

Package ‘ctmva’

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Type Package

Title Continuous-Time Multivariate Analysis

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Description Implements a basis function or functional data analysis framework for several techniques of multivariate analysis in continuous-time setting. Specifically, we introduced continuous-time analogues of several classical techniques of multivariate analysis, such as principal component analysis, canonical correlation analysis, Fisher linear discriminant analysis, K-means clustering, and so on. Details are in Biplab Paul, Philip T. Reiss and Erjia Cui (2023) ``Continuous-time multivariate analysis" <[doi:10.48550/arXiv.2307.09404](https://doi.org/10.48550/arXiv.2307.09404)>.

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ctmva-package	<i>Continuous-time multivariate analysis</i>
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Description

Implements continuous-time analogues of several classical techniques of multivariate analysis. The inputs are "[fd](#)" (functional data) objects from the **fd**a package.

Author(s)

Bi-plab Paul <paul.bi-plab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

References

Paul, B., Reiss, P. T. and Cui, E. (2023). Continuous-time multivariate analysis. arXiv:2307.09404 [stat.ME]

cca.ct	<i>Continuous-time canonical correlation analysis</i>
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Description

A continuous-time version of canonical correlation analysis (CCA).

Usage

cca.ct(fdobj1, fdobj2)

Arguments

fdobj1, fdobj2 a pair of continuous-time multivariate data sets, of class "[fd](#)"

Value

A list consisting of

vex1, vex2	matrices defining the canonical variates. The first columns of each give the coefficients defining the first pair of canonical variates; and so on.
cor	canonical correlations, i.e., correlations between the pairs of canonical variates

Note

Columns of the output matrix vex2 are flipped as needed to ensure positive correlations.

Author(s)

Biplab Paul <paul.biplab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[cancor](#), for classical CCA

Examples

```
## Not run:

# CCA relating Canadian daily temperature and precipitation data
require(fda)
data(CanadianWeather)
daybasis <- create.bspline.basis(c(0,365), nbasis=80)
tempfd <- smooth.basis(day.5, CanadianWeather$dailyAv[,,"Temperature.C"], daybasis)$fd
precfd <- smooth.basis(day.5, CanadianWeather$dailyAv[,,"log10precip"], daybasis)$fd
tpcor <- cca.ct(tempfd, precfd)
par(mfrow=1:2)
barplot(tpcor$vex1[,1], horiz=TRUE, las=1, main="Temperature",
        sub="First canonical coefficients vector")
barplot(tpcor$vex2[,1], horiz=TRUE, las=1, main="Log precipitation",
        sub="First canonical coefficients vector")

## End(Not run)
```

Description

Subtracts the (continuous-time) mean of each of the variables. This is analogous to column-centering an $n \times p$ data matrix.

Usage

```
center.ct(fdobj)
```

Arguments

fdobj continuous-time multivariate data set of class "[fd](#)"

Value

A centered version of the input data.

Author(s)

Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[standardize.ct](#)

cor.ct

Continuous-time correlation or cross-correlation matrix

Description

Computes the correlation matrix of a continuous-time multivariate data set represented as an [fd](#) object; or the cross-correlation matrix of two such data sets.

Usage

```
cor.ct(fdobj1, fdobj2 = fdobj1, common_trend = FALSE)
```

Arguments

fdobj1 continuous-time multivariate data set of class "[fd](#)"
fdobj2 an optional second data set
common_trend logical: centering wrt mean function if TRUE, without centering if FALSE (the default)

Value

A matrix of (cross-) correlations

Author(s)

Biplot Paul <paul.biplot497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[center.fd](#), for centering of "fd" objects; [inprod.cent](#)

Examples

```
## Not run:

# Canadian temperature data

require(fda)
require(corrplot)
data(CanadianWeather)
daybasis <- create.fourier.basis(c(0,365), nbasis=55)
tempfd <- smooth.basis(day.5, CanadianWeather$dailyAv[,,"Temperature.C"], daybasis)$fd

## The following yields a matrix of correlations that are all near 1:
rawcor <- cor.ct(tempfd)
corrplot(rawcor, method = 'square', type = 'lower', tl.col="black", tl.cex = 0.6)
## This occurs due to a strong seasonal trend that is common to all stations
## Removing this common trend leads to a more interesting result:
dtcor <- cor.ct(tempfd, common_trend = TRUE)
ord <- corrMatOrder(dtcor)
dtcord <- dtcor[ord,ord]
corrplot(dtcord, method = 'square', type = 'lower', tl.col="black", tl.cex = 0.6)

## End(Not run)
```

cov.ct

Continuous-time covariance or cross-covariance matrix

Description

Computes the covariance matrix of a continuous-time multivariate data set represented as an [fd](#) object; or the cross-covariance matrix of two such data sets.

Usage

```
cov.ct(fdobj1, fdobj2 = fdobj1, common_trend = FALSE)
```

Arguments

fdobj1	continuous-time multivariate data set of class " fd "
fdobj2	an optional second data set
common_trend	logical: centering with respect to the mean function if TRUE, without centering if FALSE (the default)

Value

A matrix of (cross-) covariances

Author(s)

Biplab Paul <paul.biplab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[cor.ct](#)

Examples

```
# see example for cor.ct, which works similarly
```

inprod.cent

Centered inner product matrix for a basis or pair of bases

Description

Several methods of continuous-time multivariate analysis require a matrix of inner products of pairs of centered functions from a basis, such as a B-spline basis, or pairs consisting of one function from each of two bases. This function computes such matrices via 7-point Newton-Cotes integration, which is exact for cubic B-splines. For a Fourier basis with the inner product taken over the entire range, a simple closed form is used instead of integration.

Usage

```
inprod.cent(basis1, basis2 = basis1, rng = NULL)
```

Arguments

basis1	basis object from the fda package.
basis2	an optional second basis
rng	time range. By default, the entire range spanned by the basis, or the intersection of the ranges of the two bases.

Value

Matrix of inner products of each pair of centered basis functions.

Author(s)

Biplab Paul <paul.biplab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

`create.bspline.basis` from package `fda`, for the most commonly used basis object type.

Examples

```
## Not run:

require(fda)
bbasis6 <- create.bspline.basis(nbasis=6)
inprod.cent(bbasis6)
fbasis7 <- create.fourier.basis(nbasis=7)
inprod.cent(fbasis7)

## End(Not run)
```

kmeans.ct

Continuous-time k-means clustering

Description

A continuous-time version of k-means clustering in which each cluster is a time segments or set of time segments.

Usage

```
kmeans.ct(
  fdobj,
  k,
  common_trend = FALSE,
  init.pts = NULL,
  tol = 0.001,
  max.iter = 100
)
```

Arguments

<code>fdobj</code>	continuous-time multivariate data set of class " <code>fd</code> "
<code>k</code>	number of clusters
<code>common_trend</code>	logical: Should the curves be centered with respect to the mean function? Defaults to FALSE.
<code>init.pts</code>	a set of <code>k</code> time points. The observations at these time points serve as initial values for the <code>k</code> means. Randomly generated if not supplied.
<code>tol</code>	convergence tolerance for the <code>k</code> means
<code>max.iter</code>	maximum number of iterations

Value

Object of class "kmeans.ct", a list consisting of

<code>fdobj</code>	the supplied <code>fdobj</code>
<code>means</code>	means of the <code>k</code> clusters
<code>transitions</code>	transition points between segments
<code>cluster</code>	cluster memberships in the segments defined by the transitions
<code>size</code>	length of each cluster, i.e. sum of lengths of subintervals making up each cluster
<code>totisd</code>	total integrated sum of distances from the overall mean; this is the analogue of <code>totss</code> from <code>link{kmeans}</code>
<code>withinisd</code>	within-cluster integrated sum of distances, i.e. integrated sum of distances from each cluster mean
<code>tot.withinisd</code>	total within-cluster integrated sum of distances, i.e. <code>sum(withinisd)</code>
<code>betweenisd</code>	between-cluster integrated sum of distances, i.e. <code>totisd-tot.withinss</code>

Author(s)

Biplab Paul <paul.biplab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[kmeans](#), [plot.kmeans.ct](#)

Examples

```
## Not run:

require(fda)
data(CanadianWeather)
daybasis <- create.bspline.basis(c(0,365), nbasis=55)
tempfd <- smooth.basis(day.5, CanadianWeather$dailyAv[, "Temperature.C"], daybasis)$fd
ktemp3 <- kmeans.ct(tempfd, 3)
plot(ktemp3, axes=FALSE)
axesIntervals(); box()
plot(silhouette.ct(ktemp3), axes=FALSE)
axesIntervals(); box()

## End(Not run)
```

lda.ct	<i>Continuous-time Fisher's linear discriminant analysis</i>
--------	--

Description

A continuous-time version of Fisher's LDA, in which segments of the time interval take the place of groups of observations.

Usage

```
lda.ct(fdobj, partition, part.names = NULL)
```

Arguments

fdobj	continuous-time multivariate data set of class " fd "
partition	a priori break points dividing the time interval into segments
part.names	optional character vector of names for the segments

Details

The means and scaling components of the output are similar to [lda](#), but unlike that function, `lda.ct` performs only *Fisher's* LDA and cannot incorporate priors or perform classification.

Value

Object of class "`lda.ct`", a list consisting of

means	means of the variables within each segment
scaling	matrix of coefficients defining the discriminants (as in lda)
values	eigenvalues giving the ratios of between to within sums of squares
partition	the supplied partition
fdobj	linear discriminants represented as an " fd " object
nld	number of linear discriminants

Author(s)

Biplab Paul <paul.biplab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[plot.lda.ct](#); [lda](#), for the classical version

Examples

```
## see end of example in ?pca.ct
```

meanbasis	<i>Compute means of basis functions</i>
-----------	---

Description

Given a basis object as defined in the **fda** package (see [basisfd](#)), this function simply computes the vector of means of the basis functions. Used internally.

Usage

```
meanbasis(basis, rng = NULL)
```

Arguments

basis	a basis object of class " basisfd "
rng	time range. By default, the entire interval spanned by the basis. Must be left NULL for Fourier bases.

Value

Vector of means of the basis functions

Author(s)

Bi-lab Paul <paul.biplab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

Examples

```
require(fda)
bbasis6 <- create.bspline.basis(nbasis=6)
meanbasis(bbasis6)
meanbasis(bbasis6, c(.3,.6))
fbasis11 <- create.fourier.basis(nbasis=11)
meanbasis(fbasis11)
```

pca.ct	<i>Continuous-time principal component analysis</i>
--------	---

Description

A continuous-time version of principal component analysis.

Usage

```
pca.ct(fdobj, cor = FALSE, common_trend = FALSE)
```

Arguments

<code>fdobj</code>	continuous-time multivariate data set of class " <code>fd</code> "
<code>cor</code>	logical: use correlation matrix if TRUE, covariance if FALSE (the default)
<code>common_trend</code>	logical: Should the curves be centered with respect to the mean function? Defaults to FALSE.

Value

Returns a list including:

<code>var</code>	variances of the principal components.
<code>loadings</code>	the matrix of loadings (i.e., its columns are the eigenvectors of the continuous-time covariance).
<code>scorefd</code>	score functions.

Author(s)

Biplab Paul <paul.biplab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[cov.ct](#); [princomp](#), for the classical version

Examples

```
## Not run:

# Data for one session from a classic EEG data set
require(fda)
require(eegkit)
data(eegdata)
data(eegcoord)
longdat <- subset(eegdata, subject=="co2a0000369" & trial==0)
widedat <- reshape(longdat, direction="wide", drop=c("subject","group","condition","trial"),
  v.names="voltage", idvar="channel")

# Convert time series for 64 channels to a functional data object
bsb <- create.bspline.basis(c(0,255),nbasis=30)
fdo <- Data2fd(argvals=0:255, y=t(as.matrix(widedat[, -1])), basisobj=bsb)
plot(fdo)

# Now do PCA and display first loadings for 3 PC's,
# along with percent variance explained by each
pcc <- pca.ct(fdo)
pve <- 100*pcc$var/sum(pcc$var)
par(mfrow=c(1,3))
cidx <- match(widedat[,1],rownames(eegcoord))
eegspace(eegcoord[cidx,4:5],pcc$loadings[,1], colorlab="PC1 loadings",
  main=paste0(round(pve[1],0), "%"), mar=c(17,3,12,2), cex.main=2)
eegspace(eegcoord[cidx,4:5],pcc$loadings[,2], colorlab="PC2 loadings",
```

```

        main=paste0(round(pve[2],0), "%"), mar=c(17,3,12,2), cex.main=2)
eegspace(eegcoord[cidx,4:5],pcc$loadings[,3], colorlab="PC3 loadings",
        main=paste0(round(pve[3],0), "%"), mar=c(17,3,12,2), cex.main=2)

# Linear discriminant analysis: discriminating among the 1st, 2nd and 3rd portions
# of the time interval
ld <- lda.ct(fdo, c(85,170))
plot(ld)
eegspace(eegcoord[cidx,4:5],ld$scaling[,1], colorlab="LD1 coefficients",
        mar=c(17,3,12,2), cex.main=2)
eegspace(eegcoord[cidx,4:5],ld$scaling[,2], colorlab="LD2 coefficients",
        mar=c(17,3,12,2), cex.main=2)

## End(Not run)

```

plot.kmeans.ct	<i>Plot a kmeans.ct object</i>
----------------	--------------------------------

Description

Plots a continuous-time k-means clustering object generated by a call to [kmeans.ct](#).

Usage

```

## S3 method for class 'kmeans.ct'
plot(
  x,
  plottype = "functions",
  mark.transitions = TRUE,
  col = NULL,
  lty = NULL,
  xlab = "Time",
  ylab = NULL,
  legend = TRUE,
  ncol.legend = 1,
  cex.legend = 1,
  ...
)

```

Arguments

x	clustering object produced by kmeans.ct
plottype	either "functions" (the default), to display each variable as a smooth function of time, or "distance", to plot distances from the k cluster means versus time.
mark.transitions	logical: Should transitions between clusters be marked with vertical lines? Defaults to TRUE.

col	plot colors
lty	line type
xlab, ylab	x- and y-axis labels
legend	either a logical variable (whether a legend should be included) or a character vector to appear in the legend. Default is TRUE.
ncol.legend	number of columns for legend
cex.legend	character expansion factor for legend
...	other arguments passed to matplot

Value

None; a plot is generated.

Author(s)

Philip Tzvi Reiss <reiss@stat.haifa.ac.il> and Biplab Paul <paul.biplab497@gmail.com>

See Also

[kmeans.ct](#), which includes an example

plot.lda.ct	<i>Plot an lda.ct object</i>
-------------	------------------------------

Description

Plots the Fisher's linear discriminant functions generated by a call to [lda.ct](#).

Usage

```
## S3 method for class 'lda.ct'
plot(x, ylab = "Discriminants", xlab = "Time", which = NULL, col = NULL, ...)
```

Arguments

x	linear discriminant analysis object produced by lda.ct
ylab, xlab	y- and x-axis labels
which	which of the linear discriminants to plot
col	color vector
...	other arguments passed to matplot

Value

None; a plot is generated.

Author(s)

Bioplab Paul <paul.bioplab497@gmail.com> and Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[lda.ct](#)

Examples

```
## see the example at the end of ?pca.ct
```

plot.silhouette.ct	<i>Plot a silhouette.ct object</i>
--------------------	------------------------------------

Description

Plots the silhouette index, generated by a call to [silhouette.ct](#), for a continuous-time k-means clustering object.

Usage

```
## S3 method for class 'silhouette.ct'
plot(x, mark.transitions = TRUE, xlab = "Time", ylab = "Silhouette", ...)
```

Arguments

x	silhouette object produced by silhouette.ct
mark.transitions	logical: Should transitions between clusters be marked with vertical lines? Defaults to TRUE.
xlab, ylab	x- and y-axis labels
...	other arguments passed to plot

Value

None; a plot is generated.

Author(s)

Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[kmeans.ct](#), which includes an example; [silhouette.ct](#)

silhouette.ct*Silhouettes for continuous-time k-means clustering*

Description

Computes the silhouette index, at a grid of time points, for a continuous-time k-means clustering object produced by [kmeans.ct](#).

Usage

```
silhouette.ct(kmobj, ngrid = 5000)
```

Arguments

kmobj	continuous-time k-means clustering from kmeans.ct
ngrid	number of equally spaced grid points at which to compute the silhouette index

Value

Object of class "silhouette.ct", a list consisting of

grid	grid of ngrid points spanning the time range
value	silhouette index at each point along the grid
transitions	transition points between segments
cluster	cluster memberships in the segments defined by the transitions
mean	mean silhouette index

Note

An error is issued if the grid of time points contains one or more of the cluster transition points. This should not ordinarily occur, but if it does, it can be remedied by modifying `ngrid`.

Author(s)

Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

[kmeans.ct](#), which includes an example; [plot.silhouette.ct](#)

`standardize.ct`*Center and scale a continuous-time multivariate data set*

Description

Subtracts the (continuous-time) mean and divides by the (continuous-time) standard deviation of each of the variables. This is the continuous-time analogue of taking an $n \times p$ data matrix, subtracting the mean of each column, and dividing by the standard deviation of each column, as is done by `scale(..., center=TRUE, scale=TRUE)`.

Usage

```
standardize.ct(fdobj)
```

Arguments

`fdobj` continuous-time multivariate data set of class "`fd`"

Value

A standardized (centered and scaled) version of the input data.

Author(s)

Philip Tzvi Reiss <reiss@stat.haifa.ac.il>

See Also

`center.ct`

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