

# Package ‘silp’

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**Title** Conditional Process Analysis (CPA) via SEM Approach

**Version** 1.0.0

**Description** The ‘silp’ package utilizes the Reliability-Adjusted Product Indicator (RAPI) method to estimate effects among latent variables, thus allowing for more precise definition and analysis of mediation and moderation models. Our simulation studies reveal that while ‘silp’ may exhibit instability with smaller sample sizes and lower reliability scores (e.g.,  $n = 100$ ,  $\rho = 0.7$ ), implementing nearest positive definite matrix correction and bootstrap confidence interval estimation can significantly ameliorate this volatility. When these adjustments are applied, ‘silp’ achieves estimations akin in quality to those derived from LMS. In conclusion, the ‘silp’ package is a valuable tool for researchers seeking to explore complex relational structures between variables without resorting to commercial software.

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**Encoding** UTF-8

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MASS,  
purrr,  
semTools,  
stats,  
stringr

**NeedsCompilation** no

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**URL** <https://github.com/TomBJJJ/silp>

**BugReports** <https://github.com/TomBJJJ/silp/issues>

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generate_data	<i>generate_data</i>
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## Description

Generates data based on the simulation settings provided by Cheung et al. (2021). Note that the reliability used here is omega.

## Usage

```
generate_data(
  n_obs = 100,
  corr = 0.3,
  effect = 0.42,
  ld = c(1, 1, 1, 1),
  alp = 0.9,
  effect_x = 0.4,
  effect_z = 0.2
)
```

## Arguments

n_obs	Integer. The number of observations.
corr	Numeric. The correlation of the latent variables.
effect	Numeric. The effect of the moderator.
ld	Numeric. The factor loading of the latent variable to its indicators.
alp	Numeric. The reliability of the latent variable.
effect_x	Numeric. The direct effect of x.
effect_z	Numeric. The direct effect of z.

## Value

A dataset simulated from the argument settings.

## Examples

```
n_obs = 100
corr = 0.1
effect = 0.12
ld = c(1,1,1,1)
alp = 0.9
generate_data(n_obs, corr, effect, ld, alp)
```

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resilp	<i>resilp</i>
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**Description**

An extended function from `silp`, applying the bootstrap method to obtain standard error estimation. Note: When using `silp` with the nearest positive definite matrix (`npd = TRUE`), this function should be used to obtain reliable inference.

**Usage**

```
resilp(fit, R = 2000, progress = T)
```

**Arguments**

<code>fit</code>	A result object from <code>silp</code> .
<code>R</code>	Integer. The number of bootstrap samples. Default is 2000.
<code>progress</code>	Logical. Whether to display a progress bar. Default is FALSE.

**Value**

An object of class "Silp".

**Examples**

```
n_obs = 100
corr = 0.1
effect = 0.12
ld = c(1,1,1,1)
alp = 0.9
data = generate_data(n_obs, corr, effect, ld, alp)
model = "
  fy =~ y1 + y2 + y3 + y4
  fx =~ x1 + x2 + x3 + x4
  fz =~ z1 + z2 + z3 + z4
  fy ~ fx + fz + fx:fz
"
fit = silp(model, data)
resilp(fit, R = 10)
```

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<code>silp</code>	<i>silp</i>
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**Description**

This function extends the `lavaan` function, allowing users to define moderation effects using the symbol ":". The RAPI method is used to estimate moderation effects.

**Usage**

```
silp(model, data, center = "double", tau.eq = F, npd = F, ...)
```

**Arguments**

model	A lavaan syntax model with extension. The notation ":" implies interaction between two variables (see Example).
data	The dataset for lavaan SEM.
center	Character. Whether single or double mean centering is used for the product indicator. Default is "double".
tau.eq	Logical. Specifies the type of reliability used to estimate error variance. If TRUE, Cronbach's alpha reliability is used. If FALSE, omega reliability is used. Default is FALSE.
npd	Logical. Specifies the type of input used in lavaan SEM. Default is FALSE for raw data or TRUE for a covariance matrix. Applying a covariance matrix can resolve problems of a non-positive definite covariance matrix. If TRUE, resilp should be used to obtain reliable inference.
...	Other parameters passed to the lavaan SEM function.#'

**Value**

An "Silp" class object.

**Examples**

```
n_obs = 100
corr = 0.1
effect = 0.12
ld = c(1,1,1,1)
alp = 0.9
data = generate_data(n_obs, corr, effect, ld, alp)
model = "
  fy =~ y1 + y2 + y3 + y4
  fx =~ x1 + x2 + x3 + x4
  fz =~ z1 + z2 + z3 + z4
  fy ~ fx + fz + fx:fz
"
silp(model, data)
```

---

Silp-class

*Define silp class*

---

**Description**

Define silp class

**Slots**

raw\_model The user-specified lavaan syntax model.  
 rapi\_model The revised model with the RAPI method.  
 time The operation time for silp (in seconds).  
 npd Logical. Whether the nearest positive definite matrix is used.  
 raw\_data The input data.

fa An object of class lavaan representing the CFA result.  
reliability The reliability index.  
composite\_data The composite data for RAPI.  
pa The result of silp.  
boot The results of resilp from R bootstrap samples.  
origine The original silp estimation.  
time\_resilp The operation time for resilp (in seconds).

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summary-methods

*Methods for Class Silp in Package **silp***

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## Description

Summary Methods for Class Silp in Package **silp**.

## Methods

signature(object = "Silp", method = "Bootstrap") Returns the summary result of 'silp' or 'resilp'. This method is for 'resilp' only. If method = "Bootstrap", the percentile bootstrap result is presented. If method = "BC\_b", the bias-corrected bootstrap result is presented.

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