

# Package ‘pompom’

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**Type** Package

**Title** Person-Oriented Method and Perturbation on the Model

**Version** 0.2.1

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**Description** An implementation of a hybrid method of person-oriented method and perturbation on the model. Pompom is the initials of the two methods. The hybrid method will provide a multivariate intraindividual variability metric (iRAM). The person-oriented method used in this package refers to uSEM (unified structural equation modeling, see Kim et al., 2007, Gates et al., 2010 and Gates et al., 2012 for details). Perturbation on the model was conducted according to impulse response analysis introduced in Lutkepohl (2007).

Kim, J., Zhu, W., Chang, L., Bentler, P. M., & Ernst, T. (2007) <doi:10.1002/hbm.20259>.

Gates, K. M., Mole-

naar, P. C. M., Hillary, F. G., Ram, N., & Rovine, M. J. (2010) <doi:10.1016/j.neuroimage.2009.12.117>.

Gates, K. M., & Molenaar, P. C. M. (2012) <doi:10.1016/j.neuroimage.2012.06.026>.

Lutkepohl, H. (2007, ISBN:3540262393).

**License** GPL-2

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**Depends** R (>= 3.0.0)

**Imports** lavaan (>= 0.5-23.1097), ggplot2 (>= 2.2.1), reshape2 (>= 1.4.2), qgraph, utils

**Suggests** knitr, rmarkdown, testthat

**VignetteBuilder** knitr

**NeedsCompilation** no

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bootstrap_iRAM_2node	<i>Bootstrapped iRAM (including replications of iRAM and corresponding time profiles) for the bivariate time-series (simts2node)</i>
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### Description

Bootstrapped iRAM (including replications of iRAM and corresponding time profiles) for the bivariate time-series (simts2node)

### Usage

```
bootstrap_iRAM_2node
```

### Format

An object of class `list` of length 5.

### Details

Data bootstrapped from the estimated three-node network structure with 200 replications.

**Examples**

```
bootstrap_iRAM_2node$mean # mean of bootstrapped iRAM
bootstrap_iRAM_2node$upper # Upper bound of confidence interval of bootstrapped iRAM
bootstrap_iRAM_2node$lower # lower bound of confidence interval of bootstrapped iRAM
bootstrap_iRAM_2node$time.profile.data # time profiles generated from the bootstrapped beta matrices
bootstrap_iRAM_2node$recovery.time.reps # iRAMs generated from the bootstrapped beta matrices
```

---

```
bootstrap_iRAM_3node Bootstrapped iRAM (including replications of iRAM and corresponding time profiles) for the 3-variate time-series (simts)
```

---

**Description**

Bootstrapped iRAM (including replications of iRAM and corresponding time profiles) for the 3-variate time-series (simts)

**Usage**

```
bootstrap_iRAM_3node
```

**Format**

An object of class list of length 5.

**Details**

Data bootstrapped from the estimated three-node network structure with 200 replications.

**Examples**

```
bootstrap_iRAM_3node$mean # mean of bootstrapped iRAM
bootstrap_iRAM_3node$upper # Upper bound of confidence interval of bootstrapped iRAM
bootstrap_iRAM_3node$lower # lower bound of confidence interval of bootstrapped iRAM
bootstrap_iRAM_3node$time.profile.data # time profiles generated from the bootstrapped beta matrices
bootstrap_iRAM_3node$recovery.time.reps # iRAMs generated from the bootstrapped beta matrices
```

---

**iRAM***Generate iRAM (impulse response analysis metric) from model fit.*

---

**Description**

Generate iRAM (impulse response analysis metric) from model fit.

**Usage**

```
iRAM(  
  model.fit,  
  beta,  
  var.number,  
  lag.order = 1,  
  threshold = 0.01,  
  boot = FALSE,  
  replication = 200,  
  steps = 100  
)
```

**Arguments**

<code>model.fit</code>	model fit object generated by lavaan
<code>beta</code>	beta matrix for a point estimate
<code>var.number</code>	number of variables in the time series
<code>lag.order</code>	lag order of the model to be fit
<code>threshold</code>	threshold of calculation of recovery time (duration of perturbation), default value is 0.01
<code>boot</code>	to bootstrap, default value is FALSE
<code>replication</code>	number of replication of bootstrap, default value is 200
<code>steps</code>	number of steps of impulse response analysis, default value is 100

**Value**

iRAM matrix. Rows represent where the orthogonal impulse was given, and columns represent the response. Dimension is `var.number` by `var.number`.

**References**

Lütkepohl, H. (2007). New introduction to multiple time-series analysis. Berlin: Springer.

**Examples**

```
boot.iRAM <- iRAM(model.fit = usemodelfit,
  beta = NULL,
  var.number = 3,
  lag.order = 1,
  threshold = 0.01,
  boot = TRUE,
  replication = 200,
  steps = 100
)
boot.iRAM$mean
```

---

iRAM_equilibrium	<i>Generate iRAM (impulse response analysis metric) in the equilibrium form.</i>
------------------	--

---

**Description**

Generate iRAM (impulse response analysis metric) in the equilibrium form.

**Usage**

```
iRAM_equilibrium(beta.matrix, var.number, lag.order)
```

**Arguments**

beta.matrix	beta matrix for a point estimate
var.number	number of variables in the time series
lag.order	lag order of the model to be fit

**Value**

a list of equilibria. First numeric number in the variable name indicate where the impulse was given, and the second numeric number indicate the response, e.g., e12 indicates equilibrium of node 2 when node 1 is given an impulse.

**Examples**

```
iRAM_evalue <- iRAM_equilibrium(beta.matrix = true_beta_3node,
  var.number = 3,
  lag.order = 1
)
iRAM_evalue
```

---

model_summary	<i>Provide model summary.</i>
---------------	-------------------------------

---

**Description**

Provide model summary.

**Usage**

```
model_summary(model.fit, var.number, lag.order)
```

**Arguments**

model.fit	model fit object generated by lavaan
var.number	number of variables in the time-series
lag.order	lag order of model

**Details**

Model fit criteria: 3 out of 4 rule, meaning 3 out of 4 criteria should be satisfied, including CFI and TLI should be greater than 0.95, RMSEA and SRMR should be less than 0.08.

**Value**

beta matrix estimates  
matrix of standard error of beta  
matrix of psi estimates  
fit statistics CFI  
fit statistics TLI  
fit statistics RMSEA  
fit statistics SRMR

**Examples**

```
mdl <- model_summary(model.fit = usemodelfit,  
                     var.number = 3,  
                     lag.order = 1)  
  
mdl$beta  
mdl$beta.se  
mdl$psi  
mdl$cfi
```



---

plot\_integrated\_time\_profile

*Plot the time profiles in the integrated form*

---

### **Description**

Plot the time profiles in the integrated form

### **Usage**

```
plot_integrated_time_profile(beta.matrix, var.number, lag.order = 1)
```

### **Arguments**

beta.matrix	matrix of temporal relations, containing both lag-1 and contemporaneous
var.number	number of variables in the time series
lag.order	lag order of the model to be fit

### **Examples**

```
plot_integrated_time_profile(beta.matrix = true_beta_3node,  
                             var.number = 3,  
                             lag.order = 1)
```

---

plot\_iRAM\_dist

*Plot distribution of recovery time based on bootstrapped version of iRAM*

---

### **Description**

Plot distribution of recovery time based on bootstrapped version of iRAM

### **Usage**

```
plot_iRAM_dist(recovery.time.reps)
```

### **Arguments**

recovery.time.reps	bootstrapped version of recovery time
--------------------	---------------------------------------

### Examples

```
plot_iRAM_dist(bootstrap_iRAM_3node$recovery.time.reps)
```

---

plot\_network\_graph      *Plot the network graph*

---

### Description

Plot the network graph

### Usage

```
plot_network_graph(beta, var.number)
```

### Arguments

beta                    matrix of temporal relations, cotaining both lag-1 and contemporaneous  
var.number            number of variables in the time series

### Examples

```
plot_network_graph(beta = true_beta_3node,  
                    var.number = 3)
```

---

plot\_time\_profile      *Plot time profiles given a time-series generated by impulse response analysis*

---

### Description

Plot time profiles given a time-series generated by impulse response analysis

### Usage

```
plot_time_profile(time.series.data, var.number, threshold = 0.01, xupper = 20)
```

**Arguments**

<code>time.series.data</code>	data of impulse response in long format
<code>var.number</code>	number of variables in the time-series
<code>threshold</code>	threshold of asymptote of equilibrium
<code>xupper</code>	upper limit of x-axis

**Examples**

```
plot_time_profile(time.series.data = bootstrap_iRAM_2node$time.profile.data,  
                 var.number = 2,  
                 threshold= .01,  
                 xupper = 20)
```

---

`simts_2node`*Simulated bivariate time-series data*

---

**Description**

Simulated bivariate time-series data

**Usage**

```
simts_2node
```

**Format**

An object of class `data.frame` with 200 rows and 2 columns.

**Details**

Data simulated from a given three-node network structure with 200 measurements. Network structure is shown in the dataset `true.beta`. Process noise has mean of 0 and SD .1.

**Examples**

```
data(simts_2node)
```

---

simts_3node	<i>Simulated 3-variate time-series data</i>
-------------	---

---

**Description**

Simulated 3-variate time-series data

**Usage**

```
simts_3node
```

**Format**

An object of class `data.frame` with 100 rows and 3 columns.

**Details**

Data simulated from a given three-node network structure with 200 measurements. Network structure is shown in the dataset `true.beta`. Process noise has mean of 0 and SD .1.

**Examples**

```
data(simts_3node)
```

---

true_beta_2node	<i>The true beta matrix (4 by 4) used in simulation.</i>
-----------------	--

---

**Description**

The true beta matrix (4 by 4) used in simulation.

**Usage**

```
true_beta_2node
```

**Format**

An object of class `matrix` (inherits from `array`) with 4 rows and 4 columns.

**Details**

```
true_beta_2node <- matrix(c(0,0,0,0, 0,0,0,0, 0.2,-.4,0,-0.25, 0,0.3,-0.2,0), nrow = 4, ncol = 4, by-row = TRUE)
```

**Examples**

```
true_beta_2node
```

---

```
true_beta_3node
```

*The true beta matrix (6 by 6) used in simulation.*

---

**Description**

The true beta matrix (6 by 6) used in simulation.

**Usage**

```
true_beta_3node
```

**Format**

An object of class `matrix` (inherits from `array`) with 6 rows and 6 columns.

**Details**

```
true_beta_3node <- matrix(c(0,0,0,0,0,0, 0,0,0,0,0,0, 0,0,0,0,0,0, 0.2,0,0.25,0,0,0.6, 0,0.3,0,-0.2,0,-0.6, 0,-0.2,0.3,0,0,0), nrow = 6, ncol = 6, byrow = TRUE)
```

**Examples**

```
true_beta_3node
```

---

```
uSEM
```

*Fit a multivariate time series with uSEM (unified Structural Equation Model).*

---

**Description**

Fit a multivariate time series with uSEM (unified Structural Equation Model).

**Usage**

```
uSEM(var.number,
      data,
      lag.order = 1,
      verbose = FALSE,
      trim = FALSE)
```

**Arguments**

<code>var.number</code>	number of variables in the time series
<code>data</code>	time series data, must be in long format
<code>lag.order</code>	lag order of the model to be fit, default value is 1. Note: Higher order (greater than 1) might not run.
<code>verbose</code>	print intermediate model fit (iterations), default value is FALSE
<code>trim</code>	to trim the insignificant betas (just one step, not iterative), default value is FALSE

**Details**

The purpose of uSEM is to quantify the temporal relations (both contemporaneous and lag-1) between variables. Model specification and estimation can be found in the references.

**Value**

model fit object generated by lavaan

**References**

Kim, J., Zhu, W., Chang, L., Bentler, P. M., & Ernst, T. (2007). Unified Structural Equation Modeling Approach for the Analysis of Multisubject, Multivariate Functional MRI Data. *Human Brain Mapping, 93*, 85–93. doi:10.1002/hbm.20259

Gates, K. M., & Molenaar, P. C. M. (2012). Group search algorithm recovers effective connectivity maps for individuals in homogeneous and heterogeneous samples. *NeuroImage 63*(1), 310-319. doi: 10.1016/j.neuroimage.2012.06.026

Gates, K. M., Molenaar, P. C. M., Hillary, F. G., Ram, N., & Rovine, M. J. (2010). Automatic search for fMRI connectivity mapping: An alternative to Granger causality testing using formal equivalences among SEM path modeling, VAR, and unified SEM. *NeuroImage, 50*(3), 1118–1125. doi: 10.1016/j.neuroimage.2009.12.117

**Examples**

```
model.fit <- uSEM(var.number = 3,  
                 data = simts_3node,  
                 lag.order = 1,  
                 verbose = FALSE,  
                 trim = FALSE)  
model.fit
```

---

`usemmodelfit`*Model fitbased on simulated time-series by uSEM.*

---

**Description**

Model fitbased on simulated time-series by uSEM.

**Usage**

```
usemmodelfit
```

**Format**

An object of class `lavaan` of length 1.

**Examples**

```
data(usemmodelfit)
```

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