

Package ‘rties’

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Title Modeling Interpersonal Dynamics

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Description The name of this package grew out of our research on temporal interpersonal emotion systems (TIES), hence 'rties'. It provides tools for using a set of models to investigate temporal processes in bivariate (e.g., dyadic) systems. The general approach is to model, one dyad at a time, the dynamics of a variable that is assessed repeatedly from both partners, extract the parameter estimates for each dyad, and then use those parameter estimates as input to a latent profile analysis to extract groups of dyads with qualitatively distinct dynamics. Finally, the profile memberships can be used to either predict, or be predicted by, another variable of interest. Currently, 2 models are supported: 1) inertia-coordination, and 2) a coupled-oscillator. Extended documentation is provided in vignettes. Theoretical background can be found in Butler (2011) <doi:10.1177/1088868311411164> and Butler & Barnard (2019) <doi:10.1097/PSY.0000000000000703>.

Depends R (>= 2.10)

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actorPartnerDataCross *Takes individual cross-sectional data from dyads and turns it into actor-partner format.*

Description

Need to use a person ID that has first person in dyad numbered 1-n and second person in dyad = ID + some number larger than the number of dyads. Need dyad ID numbered same as for person ID for the first person in the dyad. Both members in each dyad need to have the same number of rows (rows of missing data are ok)

Usage

```
actorPartnerDataCross(basedata, dyadId, personId)
```

Arguments

basedata	A dataframe with cross-sectional dyadic data.
dyadId	The name of variable indicating dyad ID.
personId	The name of the variable indicating peron ID.

Value

A dataframe in actor-partner format.

Examples

```
data <- rties_ExampleDataShort
newData1 <- data[data$time==1, ] # make a cross-sectional dataframe
newData2 <- actorPartnerDataCross(basedata=newData1, dyadId="couple", personId="couple")
head(newData2)
```

actorPartnerDataTime *Takes individual repeated measures data from dyads and turns it into actor-partner format.*

Description

Need to use a person ID that has first person in dyad numbered 1-n and second person in dyad = ID + some number larger than the number of dyads. Need dyad ID numbered same as for person ID for the first person in the dyad. Both members in each dyad need to have the same number of rows (rows of missing data are ok).

Usage

```
actorPartnerDataTime(basedata, dyadId, personId)
```

Arguments

basedata	A dataframe with repeated measures dyadic data
dyadId	The name of variable indicating dyad ID.
personId	The name of the variable indicating peron ID.

Value

A dataframe in actor-partner format.

Examples

```
data <- rties_ExampleDataShort
newData <- actorPartnerDataTime(basedata=data, dyadId="couple", personId="couple")
head(newData)
```

autoCorPlots	<i>Produces auto-correlation plots of the observed state variable for lags of -+ 20 time steps for each dyad.</i>
--------------	---

Description

Produces auto-correlation plots of the observed state variable for lags of -+ 20 time steps for each dyad.

Usage

```
autoCorPlots(basedata, dyadId, personId, obs_name, time_name)
```

Arguments

basedata	A user provided dataframe.
dyadId	The name of the column in the dataframe that has the dyad-level identifier.
personId	The name of the column in the dataframe that has the person-level identifier.
obs_name	The name of the column in the dataframe that has the time-varying observable (e.g., the variable for which dynamics will be assessed).
time_name	The name of the column in the dataframe that indicates sequential temporal observations.

Value

Prints the plots to the screen.

Examples

```
data <- rties_ExampleDataShort
autoCorPlots(basedata=data, dyadId="couple", personId="person", obs_name="dial", time_name="time")
```

cloCoupledOde	<i>Provides the equation for a coupled oscillator model for the differential equation solver (ode) to plot</i>
---------------	--

Description

Provides the equation for a coupled oscillator model for the differential equation solver (ode) to plot

Usage

```
cloCoupledOde(t, state, parameters)
```

Arguments

t	A parameter used by the ode and passed by functions calling cloCoupleOde
state	Another parameter used by the ode and passed by functions calling cloCoupleOde
parameters	Another parameter used by the ode and passed by functions calling cloCoupleOde

Value

A list with the rates of change for each state variable.

cloPlotTraj	<i>Plots the bivariate state variable's clo model-predicted temporal trajectories for each latent profile of clo parameters.</i>
-------------	--

Description

Plots the bivariate state variable's clo model-predicted temporal trajectories for each latent profile of clo parameters.

Usage

```

cloPlotTraj(
  prepData,
  paramEst,
  n_profiles,
  dist0name = NULL,
  dist1name = NULL,
  plot_obs_name = NULL,
  minMax = NULL,
  time_length = NULL,
  printPlots = T
)

```

Arguments

prepData	A dataframe that was produced with the "dataPrep" function.
paramEst	A dataframe created by indivClo containing the clo parameter estimates for each dyad.
n_profiles	The number of latent profiles.
dist0name	An optional name for the level-0 of the distinguishing variable (e.g., "Women"). Default is dist0.
dist1name	An optional name for the level-1 of the distinguishing variable (e.g., "Men"). Default is dist1
plot_obs_name	An optional name for the observed state variable to appear on plots (e.g., "Emotional Experience").
minMax	An optional vector with desired minimum and maximum quantiles to be used for setting the y-axis range on the plots, e.g., minMax <- c(.1, .9) would set the y-axis limits to the 10th and 90th percentiles of the observed state variables. If not provided, the default is to use the minimum and maximum observed values of the state variables.
time_length	An optional value specifying how many time points to plot across. Default is the 75th percentile for the observed time variable.
printPlots	If true (the default) plots are displayed on the screen.

Value

The function returns the plots as a list.

Examples

```
# See vignettes for examples.
```

cloResids	<i>Produces histograms of the residuals from the oscillator model for each dyad.</i>
-----------	--

Description

Produces histograms of the residuals from the oscillator model for each dyad.

Usage

```
cloResids(derivData, whichModel, printPlots = T)
```

Arguments

derivData	A dataframe that was produced with the "estDerivs" function.
whichModel	Whether the model to be estimated is the uncoupled-oscillator ("uncoupled") or the coupled-oscillator ("coupled").
printPlots	If true (the default) plots are displayed on the screen.

Value

A list with the histograms of the residuals for each dyad.

Examples

```
# See vignettes for examples.
```

cloUncoupledOde	<i>Provides the equation for an un-coupled oscillator model for the differential equation solver (ode) to plot</i>
-----------------	--

Description

Provides the equation for an un-coupled oscillator model for the differential equation solver (ode) to plot

Usage

```
cloUncoupledOde(t, state, parameters)
```

Arguments

t	A parameter used by the ode and passed by functions calling cloCoupleOde
state	Another parameter used by the ode and passed by functions calling cloCoupleOde
parameters	Another parameter used by the ode and passed by functions calling cloCoupleOde #'

Value

A list with the rates of change for each state variable.

crossCorPlots	<i>Produces cross-correlation plots of the observed state variable for lags of -+ 20 time steps for each dyad.</i>
---------------	--

Description

Produces cross-correlation plots of the observed state variable for lags of -+ 20 time steps for each dyad.

Usage

```
crossCorPlots(basedata, personId, dyadId, obs_name, time_name)
```

Arguments

basedata	A user provided dataframe.
personId	The name of the column in the dataframe that has the person-level identifier.
dyadId	The name of the column in the dataframe that has the dyad-level identifier.
obs_name	The name of the column in the dataframe that has the time-varying observable (e.g., the variable for which dynamics will be assessed).
time_name	The name of the column in the dataframe that indicates sequential temporal observations.

Value

Prints the plots to the screen.

Examples

```
data <- rties_ExampleDataShort
crossCorPlots(basedata=data, dyadId="couple", personId="person", obs_name="dial", time_name="time")
```

dataPrep	<i>Reformat a user-provided dataframe in a generic form appropriate for rties modeling</i>
----------	--

Description

The dataframe must be in a specific format and include several specific variables. See the "overview_data_prep" vignette for complete details on the necessary format and follow it closely if you'd like to avoid error messages. That vignette also includes information on how to structure the data if you have two variables within people (rather than two people within dyads) or have indistinguishable dyads.

Usage

```
dataPrep(
  basedata,
  dyadId,
  personId,
  obs_name,
  dist_name,
  time_name,
  time_lag = NULL
)
```

Arguments

basedata	A user-provided dataframe that includes all variables needed for an rties analysis.
dyadId	The name of the column in the dataframe that has the dyad-level identifier.
personId	The name of the column in the dataframe that has the person-level identifier.
obs_name	The name of the column in the dataframe that has the time-varying observable (e.g., the variable for which dynamics will be assessed).
dist_name	The name of the column in the dataframe that has a variable that distinguishes the partners (e.g., sex, mother/daughter, etc) that is numeric and scored 0/1.
time_name	The name of the column in the dataframe that indicates sequential temporal observations.
time_lag	An optional argument for the number of lags for the lagged observable. If a number is provided, the observed variable is lagged that amount. The other option is to use "absMaxCC". In this case the maximum cross-correlation is found for each dyad and the lag at which that occurs is used to lag their observed variables.

Value

The function returns a dataframe that has all the variables needed for modeling system dynamics, each renamed to a generic variable name, which are:

- id = person id
- dyad = dyad id
- obs = observed state variable
- dist1 = 0/1 variable where the 1's indicate the 1's in the original distinguishing variable
- time = the variable indicating temporal sequence
- dist0 = 0/1 variable where the 1's indicate the 0's in the original distinguishing variable
- obs_deTrend = the observed state variable with each person's linear trend removed
- p_ = all the same variables, but for a person's partner rather than themselves

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person", obs_name="dial",
  dist_name="female", time_name="time", time_lag=2)
head(newData)
```

dyadic

Produces plots for sysVarIn when sysVar is dyadic.

Description

Produces plots for sysVarIn when sysVar is dyadic.

Usage

```
dyadic(basedata, sysVar_name)
```

Arguments

basedata A dataframe created internally by the "sysVarInPlots" function.
 sysVar_name The name of the variable in the dataframe that contains the system variable.

Value

A plot with the profiles on the y-axis and the system variable on the x-axis

 estDerivs

Estimates first and second derivatives of an observed state variable

Description

This function makes use of 2 functions written by Steven Boker, "gllaWMatrix" and "gllaEmbed" which are available on his website, <http://people.virginia.edu/~smb3u/>. It fits a coupled oscillator model for each dyad at different combinations of the input parameters (tau, embeds) and returns the input values and period of oscillation that maximize the R² for each dyad. It also estimates first and second derivatives of the observed state variable for each person at the input values that maximize the R² for that dyad and returns a dataframe that contains them.

Usage

```
estDerivs(prepData, taus, embeds, delta, idConvention)
```

Arguments

prepData	A dataframe that was produced with the "dataPrep" function.
taus	A vector containing the values of tau to use. Tau indicates the number of time points to lag in the lagged data matrix (see Boker, S.M., Deboeck, P.R., Edler, C., & Keel, P.K. (2010). Generalized local linear approximation of derivatives from time series. In S.M. Chow & E. Ferrer (Eds.), <i>Statistical Methods for Modeling Human Dynamics: An Interdisciplinary Dialogue</i> (pp. 161-178). New York, NY: Taylor & Francis Group). The first derivative is estimated as the mean of the two adjacent slopes across that number of lags, e.g., if tau = 2 then the estimate of the first derivative at time = t is based on the mean of the slopes left and right of time t across 2 observations each. The second derivative is the difference in the two slopes with respect to time. Tau = 1 is sensitive to noise and increasing its value acts as smoothing.
embeds	A vector containing the values of embeds to use. Embeds indicates the number of columns in the lagged data matrix. The minimum = 3 for 2nd order derivatives and higher values increase smoothing.
delta	A value indicating the inter-observation interval. For example, if delta = 2, then every second observation is used in the estimation process.
idConvention	The value that was added to the dist1 ID number to get the dist2 ID number

Value

The function returns a list including: 1) "data" which is a dataframe containing first and second derivative estimates of an observed state variable, and 2) "fitTable" which shows the maximal R² achieved for each dyad for a coupled oscillator model, along with the associated tau, embed and estimated period of oscillation.

Examples

```

data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time")
taus <-c(2,3)
embeds <- c(3,4)
delta <- 1
derivs <- estDerivs(prepData=newData, taus=taus, embeds=embeds, delta=delta, idConvention=500)
head(derivs$fitTable)
summary(derivs$fitTable[,4]) # summary of R-square
summary(derivs$fitTable[,5]) # summary of period of oscillation

```

histAll

Histograms for all numeric variables in a dataframe.

Description

Useful for checking distributions to assess normality

Usage

```
histAll(basedata)
```

Arguments

basedata A user-provided dataframe.

Value

No return value. Prints plots to the console.

Examples

```

data <- rties_ExampleDataShort
vars <- c("reltime", "ambiv", "love", "conflict")
newData <- data[vars ]
histAll(newData)

```

indiv2profilesCat	<i>Produces plots for sysVarIn when sysVar is categorical and there are 2 profiles</i>
-------------------	--

Description

Produces plots for sysVarIn when sysVar is categorical and there are 2 profiles

Usage

```
indiv2profilesCat(testModel, sysVar0name, sysVar1name)
```

Arguments

testModel	The model object created by sysVarIn for the interaction model (e.g., sysVarInteract)
sysVar0name	The name created by sysVarInPlots referring to the system variable for partner-0.
sysVar1name	The name created by sysVarInPlots referring to the system variable for partner-1.

Value

A plot produced by the interactions package.

indiv2profilesCont	<i>Produces plots for sysVarIn when sysVar is continuous and there are 2 profiles</i>
--------------------	---

Description

Produces plots for sysVarIn when sysVar is continuous and there are 2 profiles

Usage

```
indiv2profilesCont(testModel, sysVar0name, sysVar1name)
```

Arguments

testModel	The model object created by sysVarIn for the interaction model (e.g., sysVarInteract)
sysVar0name	The name created by sysVarInPlots referring to the system variable for partner-0.
sysVar1name	The name created by sysVarInPlots referring to the system variable for partner-1.

Value

A plot produced by the interactions package.

indiv3profilesCat	<i>Produces plots for sysVarIn when sysVar is categorical and there are 3 profiles</i>
-------------------	--

Description

Produces plots for sysVarIn when sysVar is categorical and there are 3 profiles

Usage

```
indiv3profilesCat(basedata, testModel, sysVar0name, sysVar1name)
```

Arguments

basedata	A dataframe created internally by the "sysVarInPlots" function.
testModel	The model object created by sysVarIn for the interaction model (e.g., sysVarInteract)
sysVar0name	The name created by sysVarInPlots referring to the system variable for partner-0.
sysVar1name	The name created by sysVarInPlots referring to the system variable for partner-1.

Value

A list of 3 plots showing the simple slopes for each of the profiles.

indiv3profilesCont	<i>Produces plots for sysVarIn when sysVar is continuous and there are 3 profiles</i>
--------------------	---

Description

Produces plots for sysVarIn when sysVar is continuous and there are 3 profiles

Usage

```
indiv3profilesCont(prob, sysVar0name, sysVar1name)
```

Arguments

prob	A dataframe created internally by the "sysVarInPlots" function.
sysVar0name	The name created by sysVarInPlots referring to the system variable for partner-0.
sysVar1name	The name created by sysVarInPlots referring to the system variable for partner-1.

Value

A list of 3 plots showing the simple slopes for each of the profiles.

indiv4profilesCat	<i>Produces plots for sysVarIn when sysVar is categorical and there are 4 profiles</i>
-------------------	--

Description

Produces plots for sysVarIn when sysVar is categorical and there are 4 profiles

Usage

```
indiv4profilesCat(basedata, testModel, sysVar0name, sysVar1name)
```

Arguments

basedata	A dataframe created internally by the "sysVarInPlots" function.
testModel	The model object created by sysVarIn for the interaction model (e.g., sysVarInteract)
sysVar0name	The name created by sysVarInPlots referring to the system variable for partner-0.
sysVar1name	The name created by sysVarInPlots referring to the system variable for partner-1.

Value

A list of 4 plots showing the simple slopes for each of the profiles.

indiv4profilesCont	<i>Produces plots for sysVarIn when sysVar is continuous and there are 4 profiles</i>
--------------------	---

Description

Produces plots for sysVarIn when sysVar is continuous and there are 4 profiles

Usage

```
indiv4profilesCont(prob, sysVar0name, sysVar1name)
```

Arguments

prob	A dataframe created internally by the "sysVarInPlots" function.
sysVar0name	The name created by sysVarInPlots referring to the system variable for partner-0.
sysVar1name	The name created by sysVarInPlots referring to the system variable for partner-1.

Value

A list of 4 plots showing the simple slopes for each of the profiles.

indivClo	<i>Estimates either an uncoupled or coupled oscillator model for each dyad.</i>
----------	---

Description

Both models predict the second derivatives of the observed state variables (with linear trends removed). For the uncoupled oscillator, the predictors are each person's own observed state variables (again with linear trends removed), as well as each person's own first derivatives of the observed state variables (again with linear trends removed). For the coupled oscillator, the predictors are each person's own and partner's observed state variables (again with linear trends removed), as well as each person's own and partner's first derivatives of the observed state variables (again with linear trends removed).

Usage

```
indivClo(derivData, whichModel)
```

Arguments

derivData	A dataframe that was produced with the "estDerivs" function.
whichModel	Whether the model to be estimated is the "uncoupled" or "coupled" oscillator.

Value

The function returns a list including: 1) the adjusted R^2 for the model for each dyad (called "R2"), and 2) the parameter estimates for the model for each dyad (called "params", for use in either predicting, or being predicted by, the system variable).

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time")
taus <- c(2,3)
embeds <- c(3,4)
delta <- 1
derivs <- estDerivs(prepData=newData, taus=taus, embeds=embeds, delta=delta, idConvention=500)
clo <- indivClo(derivData=derivs$data, whichModel="coupled")
summary(clo$R2)
head(clo$params)
```

indivCloCompare	<i>Compares model fit for the uncoupled and coupled oscillator for each dyad's state trajectories using an R-square comparison.</i>
-----------------	---

Description

Fits an uncoupled and coupled oscillator model to each dyad's observed state variables and returns the adjusted R-squares, along with the difference between them (coupled - uncoupled, so positive values indicate better fit for the more complex model).

Usage

```
indivCloCompare(derivData)
```

Arguments

derivData	A dataframe that was produced with the "estDerivs" function. #' @examples <pre>data <- rties_ExampleDataShort newData <- dataPrep(basedata=data, dyadId="couple", personId="person", obs_name="dial", dist_name="female", time_name="time") taus <- c(2,3) embeds <- c(3,4) delta <- 1 derivs <- estDerivs(prepData=newData, taus=taus, embeds=embeds, delta=delta, idConvention=500) compare <- indivCloCompare(derivData=derivs\$data) summary(compare\$R2couple)</pre>
-----------	---

Value

The function returns a named list including: 1) the adjusted R² for the uncoupled model for each dyad (called "R2uncouple"), 2) the adjusted R² for the coupled model for each dyad (called "R2couple"), and 3) the difference between the R-squares for each dyad (coupled - uncoupled, called "R2dif").

indivCloPlots	<i>Produces plots of either an uncoupled or coupled oscillator model-predicted trajectories overlaid on raw data for each dyad.</i>
---------------	---

Description

The observed and CLO-model predicted state variables (with linear trends removed) are plotted for each dyad individually.

Usage

```

indivCloPlots(
  derivData,
  whichModel,
  idConvention,
  dist0name = NULL,
  dist1name = NULL,
  plot_obs_name = NULL,
  minMax = NULL,
  printPlots = T
)

```

Arguments

<code>derivData</code>	A dataframe that was produced with the "estDerivs" function.
<code>whichModel</code>	Whether the model to be estimated is the "uncoupled" or "coupled" oscillator.
<code>idConvention</code>	The number that was added to the dist0 partner to get the ID number for the dist1 partner.
<code>dist0name</code>	An optional name for the level-0 of the distinguishing variable (e.g., "Women"). Default is dist0.
<code>dist1name</code>	An optional name for the level-1 of the distinguishing variable (e.g., "Men"). Default is dist1.
<code>plot_obs_name</code>	An optional name for the observed state variables being plotted (e.g., "Emotional Experience"). Default is observed.
<code>minMax</code>	An optional vector with desired minimum and maximum quantiles to be used for setting the y-axis range on the plots, e.g., <code>minMax <- c(.1, .9)</code> would set the y-axis limits to the 10th and 90th percentiles of the observed state variables. Default is to use the minimum and maximum observed values of the state variables.
<code>printPlots</code>	If true (the default) plots are displayed on the screen.

Value

A list plots of the predicted values against the observed values for each dyad.

Examples

```
# See vignettes for examples.
```

indivInertCoord	<i>Estimates versions of the inertia-coordination model for each dyad.</i>
-----------------	--

Description

The user specifies which of 3 models are to be estimated. Each model predicts the observed state variables (with linear trends removed) from either: 1) Inertia only ("inert")- each person's intercept and each person's own observed state variable lagged at the amount specified during the dataPrep step (again with linear trends removed), 2) Coordination only ("coord")- each person's intercept and each person's partner's state variable lagged at the amount specified (again with linear trends removed), or 3) Full inertia-coordination model ("inertCoord") - each person's intercept, each person's own observed state variable lagged at the amount specified during the dataPrep step (again with linear trends removed), and each person's partner's state variable lagged at the amount specified (again with linear trends removed).

Usage

```
indivInertCoord(prepData, whichModel)
```

Arguments

prepData	A dataframe that was produced with the "dataPrep" function.
whichModel	Whether the model to be estimated is the inertia only model ("inert"), the coordination only model ("coord"), or the full inertia-coordination model ("inertCoord").

Value

The function returns a dataframe containing the parameter estimates, called "params", for use in the latent profile analysis.

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
ic <- indivInertCoord(prepData=newData, whichModel="inertCoord")
head(ic$params)
```

indivInertCoordCompare

Compares model fit for the inertia-only, coordination-only and full inertia-coordination model for each dyad's state trajectories using an R-square comparison.

Description

Fits inertia-only, coordination-only and full inertia-coordination models to each dyad's observed state variables and returns the adjusted R-squares, along with the differences between them, so positive values indicate better fit for the first model in the comparison. The 3 comparisons are inertia minus coordination, full model minus inertia, and full model minus coordination.

Usage

```
indivInertCoordCompare(prepData)
```

Arguments

prepData A dataframe that was produced with the "dataPrep" function.

Value

The function returns a named list including: 1) the adjusted R^2 for the inertia model for each dyad (called "R2inert"), 2) the adjusted R^2 for the coordination model for each dyad (called "R2coord"), 3) the adjusted R^2 for the full inertia-coordination model for each dyad (called "R2inertCoord"), 4) the difference between the R-squares for each dyad for inertia minus coordination (called "R2dif_IC_I"), 5) the difference for the full model minus inertia (called "R2dif_IC_I"), and 6) the difference for the full model minus coordination (called "R2dif_IC_C")

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
compare <- indivInertCoordCompare(prepData=newData)
summary(compare$R2inert)
summary(compare$R2coord)
summary(compare$R2inertCoord)
summary(compare$R2dif_IC_I)
```

indivInertCoordPlots *Produces plots of the inertia-coordination model-predicted trajectories overlaid on raw data for each dyad.*

Description

The observed state variables (with linear trends removed) are predicted from one of the 3 versions of the inertia-coordination model (inertia only, "inert"; coordination only, "coord"; full inertia-coordination, "inertCoord") for each dyad individually. The predicted trajectories are plotted overlaid on the observed trajectories.

Usage

```
indivInertCoordPlots(
  prepData,
  whichModel,
  dist0name = NULL,
  dist1name = NULL,
  plot_obs_name = NULL,
  minMax = NULL,
  printPlots = T
)
```

Arguments

prepData	A dataframe that was produced with the "dataPrep" function.
whichModel	Whether the model to be estimated is the inertia only model ("inert"), the coordination only model ("coord"), or the full inertia-coordination model ("inertCoord").
dist0name	An optional name for the level-0 of the distinguishing variable to appear on plots (e.g., "Women").
dist1name	An optional name for the level-1 of the distinguishing variable to appear on plots (e.g., "Men").
plot_obs_name	An optional name for the observed state variable to appear on plots (e.g., "Emotional Experience").
minMax	An optional vector with desired minimum and maximum quantiles to be used for setting the y-axis range on the plots, e.g., minMax <- c(.1, .9) would set the y-axis limits to the 10th and 90th percentiles of the observed state variables. If not provided, the default is to use the minimum and maximum observed values of the state variables.
printPlots	If true (the default) plots are displayed on the screen.

Value

A list with the plots of the predicted values against the observed values for each dyad.

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
temp <- newData[newData$dyad < 5, ]
plots <- indivInertCoordPlots(prepData=temp, whichModel="inertCoord")
```

inertCoordPlotTraj	<i>Plots the bivariate state variables' model-predicted temporal trajectories for each latent profile of inertia-coordination parameters.</i>
--------------------	---

Description

Produces sets of prototypical example plots of the state variables' predicted temporal trajectories for each latent profile obtained based on the inertia-coordination parameters. The plots are produced by using the inertia-coordination parameters to predict temporal trajectories, with random noise added at each temporal step.

Usage

```
inertCoordPlotTraj(
  prepData,
  paramEst,
  n_profiles,
  dist0name = NULL,
  dist1name = NULL,
  plot_obs_name = NULL,
  minMax = NULL,
  time_length = NULL,
  numPlots = NULL,
  seed = NULL,
  printPlots = T
)
```

Arguments

prepData	A dataframe that was produced with the "dataPrep" function.
paramEst	A dataframe created by indivInertCoord containing the inertia-coordination parameter estimates for each dyad.
n_profiles	The number of latent profiles.
dist0name	An optional name for the level-0 of the distinguishing variable (e.g., "Women"). Default is dist0.
dist1name	An optional name for the level-1 of the distinguishing variable (e.g., "Men"). Default is dist1

plot_obs_name	An optional name for the observed state variable to appear on plots (e.g., "Emotional Experience").
minMax	An optional vector with desired minimum and maximum quantiles to be used for setting the y-axis range on the plots, e.g., <code>minMax <- c(.1, .9)</code> would set the y-axis limits to the 10th and 90th percentiles of the observed state variables. If not provided, the default is to use the minimum and maximum observed values of the state variables.
time_length	An optional value specifying how many time points to plot across. Default is the 75th percentile for the time variable.
numPlots	An optional value controlling how many random examples of each profile are produced. Default is 3.
seed	An optional integer argument that sets the seed of R's random number generator to create reproducible trajectories. If used, the "numPlots" can be set to one - otherwise each plot is replicated 3 times.
printPlots	If true (the default) plots are displayed on the screen.

Value

A list with the plots of predicted trajectories for each dyad.

Examples

```
# See vignettes for examples.
```

inertCoordResids	<i>Produces histograms of the residuals from the inertia-coordination model for each dyad.</i>
------------------	--

Description

Produces histograms of the residuals from the inertia-coordination model for each dyad.

Usage

```
inertCoordResids(prepData, whichModel, printPlots = T)
```

Arguments

prepData	A dataframe that was produced with the "dataPrep" function.
whichModel	Whether the model to be estimated is the inertia only model ("inert"), the coordination only model ("coord"), or the full inertia-coordination model ("inertCoord").
printPlots	If true (the default) plots are displayed on the screen.

Value

A list with the histograms of the residuals for each dyad.

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
temp <- newData[newData$dyad < 5, ]
residPlots <- inertCoordResids(prepData=temp, whichModel="inertCoord")
```

inspectProfiles	<i>Provides information to help decide how many profiles to use for subsequent rties analyses.</i>
-----------------	--

Description

The function prints out the number of dyads in each profile for a specified number of profiles. It also prints out: 1) a figure showing the best clustering solution as indicated by BIC (e.g., the observed data separated into clusters, produced by mclust), 2) a line plot showing the content of the best solution (e.g., the mean parameter estimates for each profile) and 3) prototypical model-predicted trajectories for each profile. For the inertia-coordination model, it produces sets of prototypical examples by using the inertia-coordination parameters to predict temporal trajectories, with random noise added at each temporal step. This process is required because the inertia-coordination model only represents local dynamics and predictions bear no resemblance to observed variables without the addition of noise. An optional argument, "seed" sets the seed for the random number generator, so you can get the same plots each time. If the "seed" argument is used, then only one plot per profile is produced. For the coupled-oscillator, this step is not necessary and one prototypical trajectory is plotted for each profile.

Usage

```
inspectProfiles(
  whichModel,
  prepData,
  paramEst,
  n_profiles,
  dist0name = NULL,
  dist1name = NULL,
  plot_obs_name = NULL,
  minMax = NULL,
  time_length = NULL,
  numPlots = NULL,
  seed = NULL
)
```


Arguments

whichModel	The name of the model that is being investigated (e.g., "inertCoord" or "clo")
prepData	A dataframe that was produced with the "dataPrep" function.
paramEst	A dataframe created by either indivInertCoord or indivClo containing the parameter estimates for each dyad.
n_profiles	The number of latent profiles.
dist0name	An optional name for the level-0 of the distinguishing variable (e.g., "Women"). Default is dist0.
dist1name	An optional name for the level-1 of the distinguishing variable (e.g., "Men"). Default is dist1
plot_obs_name	An optional name for the observed state variable to appear on plots (e.g., "Emotional Experience").
minMax	An optional vector with desired minimum and maximum quantiles to be used for setting the y-axis range on the plots, e.g., minMax <- c(.1, .9) would set the y-axis limits to the 10th and 90th percentiles of the observed state variables. If not provided, the default is to use the minimum and maximum observed values of the state variables.
time_length	An optional value specifying how many time points to plot across. Default is the 75th percentile for the time variable.
numPlots	Only relevant for the inertCoord model. An optional value controlling how many random examples of each profile are produced. Default is 3.
seed	Only relevant for the inertCoord model. An optional integer argument that sets the seed of R's random number generator to create reproducible trajectories. If used, the "numPlots" can be set to one - otherwise each plot is replicated 3 times.

Value

A dataframe called "profileData" that contains the profile classification for each dyad.

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time")
taus <-c(2,3)
embeds <- c(3,4)
delta <- 1
derivs <- estDerivs(prepData=newData, taus=taus, embeds=embeds, delta=delta, idConvention=500)
clo <- indivClo(derivData=derivs$data, whichModel="coupled")
profiles <- inspectProfiles(whichModel="clo", prepData=newData, paramEst=clo$params, n_profiles=2)
head(profiles)
```

makeCrossCorr	<i>Calculates cross-correlations for a given variable and returns a dataframe with the largest absolute cross-correlation and its lag added for each dyad (e.g., it returns either the most negative or most positive cross-correlation, whichever is larger in absolute terms).</i>
---------------	--

Description

Calculates cross-correlations for a given variable and returns a dataframe with the largest absolute cross-correlation and its lag added for each dyad (e.g., it returns either the most negative or most positive cross-correlation, whichever is larger in absolute terms).

Usage

```
makeCrossCorr(basedata, dyadId, personId, obs_name, dist_name)
```

Arguments

basedata	The original dataframe provided by the user that includes all variables needed for an rties analysis, including potential system and control variables, etc.
dyadId	The name of the column in the dataframe that has the couple-level identifier.
personId	The name of the column in the dataframe that has the person-level identifier.
obs_name	The name of the column in the dataframe that has the time-varying observable (e.g., the variable for which dynamics will be assessed).
dist_name	The name of the column in the dataframe that has a variable that distinguishes the partners (e.g., sex, mother/daughter, etc) that is numeric and scored 0/1.

Value

A cross-sectional version of the original dataframe with maximal absolute-value cross-correlations and their lags added for each dyad.

Examples

```
data <- rties_ExampleDataShort
newData <- makeCrossCorr(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female")
head(newData)
```

makeDist	<i>Create a distinguishing variable (called "dist") for non-distinguishable dyads by assigning the partner who is lower on a chosen variable a 0 and the partner who is higher on the variable a 1.</i>
----------	---

Description

Create a distinguishing variable (called "dist") for non-distinguishable dyads by assigning the partner who is lower on a chosen variable a 0 and the partner who is higher on the variable a 1.

Usage

```
makeDist(basedata, dyadId, personId, time_name, dist_name)
```

Arguments

basedata	A user-provided dataframe.
dyadId	The name of the column in the dataframe that has the couple-level identifier.
personId	The name of the column in the dataframe that has the person-level identifier.
time_name	The name of the column in the dataframe that indicates sequential temporal observations.
dist_name	The name of the column in the dataframe that holds the variable to use for distinguishing the partners. For example, if "influence" was the variable, for each dyad the partner scoring lower on "influence" would be given a score of 0 on "dist" and the partner scoring higher on "influence" would be given a score of 1 on "dist"

Value

The function returns the original dataframe with an additional variable, called "dist" that distinguishes between partners based on the user-specified variable indicated by "dist_name"

Examples

```
data <- rties_ExampleDataShort
newData <- makeDist(basedata=data, dyadId="couple", personId="person",
  time_name="time", dist_name="relstress")
summary(newData$dist)
```

makeFullData	<i>Combines profile membership data from the latent profile analysis with other data for using the profile membership to predict and be predicted by the system variable.</i>
--------------	---

Description

Combines profile membership data from the latent profile analysis with other data for using the profile membership to predict and be predicted by the system variable.

Usage

```
makeFullData(basedata, dyadId, personId, dist_name, lpaData, params)
```

Arguments

basedata	The original dataframe provided by the user that includes all variables needed for an rties analysis, including potential system and control variables, etc.
dyadId	The name of the column in the dataframe that has the couple-level identifier.
personId	The name of the column in the dataframe that has the person-level identifier.
dist_name	The name of the column in the dataframe that has a variable that distinguishes the partners (e.g., sex, mother/daughter, etc) that is numeric and scored 0/1.
lpaData	The object returned by the "inspectProfiles" function
params	The list called "params" returned by one of the "indiv" functions (e.g., indivInertCoord or indivClo)

Value

A dataframe that contains all variables needed for using the profiles to predict, or be predicted by, the system variable.

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
ic <- indivInertCoord(prepData=newData, whichModel="inertCoord")
profiles <- inspectProfiles(whichModel="inertCoord", prepData=newData,
  paramEst=ic$params, n_profiles=2)
fullData <- makeFullData(basedata=data, dyadId="couple", personId="person",
  dist_name="female", lpaData=profiles, params=ic$params)
head(fullData)
```

Max_Min_CCF_Signed *A helper function for makeCrossCorr*

Description

A helper function for makeCrossCorr

Usage

```
Max_Min_CCF_Signed(a, b)
```

Arguments

a First time-series used in the cross-correlation
b Second time-series used in the cross-correlation

Value

A list of maximum absolute value cross-correlations and the lag at which they occurred.

plotDataByProfile *Plots of de-trended observed variable over time, with dyads separated into groups based on LPA profile membership.*

Description

Plots of de-trended observed variable over time, with dyads separated into groups based on LPA profile membership.

Usage

```
plotDataByProfile(  
  prepData,  
  fullData,  
  n_profiles,  
  dist0name = NULL,  
  dist1name = NULL,  
  plot_obs_name = NULL,  
  printPlots = T  
)
```

Arguments

prepData	A dataframe created by the dataPrep function.
fullData	A dataframe created by the makeFullData function.
n_profiles	The number of profiles that were estimated.
dist0name	An optional name for the level-0 of the distinguishing variable (e.g., "Women"). Default is dist0.
dist1name	An optional name for the level-1 of the distinguishing variable (e.g., "Men"). Default is dist1.
plot_obs_name	An optional name for the observed state variable to appear on plots (e.g., "Emotional Experience").
printPlots	If true (the default) plots are displayed on the screen.

Value

A list of plots.

Examples

```
# See vignettes for examples.
```

plotRaw	<i>Plots of observed variable over time by dyad.</i>
---------	--

Description

Produces plots of the observed variable for each dyad over time to check for data errors, etc.

Usage

```
plotRaw(
  basedata,
  dyadId,
  obs_name,
  dist_name,
  time_name,
  dist0name = NULL,
  dist1name = NULL,
  plot_obs_name = NULL,
  printPlots = T
)
```

Arguments

basedata	A user provided dataframe.
dyadId	The name of the column in the dataframe that has the dyad-level identifier.
obs_name	The name of the column in the dataframe that has the time-varying observable (e.g., the variable for which dynamics will be assessed).
dist_name	The name of the column in the dataframe that has a variable that distinguishes the partners (e.g., sex, mother/daughter, etc) that is numeric and scored 0/1.
time_name	The name of the column in the dataframe that indicates sequential temporal observations.
dist0name	An optional name for the level-0 of the distinguishing variable to appear on plots (e.g., "Women").
dist1name	An optional name for the level-1 of the distinguishing variable to appear on plots (e.g., "Men").
plot_obs_name	An optional name for the observed state variable to appear on plots (e.g., "Emotional Experience").
printPlots	If true (the default) plots are displayed on the screen.

Value

A list of plots.

Examples

```
data <- rties_ExampleDataShort
plotRaw(basedata=data, dyad="couple", obs_name="dial", dist_name="female", time_name="time")
```

removeDyads	<i>Remove data for specified dyads from a dataframe</i>
-------------	---

Description

Useful for cleaning data if some dyads have extensive missing or otherwise problematic data.

Usage

```
removeDyads(basedata, dyads, dyadId)
```

Arguments

basedata	A user provided dataframe.
dyads	A vector of dyad IDs to remove.
dyadId	The variable in the dataframe specifying dyad ID.

Value

A dataframe with the data for the specified dyads removed.

Examples

```
data <- rties_ExampleDataShort
dyads <- c(3, 12)
newData <- removeDyads(basedata=data, dyads=dyads, dyadId="couple")
```

rties_ExampleDataFull *Data for examples in the vignettes.*

Description

A dataset containing variables for the examples in the vignettes.

Usage

```
rties_ExampleDataFull
```

Format

An object of class `data.frame` with 78858 rows and 19 columns.

rties_ExampleDataShort
Data for the function examples.

Description

A dataset containing variables for the examples of the functions.

Usage

```
rties_ExampleDataShort
```

Format

An object of class `data.frame` with 17884 rows and 19 columns.

 rties_ExampleData_Demo

Data for demonstrating rties models.

Description

A dataset containing a minimal set of variables for demonstrating rties analyses.

Usage

```
rties_ExampleData_Demo
```

Format

An object of class `data.frame` with 12 rows and 6 columns.

<code>sysVarIn</code>	<i>Provides results for predicting couples' latent profile membership from the system variable.</i>
-----------------------	---

Description

If there are 2 profiles, then glm binomial regression models are used. If there are more than 2 profiles then multinomial regression is used (from the `nnet` package). The system variable can be either dyadic (`sysVarType = "dyadic"`), where both partners have the same score (e.g., relationship length) or individual (`sysVarType = "indiv"`), where the partners can have different scores (e.g., age). For dyadic system variables, a couple's shared score is the only predictor of their profile membership (called "sysVar"). For individual system variables, two models are tested, one with the main effects of both partner's system variable ("sysVarMain") and one with the main effects and their interaction ("sysVarInteract"). In both cases an intercept-only model is included as a comparison point (called "base"). The function returns a list of the full model results.

Usage

```
sysVarIn(fullData, sysVar_name, sysVarType, n_profiles)
```

Arguments

<code>fullData</code>	A dataframe created by the <code>makeFullData</code> function.
<code>sysVar_name</code>	The name of the variable in the dataframe that contains the system variable to be predicted by profile membership.
<code>sysVarType</code>	Whether the system variable is "dyadic", which means both partners have the same score, or "indiv" which means the partners can have different scores
<code>n_profiles</code>	The number of latent profiles.

Value

A list including the glm or multinom objects containing the full results for each model (called "models").

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
ic <- indivInertCoord(prepData=newData, whichModel="inertCoord")
profiles <- inspectProfiles(whichModel="inertCoord", prepData=newData,
  paramEst=ic$params, n_profiles=2)
fullData <- makeFullData(basedata=data, dyadId="couple", personId="person",
  dist_name="female", lpaData=profiles, params=ic$params)
sysIn <- sysVarIn(fullData=fullData, sysVar_name="conflict", sysVarType="indiv", n_profiles=2)
summary(sysIn$models$sysVarMain)
```

 sysVarInPlots

Produces plots for interpreting the results from sysVarIn.

Description

Produces plots for interpreting the results from sysVarIn.

Usage

```
sysVarInPlots(
  fullData,
  sysVar_name,
  sysVarType,
  n_profiles,
  testModel = NULL,
  dist0name = NULL,
  dist1name = NULL,
  printPlots = T
)
```

Arguments

fullData	A dataframe created by the "makeFullData" function.
sysVar_name	The name of the variable in the dataframe that contains the system variable.
sysVarType	Whether the system variable is "dyadic", which means both partners have the same score, or "indiv" which means the partners can have different scores
n_profiles	The number of latent profiles.

testModel	The name of the model that is being interpreted (e.g., sysIn\$models\$sysVarInteract). Only needed when the system variable is "indiv" (e.g., individual scores for each partner)
dist0name	An optional name for the level-0 of the distinguishing variable (e.g., "Women"). Default is dist0.
dist1name	An optional name for the level-1 of the distinguishing variable (e.g., "Men"). Default is dist1
printPlots	If true (the default) plots are displayed on the screen.

Value

Single plots or a list of plots (depending on the model that is being interpreted).

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
ic <- indivInertCoord(prepData=newData, whichModel="inertCoord")
profiles <- inspectProfiles(whichModel="inertCoord", prepData=newData,
  paramEst=ic$params, n_profiles=2)
fullData <- makeFullData(basedata=data, dyadId="couple", personId="person",
  dist_name="female", lpaData=profiles, params=ic$params)
sysIn <- sysVarIn(fullData=fullData, sysVar_name="conflict", sysVarType="indiv", n_profiles=2)
sysVarInPlots(fullData=fullData, sysVar_name="conflict", sysVarType="indiv",
  n_profiles=2, testModel=sysIn$models$sysVarInteract)
```

sysVarInResults	<i>Produces results from sysVarIn.</i>
-----------------	--

Description

Produces results from sysVarIn.

Usage

```
sysVarInResults(baseModel, testModel, n_profiles)
```

Arguments

baseModel	The name of the model that was produced by sysVarIn to be used as the null model for comparison (e.g., sysIn\$models\$base).
testModel	The name of the model that was produced by sysVarIn that you want results for (e.g., sysIn\$models\$sysVarMain or sysIn\$models\$sysVarInteract).

`n_profiles` The number of latent profiles. #' @examples data <- rties_ExampleDataShort
 newData <- dataPrep(basedata=data, dyadId="couple", personId="person", obs_name="dial",
 dist_name="female", time_name="time", time_lag=2) ic <- indivInertCoord(prepData=newData,
 whichModel="inertCoord") profiles <- inspectProfiles(whichModel="inertCoord",
 prepData=newData, paramEst=ic\$params, n_profiles=2) fullData <- makeFull-
 Data(basedata=data, dyadId="couple", personId="person", dist_name="female",
 lpaData=profiles, params=ic\$params) sysIn <- sysVarIn(fullData=fullData, sys-
 Var_name="conflict", sysVarType="indiv", n_profiles=2) sysVarInResults(baseModel=sysIn\$models\$bas-
 testModel=sysIn\$models\$sysVarMain, n_profiles=2)

Value

A list of results including a comparison of the test model to the null (either a LRT or Chisquare test depending on the model), a summary of the parameter estimates, exponentiated parameter estimates (e.g., odds ratios), and p values for the parameter estimates.

<code>sysVarOut</code>	<i>Provides results for predicting the system variable from the latent profiles of the dynamic parameters.</i>
------------------------	--

Description

The system variable can be either dyadic (`sysVarType = "dyadic"`), where both partners have the same score (e.g., relationship length) or individual (`sysVarType = "indiv"`), where the partners can have different scores (e.g., age). For dyadic system variables, the only predictor is profile membership and the model is a regular regression model since all variables are at the level of the dyad. If the system variable is individual then the model is a random-intercept dyadic model and 3 models are estimated: 1) the main effect of profile membership, 2) main effects of profile membership and the distinguishing variable, and 3) the interaction of profile membership and the distinguishing variable. If the system variable is not normally distributed, any of the generalized linear models supported by `glm` (for dyadic system variables) or `glmer` (for individual system variables) are available by specifying the "family" distribution.

Usage

```
sysVarOut(  
  fullData,  
  sysVar_name,  
  sysVarType,  
  dist0name = NULL,  
  dist1name = NULL,  
  family = NULL  
)
```

Arguments

fullData	A dataframe created by the "makeFullData" function.
sysVar_name	The name of the variable in the dataframe that contains the system variable to be predicted by profile membership.
sysVarType	Whether the system variable is "dyadic", which means both partners have the same score, or "indiv" which means the partners can have different scores
dist0name	An optional name for the level-0 of the distinguishing variable (e.g., "Women"). Default is dist0.
dist1name	An optional name for the level-1 of the distinguishing variable (e.g., "Men"). Default is dist1
family	An optional argument specifying the error distribution and link function to be used in the model. Any of the "family" options supported by glm (for dyadic system variables) or glmer (for individual system variables) are available. Default is gaussian.

Value

For normally distributed system variables, the function returns a list including the lm or lme objects containing the full results for each model (called "models"). Similarly, for non-normal system variables, the function returns a list of the glm or glmer objects containing the full results for the models.

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
ic <- indivInertCoord(prepData=newData, whichModel="inertCoord")
profiles <- inspectProfiles(whichModel="inertCoord", prepData=newData,
  paramEst=ic$params, n_profiles=2)
fullData <- makeFullData(basedata=data, dyadId="couple", personId="person",
  dist_name="female", lpaData=profiles, params=ic$params)
sysOut <- sysVarOut(fullData=fullData, sysVar_name="conflict", sysVarType="indiv")
summary(sysOut$models$profilePlusDist)
```

 sysVarOutPlots

Produces plots for interpreting the results from sysVarIn.

Description

Produces plots for interpreting the results from sysVarIn.

Usage

```

sysVarOutPlots(
  fullData,
  sysVar_name,
  sysVarType,
  testModel,
  dist0name = NULL,
  dist1name = NULL,
  binomial = F
)

```

Arguments

fullData	A dataframe created by the "makeFullData" function.
sysVar_name	The name of the variable in the dataframe that contains the system variable.
sysVarType	Whether the system variable is "dyadic", which means both partners have the same score, or "indiv" which means the partners can have different scores
testModel	The name of the model that is being interpreted (e.g., sysIn\$models\$sysVarInteract).
dist0name	An optional name for the level-0 of the distinguishing variable (e.g., "Women"). Default is dist0.
dist1name	An optional name for the level-1 of the distinguishing variable (e.g., "Men"). Default is dist1
binomial	Whether the system variable is binomial. Default is false.

Value

Single plots or a list of plots (depending on the model that is being interpreted).

Examples

```
# See vignettes for examples.
```

sysVarOutResults	<i>Produces results from sysVarOut.</i>
------------------	---

Description

Produces results from sysVarOut.

Usage

```
sysVarOutResults(baseModel, testModel, Gaussian = TRUE)
```

Arguments

baseModel	The name of the model that was produced by sysVarOut to be used as the null model for comparison (e.g., sysOut\$models\$base).
testModel	The name of the model that was produced by sysVarOut that you want results for (e.g., sysOut\$models\$profile, sysOut\$models\$profilePlusDist, sysOut\$models\$profileByDist).
Gaussian	Whether the system variable is Gaussian. Default is true.

Value

A list of results including an LRT comparison of the test model to the null, an omnibus anova test for the parameters in the model (this is identical to the LRT test for Gaussian dyadic system variables), a summary of the parameter estimates, and exponentiated parameter estimates (e.g., odds ratios) if Gaussian = FALSE.

Examples

```
data <- rties_ExampleDataShort
newData <- dataPrep(basedata=data, dyadId="couple", personId="person",
  obs_name="dial", dist_name="female", time_name="time", time_lag=2)
ic <- indivInertCoord(prepData=newData, whichModel="inertCoord")
profiles <- inspectProfiles(whichModel="inertCoord", prepData=newData,
  paramEst=ic$params, n_profiles=2)
fullData <- makeFullData(basedata=data, dyadId="couple", personId="person",
  dist_name="female", lpaData=profiles, params=ic$params)
sysOut <- sysVarOut(fullData=fullData, sysVar_name="conflict", sysVarType="indiv")
sysVarOutResults(baseModel=sysOut$models$base, testModel=sysOut$models$profileByDist)
```

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