative weights
<code>init</code>	(optional) initial configuration
<code>ndim</code>	number of dimensions of the target space
<code>itmaxi</code>	number of iterations. default is 10000.
<code>...</code>	additional arguments to be passed to the fitting procedure
<code>stressweight</code>	weight to be used for the fit measure; defaults to 1
<code>cordweight</code>	weight to be used for the cordillera; defaults to 0.5
<code>q</code>	the norm of the cordillera; defaults to 1
<code>minpts</code>	the minimum points to make up a cluster in OPTICS; defaults to <code>ndim+1</code>
<code>epsilon</code>	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
<code>rang</code>	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
<code>verbose</code>	numeric value that prints information on the fitting process; >2 is extremely verbose

scale	should the configuration be scale adjusted
normed	should the cordillera be normed; defaults to TRUE
stresstype	which stress to report? Defaults to explicitly normed stress

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (κ , λ)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_rstress	<i>PCOPS version of rstress</i>
-------------	---------------------------------

Description

PCOPS version of rstress

Usage

```
cop_rstress(
  dis,
  theta = c(1, 1, 1),
  weightmat = 1 - diag(nrow(dis)),
  init = NULL,
  ndim = 2,
  itmaxi = 10000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  scale = "sd",
  normed = TRUE,
  stresstype = c("default", "stress1", "rawstress", "normstress", "enormstress",
    "enormstress1")
)
```

Arguments

<code>dis</code>	numeric matrix or dist object of a matrix of proximities
<code>theta</code>	the theta vector of powers; this is either a scalar of the kappa transformation for the fitted distances proximities, or a vector where the first is the kappa argument for the fitted distances and the second the lambda argument, the third the nu argument (here internally fixed to 1). Defaults to 1 1 1
<code>weightmat</code>	(optional) a matrix of nonnegative weights
<code>init</code>	(optional) initial configuration
<code>ndim</code>	number of dimensions of the target space
<code>itmaxi</code>	number of iterations. default is 10000.
<code>...</code>	additional arguments to be passed to the fitting procedure
<code>stressweight</code>	weight to be used for the fit measure; defaults to 1
<code>cordweight</code>	weight to be used for the cordillera; defaults to 0.5
<code>q</code>	the norm of the cordillera; defaults to 1
<code>minpts</code>	the minimum points to make up a cluster in OPTICS; defaults to $ndim+1$
<code>epsilon</code>	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
<code>rang</code>	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
<code>verbose</code>	numeric value that prints information on the fitting process; >2 is extremely verbose
<code>scale</code>	should the configuration be scale adjusted
<code>normed</code>	should the cordillera be normed; defaults to TRUE
<code>stresstype</code>	which stress to report? Defaults to explicitly normed stress

Value

A list with the components

- `stress`: the stress
- `stress.m`: default normalized stress
- `copstress`: the weighted loss value
- `OC`: the Optics cordillera value
- `parameters`: the parameters used for fitting (kappa, lambda)
- `fit`: the returned object of the fitting procedure
- `cordillera`: the cordillera object

 cop_sammon

PCOPS version of sammon mapping

Description

PCOPS version of sammon mapping

Usage

```
cop_sammon(
  dis,
  theta = c(1, 1, -1),
  ndim = 2,
  init = NULL,
  weightmat = NULL,
  itmaxi = 100,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  scale = "sd",
  normed = TRUE,
  stresstype = "default"
)
```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; this is either a scalar of the lambda transformation for the observed proximities, or a vector where the first is the kappa argument for the fitted distances (here internally fixed to 1) and the second the lambda argument and the third the nu argument (here internally fixed to -1). Defaults to 1 1 -1
ndim	number of dimensions of the target space
init	(optional) initial configuration
weightmat	(optional) a matrix of nonnegative weights
itmaxi	number of iterations. default is 1000.
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5

q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value that prints information on the fitting process; >2 is extremely verbose
scale	should the configuration be scale adjusted
normed	should the cordillera be normed; defaults to TRUE
stresstype	which stress to report. Only takes smacofs default stress currently.

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_sammon2

COPS versions of Sammon mapping models (via smacofSym)

Description

COPS versions of Sammon mapping models (via smacofSym)

Usage

```
cop_sammon2(
  dis,
  theta = c(1, 1, -1),
  ndim = 2,
  weightmat = NULL,
  init = NULL,
  itmaxi = 1000,
  ...,
  stressweight = 1,
```

```

cordweight = 0.5,
q = 1,
minpts = ndim + 1,
epsilon = 10,
rang = NULL,
verbose = 0,
normed = TRUE,
scale = "sd",
stresstype = "default"
)

```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; this is either a scalar of the lambda transformation for the observed proximities, or a vector where the first is the kappa argument for the fitted distances (here internally fixed to 1) and the second the lambda argument, the third the nu argument (fixed to -1). Defaults to 1 1 -1.
ndim	number of dimensions of the target space
weightmat	(optional) a matrix of nonnegative weights (NOT the sammon weights)
init	(optional) initial configuration
itmaxi	number of iterations. default is 1000.
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value that prints information on the fitting process; >2 is extremely verbose
normed	should the cordillera be normed; defaults to TRUE
scale	should the configuration be scale adjusted
stresstype	which stress to report. Only takes smacofs default stress currently.

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress

- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_smacofSphere *PCOPS versions of smacofSphere models*

Description

PCOPS versions of smacofSphere models

Usage

```
cop_smacofSphere(
  dis,
  theta = c(1, 1, 1),
  ndim = 2,
  weightmat = NULL,
  init = NULL,
  itmaxi = 1000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  normed = TRUE,
  scale = "sd",
  stresstype = "default"
)
```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; this is either a scalar of the lambda transformation for the observed proximities, or a vector where the first is the kappa argument for the fitted distances (here internally fixed to 1) and the second the lambda argument and the third the nu argument. Defaults to 1 1 1
ndim	number of dimensions of the target space
weightmat	(optional) a matrix of nonnegative weights
init	(optional) initial configuration

itmaxi	number of iterations. default is 1000.
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the corrdillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value hat prints information on the fitting process; >2 is extremely verbose
normed	should the cordillera be normed; defaults to TRUE
scale	should the configuration be scale adjusted
stresstype	which stress to report. Only takes smacofs default stress currently.

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

cop_smacofSym

PCOPS versions of smacofSym models

Description

PCOPS versions of smacofSym models

Usage

```

cop_smacofSym(
  dis,
  theta = c(1, 1, 1),
  ndim = 2,
  weightmat = NULL,
  init = NULL,
  itmaxi = 1000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  normed = TRUE,
  scale = "sd",
  stresstype = "default"
)

```

Arguments

dis	numeric matrix or dist object of a matrix of proximities
theta	the theta vector of powers; this is either a scalar of the lambda transformation for the observed proximities, or a vector where the first is the kappa argument for the fitted distances (here internally fixed to 1) and the second the lambda argument and the third the nu argument (here internally fixed to 1). Defaults to 1 1 1
ndim	number of dimensions of the target space
weightmat	(optional) a matrix of nonnegative weights
init	(optional) initial configuration
itmaxi	number of iterations. default is 1000
...	additional arguments to be passed to the fitting procedure
stressweight	weight to be used for the fit measure; defaults to 1
cordweight	weight to be used for the cordillera; defaults to 0.5
q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
verbose	numeric value that prints information on the fitting process; >2 is extremely verbose

normed	should the cordillera be normed; defaults to TRUE
scale	should the configuration be scale adjusted
stresstype	which stress to report. Only takes smacofs default stress currently.

Value

A list with the components

- stress: the stress
- stress.m: default normalized stress
- copstress: the weighted loss value
- OC: the Optics cordillera value
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure (which has all smacofB elements and some more)
- cordillera: the cordillera object

cop_sstress	<i>PCOPS version of sstress</i>
-------------	---------------------------------

Description

PCOPS version of sstress

Usage

```
cop_sstress(
  dis,
  theta = c(2, 1, 1),
  weightmat = 1 - diag(nrow(dis)),
  init = NULL,
  ndim = 2,
  itmaxi = 10000,
  ...,
  stressweight = 1,
  cordweight = 0.5,
  q = 1,
  minpts = ndim + 1,
  epsilon = 10,
  rang = NULL,
  verbose = 0,
  scale = "sd",
  normed = TRUE,
  stresstype = c("default", "stress1", "rawstress", "normstress", "enormstress",
    "enormstress1")
)
```

Arguments

<code>dis</code>	numeric matrix or dist object of a matrix of proximities
<code>theta</code>	the theta vector of powers; this is either a scalar of the lambda transformation for the observed proximities, or a vector where the first is the kappa argument for the fitted distances (here internally fixed to 2) and the second the lambda argument and the third the nu argument (internally fixed to 1). Defaults to 2 1 1
<code>weightmat</code>	(optional) a matrix of nonnegative weights
<code>init</code>	(optional) initial configuration
<code>ndim</code>	number of dimensions of the target space
<code>itmaxi</code>	number of iterations. default is 10000.
<code>...</code>	additional arguments to be passed to the fitting procedure
<code>stressweight</code>	weight to be used for the fit measure; defaults to 1
<code>cordweight</code>	weight to be used for the cordillera; defaults to 0.5
<code>q</code>	the norm of the cordillera; defaults to 1
<code>minpts</code>	the minimum points to make up a cluster in OPTICS; defaults to $ndim+1$
<code>epsilon</code>	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
<code>rang</code>	range of the distances (min distance minus max distance). If NULL (default) the cordillera will be normed to each configuration's maximum distance, so an absolute value of goodness-of-clusteredness.
<code>verbose</code>	numeric value that prints information on the fitting process; >2 is extremely verbose
<code>scale</code>	should the configuration be scale adjusted
<code>normed</code>	should the cordillera be normed; defaults to TRUE
<code>stresstype</code>	which stress to report? Defaults to explicitly normed stress

Value

A list with the components

- `stress`: the stress
- `stress.m`: default normalized stress
- `copstress`: the weighted loss value
- `OC`: the Optics cordillera value
- `parameters`: the parameters used for fitting (kappa, lambda)
- `fit`: the returned object of the fitting procedure
- `cordillera`: the cordillera object

doubleCenter	<i>Double centering of a matrix</i>
--------------	-------------------------------------

Description

Double centering of a matrix

Usage

doubleCenter(x)

Arguments

x numeric matrix

Value

the double centered matrix

enorm	<i>Explicit Normalization Normalizes distances</i>
-------	--

Description

Explicit Normalization Normalizes distances

Usage

enorm(x, w = 1)

Arguments

x numeric matrix

w weight

Value

a constant

ljoptim

*(Adaptive) Version of Luus-Jakola Optimization***Description**

Adaptive means that the search space reduction factors in the number of iterations; makes convergence faster at about 100 iterations

Usage

```
ljoptim(
  x,
  fun,
  ...,
  red = ifelse(adaptive, 0.99, 0.95),
  lower,
  upper,
  acc = 1e-06,
  accd = 1e-04,
  itmax = 1000,
  verbose = 0,
  adaptive = TRUE
)
```

Arguments

x	optional starting values
fun	function to minimize
...	additional arguments to be passed to the function to be optimized
red	value of the reduction of the search region
lower	The lower constraints of the search region
upper	The upper constraints of the search region
acc	if the numerical accuracy of two successive target function values is below this, stop the optimization; defaults to 1e-6
accd	if the width of the search space is below this, stop the optimization; defaults to 1e-4
itmax	maximum number of iterations
verbose	numeric value that prints information on the fitting process; >2 is extremely verbose
adaptive	should the adaptive version be used? defaults to TRUE.

Value

A list with the components (see also [optim](#))

- `par` The position of the optimum in the search space (parameters that minimize the function; `argmin fun`)
- `value` The value of the objective function at the optimum (`min fun`)
- `counts` The number of iterations performed at convergence with entries `fnction` for the number of iterations and `gradient` which is always NA at the moment
- `convergence` 0 successful completion by the `accd` or `acc` criterion, 1 indicate iteration limit was reached, 99 is a problem
- `message` is NULL (only for compatibility or future use)

Examples

```
fbana <- function(x) {
  x1 <- x[1]
  x2 <- x[2]
  100 * (x2 - x1 * x1)^2 + (1 - x1)^2
}
res1<-ljoptim(c(-1.2,1),fbana,lower=-5,upper=5,accd=1e-16,acc=1e-16)
res1

set.seed(210485)
fwild <- function (x) 10*sin(0.3*x)*sin(1.3*x^2) + 0.00001*x^4 + 0.2*x+80
plot(fwild, -50, 50, n = 1000, main = "ljoptim() minimising 'wild function'")
res2<-ljoptim(50, fwild,lower=-50,upper=50,adaptive=FALSE,accd=1e-16,acc=1e-16)
points(res2$par,res2$value,col="red",pch=19)
res2
```

mkBmat

Auxfunction1

Description

only used internally

Usage

```
mkBmat(x)
```

Arguments

`x` matrix

Value

a matrix

mkPower	<i>Take matrix to a power</i>
---------	-------------------------------

Description

Take matrix to a power

Usage

```
mkPower(x, r)
```

Arguments

x	matrix
r	numeric (power)

Value

a matrix

pcops	<i>Profile COPS Function (aka COPS Variant 2)</i>
-------	---

Description

Metaparameter selection for MDS models based on the Profile COPS approach (COPS Variant 2). It uses copstress for hyperparameter selection. It is a special case of a STOPS model.

Usage

```
pcops(
  dis,
  loss = c("stress", "smacofSym", "smacofSphere", "strain", "sammon", "rstress",
    "powermds", "sstress", "elastic", "powersammon", "powerelastic", "powerstress",
    "sammon2", "powerstrain", "apstress", "rpowerstress"),
  weightmat = NULL,
  ndim = 2,
  init = NULL,
  theta = c(1, 1, 1),
  stressweight = 1,
  cordweight,
  q = 2,
  minpts = ndim + 1,
  epsilon = 100,
  rang,
```

```

optimmethod = c("ALJ", "pso", "SANN", "DIRECT", "DIRECTL", "stogo", "MADS", "hjk"),
lower = c(1, 1, 0.5),
upper = c(5, 5, 2),
verbose = 0,
scale = c("proc", "sd", "none", "std"),
normed = TRUE,
s = 4,
stresstype = "default",
acc = 1e-07,
itmaxo = 200,
itmaxi = 10000,
...
)

```

Arguments

<code>dis</code>	numeric matrix or dist object of a matrix of proximities
<code>loss</code>	which loss function to be used for fitting, defaults to strain. Currently allows for the following models: <ul style="list-style-type: none"> • Power transformations of observed proximities only: Strain loss or classical scaling (<code>strain</code>, <code>workhorse</code> is <code>cmdscale</code>), Kruskal's stress for symmetric matrices (<code>smacofSym</code> or <code>stress</code> and <code>smacofSphere</code> for scaling onto a sphere; <code>workhorse</code> is <code>smacof</code>), Sammon mapping (<code>sammon</code> or <code>sammon2</code>; for the earlier the <code>workhorse</code> is <code>sammon</code> from MASS for the latter it is <code>smacof</code>), elastic scaling (<code>elastic</code>, the <code>workhorse</code> is <code>smacof</code>), Takane et al's S-Stress <code>sstress</code> (<code>workhorse</code> is <code>powerstressMin</code>) • Power transformations of fitted distances only: De Leeuw's r-stress <code>rstress</code> (<code>workhorse</code> is <code>powerstressMin</code>) • Power transformations of fitted distances and observed proximities: <code>Powermds</code> (<code>powermds</code>), Sammon mapping/elastic scaling with powers (<code>powersammon</code>, <code>powerelastic</code>), <code>powerstress</code> (POST-MDS, <code>powerstress</code>), restricted <code>powerstress</code> with equal transformations for delta and d (<code>rpowerstress</code>); <code>workhorse</code> is <code>powerstressMin</code>) • Approximation to power stress: Approximated power stress (<code>apstress</code>; <code>workhorse</code> is <code>smacof</code>)
<code>weightmat</code>	(optional) a matrix of nonnegative weights; defaults to 1 for all off diagonals
<code>ndim</code>	number of dimensions of the target space
<code>init</code>	(optional) initial configuration. If not supplied, the Torgerson scaling result of the dissimilarity matrix $dis^{\theta[2]}/\text{enorm}(dis^{\theta[2]}, weightmat)$ is used.
<code>theta</code>	the theta vector of powers; the first is kappa (for the fitted distances if it exists), the second lambda (for the observed proximities if it exist), the third is nu (for the weights if it exists). If a scalar is given as argument, it will take the role designated by the loss argument (typically recycled). Defaults to 1 1 1
<code>stressweight</code>	weight to be used for the fit measure; defaults to 1
<code>cordweight</code>	weight to be used for the cordillera; if missing gets estimated from the initial configuration so that $copstress = 0$ for $theta=c(1,1)$

q	the norm of the cordillera; defaults to 1
minpts	the minimum points to make up a cluster in OPTICS; defaults to ndim+1
epsilon	the epsilon parameter of OPTICS, the neighbourhood that is checked; defaults to 10
rang	range of the minimum reachabilities to be considered. If missing it is found from the initial configuration by taking 1.5 times the maximal minimum reachability of the model with theta=c(1,1). If NULL it will be normed to each configuration's minimum and maximum distance, so an absolute value of goodness-of-clusteredness. Note that the latter is not necessarily desirable when comparing configurations for their relative clusteredness. See also cordillera .
optimmethod	What general purpose optimizer to use? Defaults to our adaptive LJ version (ALJ). Also allows particle swarm optimization with s particles ("pso") and simulated annealing ("SANN"), "DIRECT" and "DIRECTL", Hooke-Jeeves ("hjk"), StoGo ("stogo"), and "MADS". We recommend not using SANN and pso with the rstress, sstress and the power stress models. We amde good experiences with ALJ, stogo, DIRECT and DIRECTL and also MADS.
lower	The lower constraints of the search region
upper	The upper constraints of the search region
verbose	numeric value hat prints information on the fitting process; >2 is extremely verbose. Note that for models with some parameters fixed, the iteration progress of the optimizer shows different values also for the fixed parameters because due to the modular setup we always optimize over a three parameter vector. These values are inconsequential however as internally they will be fixed.
scale	should the configuration be scaled and/or centered for calculating the cordillera? "std" standardizes each column of the configurations to mean=0 and sd=1 (typically not a good idea), "sd" scales the configuration by the maximum standard deviation of any column (default), "proc" adjusts the fitted configuration to the init configuration (or the Togerson scaling solution if init=NULL). This parameter only has an effect for calculating the cordillera, the fitted and returned configuration is NOT scaled.
normed	should the cordillera be normed; defaults to TRUE
s	number of particles if pso is used
stresstype	what stress to be used for comparisons between solutions. Currently not implemented and pcops uses explicitly normalized stress for copstress (not stress-1). Stress-1 is reported by the print function though.
acc	termination threshold difference of two successive outer minimization steps.
itmaxo	iterations of the outer step (optimization over the hyperparameters; if solver allows it). Defaults to 200.
itmaxi	iterations of the inner step (optimization of the MDS). Defaults to 10000 (which is huge).
...	additional arguments to be passed to the optimization procedure

Value

A list with the components

- copstress: the weighted loss value
- OC: the OPTICS cordillera for the scaled configuration (as defined by scale)
- optim: the object returned from the optimization procedure
- stress: the stress (square root of stress.m)
- stress.m: default normalized stress
- parameters: the parameters used for fitting (kappa, lambda)
- fit: the returned object of the fitting procedure
- cordillera: the cordillera object

Examples

```
dis<-as.matrix(smacof::kinshipdelta)
set.seed(210485)
#configuration is scaled with highest column sd for calculating cordilera
res1<-pcops(dis,loss="strain",lower=0.1,upper=5,minpts=2)
res1
summary(res1)
plot(res1)
```

pdist

Squared p-distances

Description

Squared p-distances

Usage

```
pdist(x, p)
```

Arguments

x	numeric matrix
p	p>0 the Minkoswki distance

Value

squared Minkowski distance matrix

plot.cops

S3 plot method for cops objects

Description

S3 plot method for cops objects

Usage

```
## S3 method for class 'cops'
plot(x, plot.type = c("confplot"), main, asp = 1, ...)
```

Arguments

- | | |
|-----------|---|
| x | an object of class cops |
| plot.type | String indicating which type of plot to be produced: "confplot", "reachplot", "resplot", "transplot", "Shepard", "stressplot" (see details) |
| main | the main title of the plot |
| asp | aspect ratio of x/y axis; defaults to NA; setting to 1 will lead to an accurate representation of the fitted distances. |
| ... | Further plot arguments passed: see 'plot.smacof' and 'plot' for detailed information. |
- Details:
- Configuration plot (plot.type = "confplot"): Plots the MDS configurations.
 - Reachability plot (plot.type = "confplot"): Plots the OPTICS reachability plot and the OPTICS cordillera
 - Residual plot (plot.type = "resplot"): Plots the dissimilarities against the fitted distances.
 - Linearized Shepard diagram (plot.type = "Shepard"): Diagram with the transformed observed dissimilarities against the transformed fitted distance as well as loess smooth and a least squares line.
 - Transformation Plot (plot.type = "transplot"): Diagram with the observed dissimilarities (lighter) and the transformed observed dissimilarities (darker) against the fitted distances together with loess smoothing lines
 - Stress decomposition plot (plot.type = "stressplot", only for SMACOF objects in \$fit): Plots the stress contribution in of each observation. Note that it rescales the stress-per-point (SPP) from the corresponding smacof function to percentages (sum is 100). The higher the contribution, the worse the fit.
 - Bubble plot (plot.type = "bubbleplot", only available for SMACOF objects \$fit): Combines the configuration plot with the point stress contribution. The larger the bubbles, the better the fit.

Examples

```
dis<-as.matrix(smacof::kinshipdelta)
resl<-copstressMin(dis,itmax=20)
plot(resl)
```

plot.pcop	<i>S3 plot method for p-cops objects</i>
-----------	--

Description

S3 plot method for p-cops objects

Usage

```
## S3 method for class 'pcops'
plot(x, plot.type = c("confplot"), main, asp = NA, ...)
```

Arguments

x	an object of class cops
plot.type	String indicating which type of plot to be produced: "confplot", "reachplot", "resplot", "transplot", "Shepard", "stressplot" (see details)
main	the main title of the plot
asp	aspect ratio of x/y axis; defaults to NA; setting to 1 will lead to an accurate representation of the fitted distances.
...	Further plot arguments passed: see 'plot.smacof' and 'plot' for detailed information. Details: <ul style="list-style-type: none"> • Configuration plot (plot.type = "confplot"): Plots the MDS configurations. • Reachability plot (plot.type = "confplot"): Plots the OPTICS reachability plot and the OPTICS cordillera • Residual plot (plot.type = "resplot"): Plots the dissimilarities against the fitted distances. • Linearized Shepard diagram (plot.type = "Shepard"): Diagram with the transformed observed dissimilarities against the transformed fitted distance as well as loess smooth and a least squares line. • Transformation Plot (plot.type = "transplot"): Diagram with the observed dissimilarities (lighter) and the transformed observed dissimilarities (darker) against the fitted distances together with loess smoothing lines • Stress decomposition plot (plot.type = "stressplot", only for SMACOF objects in \$fit): Plots the stress contribution in of each observation. Note that it rescales the stress-per-point (SPP) from the corresponding smacof function to percentages (sum is 100). The higher the contribution, the worse the fit. • Bubble plot (plot.type = "bubbleplot", only available for SMACOF objects \$fit): Combines the configuration plot with the point stress contribution. The larger the bubbles, the better the fit.

Examples

```
dis<-as.matrix(smacof::kinshipdelta)
resl<-pcops(dis,loss="strain",lower=0.1,upper=5,minpts=2)
plot(resl)
plot(resl,plot.type="Shepard")
```

plot.smacofP

S3 plot method for smacofP objects

Description

S3 plot method for smacofP objects

Usage

```
## S3 method for class 'smacofP'
plot(
  x,
  plot.type = "confplot",
  plot.dim = c(1, 2),
  bubscale = 5,
  col,
  label.conf = list(label = TRUE, pos = 3, col = 1, cex = 0.8),
  identify = FALSE,
  type = "p",
  pch = 20,
  asp = 1,
  main,
  xlab,
  ylab,
  xlim,
  ylim,
  legend = TRUE,
  legpos,
  loess = TRUE,
  ...
)
```

Arguments

x	an object of class smacofP
plot.type	String indicating which type of plot to be produced: "confplot", "resplot", "Shepard", "stressplot", "transplot", "bubbleplot" (see details)
plot.dim	dimensions to be plotted in confplot; defaults to c(1, 2)
bubscale	Scaling factor (size) for the bubble plot
col	vector of colors for the points

label.conf	List with arguments for plotting the labels of the configurations in a configuration plot (logical value whether to plot labels or not, label position, label color)
identify	If 'TRUE', the 'identify()' function is called internally that allows to add configuration labels by mouse click
type	What type of plot should be drawn (see also 'plot')
pch	Plot symbol
asp	Aspect ratio; defaults to 1 so distances between x and y are represented accurately; can lead to slightly weird looking plots if the variance on one axis is much smaller than on the other axis; use NA if the standard type of R plot is wanted where the ylim and xlim arguments define the aspect ratio - but then the distances seen are no longer accurate
main	plot title
xlab	label of x axis
ylab	label of y axis
xlim	scale of x axis
ylim	scale of y axis
legend	Flag whether legends should be drawn for plots that have legends
legpos	Position of legend in plots with legends
loess	should loess fit be added to Shepard plot
...	Further plot arguments passed: see 'plot.smacof' and 'plot' for detailed information.

Details:

- Configuration plot (plot.type = "confplot"): Plots the MDS configurations.
- Residual plot (plot.type = "resplot"): Plots the dissimilarities against the fitted distances.
- Linearized Shepard diagram (plot.type = "Shepard"): Diagram with the transformed observed dissimilarities against the transformed fitted distance as well as loess curve and a least squares line.
- Transformation Plot (plot.type = "transplot"): Diagram with the observed dissimilarities (lighter) and the transformed observed dissimilarities (darker) against the fitted distances together with the nonlinear regression curve
- Stress decomposition plot (plot.type = "stressplot"): Plots the stress contribution in of each observation. Note that it rescales the stress-per-point (SPP) from the corresponding smacof function to percentages (sum is 100). The higher the contribution, the worse the fit.
- Bubble plot (plot.type = "bubbleplot"): Combines the configuration plot with the point stress contribution. The larger the bubbles, the better the fit.

Examples

```
dis<-as.matrix(smacof::kinshipdelta)
res<-powerStressMin(dis)
plot(res)
plot(res,"reachplot")
```

```

plot(res,"Shepard")
plot(res,"resplot")
plot(res,"transplot")
plot(res,"stressplot")
plot(res,"bubbleplot")

```

plot3dstatic *plot3dstatic: static 3D plots*

Description

A static 3d plot S3 generic

Usage

```
plot3dstatic(x, plot.dim = c(1, 2, 3), main, xlab, ylab, zlab, col, ...)
```

Arguments

x	object
plot.dim	dimensions to plot
main	main title
xlab	label for x axis
ylab	label for y axis
zlab	label for z axis
col	color
...	other arguments

Details

A static 3d plot

plot3dstatic.cmdscale *3D plots: plot3dstatic method for class cmdscale*

Description

This methods produces a static 3D configuration plot.

Usage

```

## S3 method for class 'cmdscale'
plot3dstatic(x, plot.dim = c(1, 2, 3), main, xlab, ylab, zlab, col, ...)

```

Arguments

x	object of class cmdscale
plot.dim	vector of length 3 with dimensions to be plotted
main	plot title
xlab	label of x axis
ylab	label of y axis
zlab	label of z axis
col	color of the text labels
...	Further plot arguments passed: see 'scatterplot3d' in package 'scatterplot3d' for detailed information.

powerStressFast	<i>Power stress minimization by NEWUOA</i>
-----------------	--

Description

An implementation to minimize power stress by a derivative-free trust region optimization algorithm (NEWUOA). Much faster than majorizing as used in powerStressMin but perhaps less accurate.

Usage

```
powerStressFast(
  delta,
  kappa = 1,
  lambda = 1,
  nu = 1,
  weightmat = 1 - diag(nrow(delta)),
  init = NULL,
  ndim = 2,
  acc = 1e-12,
  itmax = 50000,
  verbose = FALSE
)
```

Arguments

delta	dist object or a symmetric, numeric data.frame or matrix of distances
kappa	power of the transformation of the fitted distances; defaults to 1
lambda	the power of the transformation of the proximities; defaults to 1
nu	the power of the transformation for weightmat; defaults to 1
weightmat	a matrix of finite weights
init	starting configuration

ndim	dimension of the configuration; defaults to 2
acc	The smallest value of the trust region radius that is allowed. If not defined, then 1e-10 will be used.
itmax	maximum number of iterations. Default is 50000.
verbose	should iteration output be printed; if > 1 then yes

Value

a smacofP object (inheriting from smacofB, see [smacofSym](#)). It is a list with the components

- delta: Observed dissimilarities, not normalized
- obsdiss: Observed dissimilarities, normalized
- confdist: Configuration dissimilarities, NOT normalized
- conf: Matrix of fitted configuration, NOT normalized
- stress: Default stress (stress 1, square root of the explicitly normalized stress on the normalized, transformed dissimilarities)
- spp: Stress per point (based on stress.en)
- ndim: Number of dimensions
- model: Name of smacof model
- niter: Number of iterations
- nobj: Number of objects
- type: Type of MDS model

and some additional components

- gamma: Empty
- stress.m: default stress for the COPS and STOP. Defaults to the explicitly normalized stress on the normalized, transformed dissimilarities
- stress.en: explicitly stress on the normalized, transformed dissimilarities and normalized transformed distances
- deltaorig: observed, untransformed dissimilarities
- weightmat: weighting matrix

See Also

[smacofSym](#)

Examples

```
dis<-smacof::kinshipdelta
res<-powerStressFast(as.matrix(dis),kappa=2,lambda=1.5)
res
summary(res)
plot(res)
```

powerStressMin *Power Stress SMACOF*

Description

An implementation to minimize power stress by minimization-majorization. Usually more accurate but slower than powerStressFast. Uses a repeat loop.

Usage

```
powerStressMin(
  delta,
  kappa = 1,
  lambda = 1,
  nu = 1,
  weightmat = 1 - diag(nrow(delta)),
  init = NULL,
  ndim = 2,
  acc = 1e-10,
  itmax = 50000,
  verbose = FALSE
)
```

Arguments

delta	dist object or a symmetric, numeric data.frame or matrix of distances
kappa	power of the transformation of the fitted distances; defaults to 1
lambda	the power of the transformation of the proximities; defaults to 1
nu	the power of the transformation for weightmat; defaults to 1
weightmat	a matrix of finite weights
init	starting configuration
ndim	dimension of the configuration; defaults to 2
acc	numeric accuracy of the iteration
itmax	maximum number of iterations. Default is 50000.
verbose	should iteration output be printed; if > 1 then yes

Value

a smacofP object (inheriting from smacofB, see [smacofSym](#)). It is a list with the components

- delta: Observed dissimilarities, not normalized
- obsdiss: Observed dissimilarities, normalized
- confdist: Configuration dissimilarities, NOT normalized
- conf: Matrix of fitted configuration, NOT normalized

- stress: Default stress (stress 1; sqrt of explicitly normalized stress)
- spp: Stress per point (based on stress.en)
- ndim: Number of dimensions
- model: Name of smacof model
- niter: Number of iterations
- nobj: Number of objects
- type: Type of MDS model

and some additional components

- stress.m: default stress for the COPS and STOP defaults to the explicitly normalized stress on the normalized, transformed dissimilarities
- stress.en: a manually calculated stress on the normalized, transformed dissimilarities and normalized transformed distances which is not correct
- deltaorig: observed, untransformed dissimilarities
- weightmat: weighting matrix

See Also

[smacofSym](#)

Examples

```
dis<-smacof::kinshipdelta
res<-powerStressMin(as.matrix(dis),kappa=2,lambda=1.5,itmax=1000)
res
summary(res)
plot(res)
```

procruster

procruster: a procrustes function

Description

procruster: a procrustes function

Usage

```
procruster(x)
```

Arguments

x numeric matrix

Value

a matrix

sammon	<i>Wrapper to sammon for S3 class</i>
--------	---------------------------------------

Description

Wrapper to sammon for S3 class

Usage

```
sammon(d, y = NULL, k = 2, ...)
```

Arguments

d	a distance structure such as that returned by 'dist' or a full symmetric matrix. Data are assumed to be dissimilarities or relative distances, but must be positive except for self-distance. This can contain missing values.
y	An initial configuration. If NULL, 'cmdscale' is used to provide the classical solution. (If there are missing values in 'd', an initial configuration must be provided.) This must not have duplicates.
k	The dimension of the configuration
...	Additional parameters passed to sammon, see sammon

Value

See [sammon](#). This wrapper only adds an extra slot to the list with the call, adds column labels to the \$points and assigns S3 classes 'sammon', 'cmdscale'. It also adds a slot obsdiss with normalized dissimilarities.

Examples

```
dis<-as.matrix(smacof::kinshipdelta)
res<-sammon(dis)
```

scale_adjust	<i>Adjusts a configuration</i>
--------------	--------------------------------

Description

Adjusts a configuration

Usage

```
scale_adjust(conf, ref, scale = c("sd", "std", "proc", "none"))
```

Arguments

conf	a configuration
ref	a reference configuration (only for scale="proc")
scale	Scale adjustment. "std" standardizes each column of the configurations to mean=0 and sd=1, "sd" scales the configuration by the maximum standard deviation of any column, "proc" adjusts the fitted configuration to the reference

Value

The scale adjusted configuration.

secularEq	<i>Secular Equation</i>
-----------	-------------------------

Description

Secular Equation

Usage

secularEq(a, b)

Arguments

a	matrix
b	matrix

Value

a matrix

sqdist	<i>Squared distances</i>
--------	--------------------------

Description

Squared distances

Usage

sqdist(x)

Arguments

x	numeric matrix
---	----------------

Value

squared distance matrix

torgerson

Torgerson scaling

Description

Torgerson scaling

Usage

```
torgerson(delta, p = 2)
```

Arguments

delta symmetric, numeric matrix of distances
p target space dimensions

Value

a n x p matrix (the configuration)

Examples

```
dis<-as.matrix(smacof::kinshipdelta)  
res<-torgerson(dis)
```

Index

- * **clustering**
 - cops, 5
 - copstressMin, 9
 - pcops, 38
- * **multivariate**
 - cop_apstress, 12
 - cop_cmdscale, 14
 - cop_elastic, 15
 - cop_powerelastic, 17
 - cop_powermds, 19
 - cop_powersammon, 20
 - cop_powerstress, 22
 - cop_rpowerstress, 23
 - cop_rstress, 25
 - cop_sammon, 27
 - cop_sammon2, 28
 - cop_smacofSphere, 30
 - cop_smacofSym, 31
 - cop_sstress, 33
 - cops, 5
 - copstress, 8
 - copstressMin, 9
 - pcops, 38
- BankingCrisesDistances, 3
- cmdscale, 3, 4
- conf_adjust, 4
- cop_apstress, 12
- cop_cmdscale, 14
- cop_elastic, 15
- cop_powerelastic, 17
- cop_powermds, 19
- cop_powersammon, 20
- cop_powerstress, 22
- cop_rpowerstress, 23
- cop_rstress, 25
- cop_sammon, 27
- cop_sammon2, 28
- cop_smacofSphere, 30
- cop_smacofSym, 31
- cop_sstress, 33
- cops, 5
- copstress, 8
- copstressMin, 5, 6, 9
- cordillera, 10, 40
- doubleCenter, 35
- enorm, 35
- ljoptim, 36
- mkBmat, 37
- mkPower, 38
- optim, 37
- pcops, 5, 6, 38
- pdist, 41
- plot.cops, 42
- plot.pcops, 43
- plot.smacofP, 44
- plot3dstatic, 46
- plot3dstatic.cmdscale, 46
- powerStressFast, 47
- powerStressMin, 49
- procruster, 50
- sammon, 51, 51
- scale_adjust, 51
- secularEq, 52
- smacofSym, 48–50
- sqdist, 52
- torgerson, 53