

# Package ‘SEMsens’

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**Type** Package

**Title** A Tool for Sensitivity Analysis in Structural Equation Modeling

**Version** 0.2.6

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**Description** Perform sensitivity analysis in structural equation modeling using meta-heuristic optimization methods (e.g., ant colony optimization and others).

The references for the proposed methods are:

(1) Harring, J. R., McNeish, D. M., & Hancock, G. R. (2017)

<doi:10.1080/10705511.2018.1506925>;

(2) Socha, K., & Dorigo, M. (2008) <doi:10.1016/j.ejor.2006.06.046>.

We also thank Dr. Krzysztof Socha for sharing his thesis and R code, which provided the base for the development of this package.

**Imports** lavaan, stats

**Depends** R (>= 4.0.0)

**License** GPL-3

**Encoding** UTF-8

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**Suggests** knitr, rmarkdown

**VignetteBuilder** rmarkdown, knitr

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**NeedsCompilation** no

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## R topics documented:

SEMsens-package . . . . .	2
gen.sens.pars . . . . .	2
sa.aco . . . . .	3
sens.tables . . . . .	6

**Index**

7

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SEMsens-package	<i>A Tool for Sensitivity Analysis in Structural Equation Modeling</i>
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**Description**

This package is to help researchers perform and report sensitivity analysis in structural equation modeling using a phantom variable approach proposed by (Harring, McNeish, & Hancock, 2017).

**Details**

The package covers sensitivity analysis using ant colony optimization and other meta-heuristic optimization methods (in development) to automatically search a phantom variable, if there is any, that meets the optimization function. The current package includes three main functions and they are `gen.sens.pars` function that generates sensitivity parameters, `sa.aco` function that performs sensitivity analysis, and `sens.tables` function that summarizes sensitivity analysis results.

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gen.sens.pars	<i>Generate Sensitivity Parameters</i>
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**Description**

This function can generate a set of path coefficients from a phantom variable to variables in a structural equation model based on given distributions of the rank of optimization target (with probability of using a distribution based on its rank).

**Usage**

```
gen.sens.pars(dist.mean, dist.rank, n.of.ants, n1, q = 1e-04, k = 50, xi = 0.5)
```

**Arguments**

<code>dist.mean</code>	List of means - coordinates
<code>dist.rank</code>	Rank of the archived values of optimization object
<code>n.of.ants</code>	Number of ants used in each iteration after the initialization of k length, default value is 10.
<code>n1</code>	Neighborhood of the search area
<code>q</code>	Locality of the search (0,1), default is 0.0001
<code>k</code>	Size of the solution archive, default is 50.
<code>xi</code>	Convergence pressure (0,Inf), suggested: (0,1), default is 0.5

**Value**

Generated sensitivity parameter values (i.e., a matrix with n.of.ants rows and n.of.sens.pars columns)

**References**

Socha, K., & Dorigo, M. (2008). Ant colony optimization for continuous domains. *European Journal of Operational Research*, 185(3), 1155-1173.

We thank Dr. Krzysztof Socha for providing us the original code (<http://iridia.ulb.ac.be/supp/IridiaSupp2008-001/>) for this function.

**Examples**

```
k <- 50 # size of archive
# Generate dist.mean and dist.rank
dist.mean <- cbind(rnorm(k), rnorm(k), rnorm(k), rnorm(k))
y <- rowMeans(dist.mean)
dist.rank <- rank(-y, ties.method = "random")
# set up neighborhood
nl <- matrix(NA, k, k-1)
for (i in 1:k){
  nl[i,] <- (1:k)[1:k != i]
}
my.sens.pars <- gen.sens.pars(dist.mean, dist.rank, n.of.ants = 10,
                             nl, q = 0.0001, k =50, xi = 0.50)
my.sens.pars
```

**Description**

This function can perform sensitivity analysis for structural equation modeling using ant colony optimization (ACO).

**Usage**

```
sa.aco(
  data = NULL,
  sample.cov,
  sample.nobs,
  model,
  sens.model,
  n.of.sens.pars = NULL,
  opt.fun,
  d = NULL,
  paths = NULL,
```

```

verbose = TRUE,
max.value = Inf,
max.iter = 1000,
e = 1e-10,
n.of.ants = 10,
k = 50,
q = 1e-04,
sig.level = 0.05,
xi = 0.5,
seed = NULL
)

```

### Arguments

<code>data</code>	The data set used for analysis.
<code>sample.cov</code>	covariance matrix
<code>sample.nobs</code>	Number of observations for covariance matrix
<code>model</code>	The analytic model of interest set up as a lavaan format.
<code>sens.model</code>	Sensitivity analysis model template for structural equation modeling with a phantom variable. This is the model of interest with phantom variable and sensitivity parameters added. See examples provided.
<code>n.of.sens.pars</code>	number of sensitivity parameters added in the <code>sens.model</code> .
<code>opt.fun</code>	Customized or preset optimization function. The argument can be customized as a function, e.g., <code>opt.fun = quote(new.par\$pvalue[paths]-old.par\$pvalue[paths])</code> , where <code>new.par</code> and <code>old.par</code> are the parameter estimates from the sensitivity analysis and analytic models, respectively. When <code>opt.fun</code> is 1, the optimization function is the average departure of new estimate from the old estimate divided by the old estimate <code>y &lt;- mean(abs(new.par\$est[paths] - old.par\$est[paths])/mean(abs(old.par\$est[paths])))</code> ; When <code>opt.fun</code> is 2, the optimization function is the standard deviation of deviance divided by the old estimate <code>y &lt;- stats::sd(new.par\$est[paths] - old.par\$est[paths])/mean(abs(old.par\$est[paths]))</code> ; When <code>opt.fun</code> is 3, the optimization function is the average p value changed or <code>y &lt;- mean(abs(new.par\$pvalue[paths] - old.par\$pvalue[paths]))</code> When <code>opt.fun</code> is 4, the optimization function is the average distance from significance level or <code>y &lt;- mean(abs(new.par\$pvalue[paths] - rep(sig.level,length(paths))))#'</code>
<code>d</code>	Domains for initial sampling, default is <code>c(-1 ,1)</code> for all. It can be specified as a list of ranges (e.g., <code>d = list(-1, 1, -1, 1)</code> for two sampling domains).
<code>paths</code>	Paths in the model to be evaluated in a sensitivity analysis.
<code>verbose</code>	Print out evaluation process if true, default is TRUE.
<code>max.value</code>	Maximal value of optimization when used as the stopping criterion
<code>max.iter</code>	Maximal number of function evaluations when used as the stopping criterion
<code>e</code>	Maximum error value used when solution quality used as stopping criterion, default is 1e-10.
<code>n.of.ants</code>	Number of ants used in each iteration after the initialization of k length, default value is 10.

k	Size of the solution archive, default is 50.
q	Locality of the search (0,1), default is 0.0001
sig.level	Significance level, default value is 0.05.
xi	Convergence pressure (0,Inf), suggested: (0,1), default is 0.5
seed	Random seed if specified, default is NULL.

## Value

Sensitivity analysis results

## References

Socha, K., & Dorigo, M. (2008). Ant colony optimization for continuous domains. *European Journal of Operational Research*, 185(3), 1155-1173.

Harring, J. R., McNeish, D. M., & Hancock, G. R. (2017). Using phantom variables in structural equation modeling to assess model sensitivity to external misspecification. *Psychological Methods*, 22(4), 616.

We thank Dr. Krzysztof Socha for providing us the original code (<http://iridia.ulb.ac.be/supp/IridiaSupp2008-001/>) that the current function is based on.

## Examples

```
library(lavaan)
# generate data
sim.model <- ' x =~ x1 + 0.8*x2 + 1.2*x3
              y =~ y1 + 0.5*y2 + 1.5*y3
              m ~ 0.5*x
              y ~ 0.5*x + 0.8*m'

set.seed(10)
data <- simulateData(sim.model, sample.nobs = 1000L)
# standardize dataset
data = data.frame(apply(data,2,scale))

# Step 1: Set up the analytic model of interest
model <- 'x =~ x1 + x2 + x3
          y =~ y1 + y2 + y3
          m ~ x
          y ~ x + m'

# Step 2: Set up sensitivity analysis model
#         the sensitivity parameters are phantom1, phantom2 and phantom3
sens.model = 'x =~ x1 + x2 + x3
              y =~ y1 + y2 + y3
              m ~ x
              y ~ x + m
              x ~ phantom1*phantom
              m ~ phantom2*phantom
              y ~ phantom3*phantom
              phantom =~ 0 # added for mean of zero
              phantom ~ 1*phantom' # added for unit variance
```

```

# Step 3: check the analytic model results and decide parameter of interests
#         for sensitivity analysis
old.model = lavaan::lavaanify(model = model, auto = TRUE, model.type="sem")
old.out = lavaan::lavaan(model = old.model, data = data)
old.par = lavaan::standardizedSolution(old.out, type="std.all")
old.par # we are interested in lines 7, 8 and 9 for the indirect and direct effects

# Step 4: perform sensitivity analysis
my.sa <- sa.aco(data, model = model, sens.model = sens.model,
               n.of.sens.pars = 3, k = 5,
               opt.fun = quote(1/abs(new.par$pvalue[9]-0.05)), #p-value
               paths = 9,
               max.iter = 10)
# Note: We run with k = 5 and max.iter = 10 for illustration purpose in 5 seconds,
# please specify them as larger numbers (e.g., default value of k = 50 and mat.iter = 1000)

# Step 5: summarize sensitivity analysis results
tables <- sens.tables(my.sa)
tables

```

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sens.tables

*Summary of sensitivity analysis results*


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## Description

This function can summarize the sensitivity analysis results from [sa.aco](#) function.

## Usage

```
sens.tables(expr = NULL, sig.level = 0.05, choice = NULL)
```

## Arguments

expr	Returned object of <a href="#">sa.aco</a> function.
sig.level	Significance level, default value is 0.05.
choice	Set up the length of summary; default is all.

## Value

List of summary tables

## Examples

```
# see examples in the \link{sa.aco} function
```

# Index

`gen.sens.pars`, [2](#), [2](#)

`sa.aco`, [2](#), [3](#), [6](#)

`SEMsens (SEMsens-package)`, [2](#)

`SEMsens-package`, [2](#)

`sens.tables`, [2](#), [6](#)