

Package ‘semhelpinghands’

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Title Helper Functions for Structural Equation Modeling

Version 0.1.10

Description An assortment of helper functions for doing structural equation modeling, mainly by 'lavaan' for now. Most of them are time-saving functions for common tasks in doing structural equation modeling and reading the output. This package is not for functions that implement advanced statistical procedures. It is a light-weight package for simple functions that do simple tasks conveniently, with as few dependencies as possible.

URL <https://sfcheung.github.io/semhelpinghands/>

BugReports <https://github.com/sfcheung/semhelpinghands/issues>

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License GPL (>= 3)

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add_exo_cov	<i>Add Covariances Between Exogenous Variables</i>
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Description

It generates the 'lavaan' model syntax for exogenous variables in a lavaan model.

Usage

```
add_exo_cov(model, FUN = "sem", print = TRUE)
```

```
auto_exo_cov(model, FUN = "sem", print = TRUE)
```

Arguments

model	The model syntax to which the covariances are to be added.
FUN	Name (as string) of the lavaan wrapper to be called. Normally should be "sem", the default.
print	Logical. Whether the generated syntax should also be printed by <code>cat()</code> . Default is TRUE.

Details

The function `lavaan::sem()` usually will set covariances between "exogenous" variables free when `fixed.x = FALSE` ("exogenous" is defined here as variables that appear on the right hand side but not on the left hand side of the `~` operator). However, if a covariance between the residual term of an endogenous variable and an exogenous variable is manually set to free, `lavaan::sem()` may not set the aforementioned covariances free. Users will need to free them manually, and there may be a lot of them in some models.

This function gets a model syntax and generates the syntax for these covariances. Users can then inspect it, modify it if necessary, and then copy and paste it to the model syntax.

Value

`add_exo_cov()` returns a one-element character vector of the syntax, with lines separated by "\n". The generated syntax is appended to the input model syntax.

`auto_exo_cov()` returns a one-element character vector of the generated syntax, with lines separated by "\n".

Functions

- `add_exo_cov()`: Add covariances between exogenous variables to the model syntax and then return the modified model syntax.
- `auto_exo_cov()`: Generate the model syntax for the covariances between exogenous variables.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```
library(lavaan)
set.seed(8976223)
n <- 100
x <- rnorm(n)
m <- .5 * x + rnorm(n, 0, sqrt(.4))
z <- rnorm(n)
y <- .4 * m + .3 * z * m + rnorm(n, 0, .5)
dat <- data.frame(x, m, z, y)
dat$zm <- dat$z * dat$m
```

```

mod0 <-
"
m ~ x
y ~ m + z + zm
m ~~ z + zm
"

fit <- sem(mod0, dat, fixed.x = FALSE)

# Add covariances. Also printed by default.
mod0_cov <- add_exo_cov(mod0)

# Fit the model
fit_cov <- sem(mod0_cov, dat, fixed.x = FALSE)

# Manually adding the covariances
mod1 <-
"
m ~ x
y ~ m + z + zm
m ~~ z + zm
z ~~ zm + x
zm ~~ x
"

fit1 <- sem(mod1, dat, meanstructure = TRUE, fixed.x = FALSE)

# Compare the results

# No manual covariances
fit

# Automatically generated covariances
fit_cov

# Manually added covariances
fit1

```

add_sig

Add Significant Test Results

Description

It inserts columns to denote whether a parameter is significant.

Usage

```
add_sig(object, ..., standardized = FALSE, na_str = "", use = "pvalue")
```

Arguments

object	A <code>lavaan</code> object or the output of <code>lavaan::parameterEstimates()</code> or <code>lavaan::standardizedSolution()</code> . May also work on an <code>est_table</code> -class object returned by functions like <code>group_by_dvs()</code> but there is no guarantee.
...	Optional arguments to be passed to <code>lavaan::parameterEstimates()</code> or <code>lavaan::standardizedSolution()</code> .
standardized	Whether standardized solution is needed. If TRUE, <code>lavaan::standardizedSolution()</code> will be called. If FALSE, the default, <code>lavaan::parameterEstimates()</code> will be called. Ignored if a table of estimates is supplied.
na_str	The string to be used for parameters with no significant tests. For example, fixed parameters. Default is "".
use	A character vector of one or more strings. If "pvalue" is in the vector, <i>p</i> -values will be used. If "ci" is in the vector, confidence intervals appeared in <code>ci.lower</code> and <code>ci.upper</code> will be used. If "boot.ci" is in the vector and the columns <code>boot.ci.lower</code> and <code>boot.ci.upper</code> are available, these columns will be used. Note that <code>ci.lower</code> and <code>ci.upper</code> can also be bootstrap confidence intervals in some tables if <code>se = "boot"</code> is used.

Details

The function calls `lavaan::parameterEstimates()` or `lavaan::standardizedSolution()` and checks the columns `pvalue`, `ci.lower` and `ci.upper`, and/or `boot.ci.lower` and `boot.ci.upper` and then inserts columns to denote for each parameter estimate whether it is significant based on the requested criteria.

Value

The output of `lavaan::parameterEstimates()` or `lavaan::standardizedSolution()`, with one or two columns inserted after the parameter estimates to denote the significant test results.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

See Also

`lavaan::parameterEstimates()` and `lavaan::standardizedSolution()`

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model <-
'
```

```

m ~ a*x
y ~ b*m
ab := a*b
'
fit <- sem(model, data = dat, fixed.x = FALSE)

# Add "*" based on 'pvalue'
add_sig(fit)

# Add "*" for standardized solution
add_sig(fit, standardized = TRUE)

# Add "*" based on confidence interval
add_sig(fit, use = "ci")

# Add "*" for standardized solution based on confidence interval
add_sig(fit, standardized = TRUE, use = "ci")

# Add "*" for standardized solution based on confidence interval
# and 'pvalue'.
add_sig(fit, standardized = TRUE, use = c("ci", "pvalue"))

# Can also accept a parameter estimates table
est <- parameterEstimates(fit)
add_sig(est)

# So it can be used with some other functions in semhelpinghands
add_sig(filter_by(est, op = "~"))

# Piping can also be used
est |> filter_by(op = "~") |>
  add_sig()

```

 annotate_matrices

Annotate the Matrices of a 'lavaan' Model

Description

Label the elements of the model matrices in a lavaan model.

Usage

```

annotate_matrices(fit)

## S3 method for class 'annotate_matrices'
print(x, ...)

```

Arguments

<code>fit</code>	The output of <code>lavaan::lavaan()</code> or its wrappers, such as <code>lavaan::cfa()</code> and <code>lavaan::sem()</code> .
<code>x</code>	A 'annotate_matrices'-class object. The output of <code>annotate_matrices()</code> .
<code>...</code>	Optional arguments. To be passed to the default print method.

Details

This function annotates the model matrices in a `lavaan` object. This function is not to be used in analysis. It is a learning tool, for learners to understand the relation between the model matrices and the model parameters.

It currently supports a single-level single-group model only.

Value

`annotate_matrices()` returns an `annotate_matrices`-class object, which is a list of model matrices, with elements annotated:

- If a parameter is free, then it is represented by "lhs-op-rhs" according to the parameter estimate data frame.
- If a parameter is fixed but appears in the parameter table, it is represented by "(lhs-op-rhs = x)", x the value it is fixed to.
- If a parameter is fixed to zero but not in the parameter table, then it is represented by 0.

The print-method return the input, x. It was called for its side-effect.

Methods (by generic)

- `print(annotate_matrices)`: The print method of the output of `annotate_matrices()`

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```
# Adapted from https://lavaan.ugent.be/tutorial/cfa.html

library(lavaan)
HS.model <- '
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9
'

fit_cfa <- cfa(HS.model,
              data = HolzingerSwineford1939)
annotate_matrices(fit_cfa)
```

compare_estimators *Refit a 'lavaan'-Model by Several Estimators*

Description

Refit a model in 'lavaan' by several lavaan-supported estimators

Usage

```
compare_estimators(object, estimators = NULL)
```

```
se_ratios(fit_list, reference = NULL)
```

Arguments

object	A lavaan object.
estimators	A character vector of the estimator supported by the estimator argument of lavaan::lavaan() and its wrappers, such as lavaan::sem() and lavaan::cfa() .
fit_list	The output of compare_estimators() .
reference	The name of the reference method (ratios will be equal to one). Must be one of the estimator used on compare_estimators() . If NULL, the first estimator will be used.

Details

The function simply uses [lapply\(\)](#) and [update\(\)](#) to rerun the analysis once for each of the estimator using `update(object, estimator = "x", x being the estimator)`.

The results can then be compared using [group_by_models\(\)](#).

Value

A list of lavaan outputs, each of them is an update of the original output using one of the estimators.

Functions

- [compare_estimators\(\)](#): Refit the model with different estimators.
- [se_ratios\(\)](#): A wrapper of [group_by_models\(\)](#) that computes the ratios of standard errors of different methods to those of one method.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>.

See Also

[group_by_models\(\)](#)

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model <-
'
m ~ a*x
y ~ b*m
ab := a*b
'

fit <- sem(model, data = dat, fixed.x = FALSE)

# Refit the model by three different estimators
fit_more <- compare_estimators(fit, estimator = c("GLS", "MLR", "ML"))

# Use group_by_models to compare the estimates
group_by_models(fit_more, col_names = c("est", "pvalue"))

# Use se_ratios to compare standard errors
se_ratios(fit_more, reference = "ML")
```

dvs_ivs

Sample Dataset: 3 Predictors and 3 Outcomes

Description

A path model with three predictors and three outcomes, for illustration.

Usage

```
dvs_ivs
```

Format

A data frame with 100 rows and 7 variables:

- y1** Outcome variable 1. Numeric.
- y2** Outcome variable 2. Numeric.
- y3** Outcome variable 3. Numeric.
- x1** Predictor 1. Numeric.
- x2** Predictor 2. Numeric.
- x3** Predictor 3. Numeric.
- gp** Group variable: "gp1" or "gp2". String.

Examples

```

data(dvs_ivs)
library(lavaan)
mod <-
"
y1 ~ x1 + x2 + x3
y2 ~ x1 + x3
y3 ~ y2 + x2
"

fit <- sem(mod, dvs_ivs)
parameterEstimates(fit)
fit_gp <- sem(mod, dvs_ivs, group = "gp")
parameterEstimates(fit_gp)

```

filter_by

Filter a Parameter Estimates Table

Description

Filter parameter estimates table and similar tables in `lavaan` by common fields such as `op` (operator).

Usage

```
filter_by(object, op = NULL, lhs = NULL, rhs = NULL, group = NULL, fit = NULL)
```

Arguments

<code>object</code>	The output of <code>lavaan::parameterEstimates()</code> , <code>lavaan::standardizedSolution()</code> , or a <code>lavaan.data.frame</code> object. May also work on an <code>est_table</code> -class object returned by functions like <code>group_by_dvs()</code> but there is no guarantee.
<code>op</code>	A character vector of the operators (<code>op</code>) for filtering. Common operators are " <code>~</code> ", " <code>~~</code> ", " <code>=~</code> ", " <code>:=</code> ", and " <code>~1</code> ".
<code>lhs</code>	A character vector of names in the lhs column.
<code>rhs</code>	A character vector of names in the rhs column.
<code>group</code>	A vector of either the group numbers in the group column of the labels of the groups. If labels are supplied, the original fit object must be supplied for extracting the group labels.
<code>fit</code>	The original fit object. Use when <code>group</code> is a vector of the group labels.

Details

This function accepts the output of `lavaan::parameterEstimates()` and `lavaan::standardizedSolution()` and filter the rows by commonly used field.

Value

The filtered version of the input object.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model <-
'
m ~ a*x
y ~ b*m
ab := a*b
'
fit <- sem(model, data = dat, fixed.x = FALSE)

model_gp <-
'
m ~ c(a1, a2)*x
y ~ c(b1, b2)*m
a1b1 := a1*b1
a2b2 := a2*b2
'
dat$gp <- sample(c("gp1", "gp2"), n, replace = TRUE)
fit_gp <- sem(model_gp, dat, group = "gp", warn = FALSE)

est <- parameterEstimates(fit)
est_gp <- parameterEstimates(fit_gp)

filter_by(est, op = "~")

filter_by(est, op = "~", lhs = "y")

filter_by(est, rhs = c("m", "x"), op = "~")

filter_by(est_gp, group = 2)

# If the fit object is supplied, can filter
# by group label
filter_by(est_gp, group = "gp2", fit = fit_gp)
filter_by(est_gp, group = "gp2", fit = fit_gp, op = "~")

# Select user-defined parameters
```

```
filter_by(est_gp, op = ":=")

# Can be used with some other functions in semhelpinghands
# Piping can also be used
est_gp |> filter_by(op = "~", group = "gp2", fit = fit_gp) |>
  add_sig()
```

fitMeasures_by_models *Fit Measures By Models*

Description

Groups fit measures into a table with models as columns.

Usage

```
fitMeasures_by_models(object_list, ...)
```

Arguments

`object_list` A named list of [lavaan](#) objects.
`...` Optional arguments to be passed to `lavaan::fitMeasures()`.

Details

It call `lavaan::fitMeasures()` to compute for each model the fit measures supported by lavaan, and combine them into a data frame. Users can then use the print method (`print.fit_by_models()`) to customize the printout.

To be consist with full lavaan output, the names used in `lavaan::fitMeasures()` are used.

This function is intended for a simple and compact table of fit measures for quick preview. For a well-organized layout, call `lavaan::fitMeasures()` and set output to "text".

Value

A data-frame-like object of the class `fit_by_models`, which has a `print` method (see `print.fit_by_models()`).

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>.

See Also

[lavaan::fitMeasures\(\)](#)

Examples

```

library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model1 <-
'
m ~ a*x
y ~ b*m
ab := a*b
'
fit1 <- sem(model1, data = dat, fixed.x = FALSE)
model2 <-
'
m ~ a*x
y ~ b*m + x
ab := a*b
'
fit2 <- sem(model2, data = dat, fixed.x = FALSE)

fitMeasures_by_models(list(no_direct = fit1,
                           direct = fit2))

```

group_by_groups

Group Estimates By Groups

Description

Groups parameter estimates or other information such as *p*-values into a table with groups as columns and parameters as rows.

Usage

```

group_by_groups(
  object,
  ...,
  col_names = "est",
  group_first = TRUE,
  group_labels = NULL,
  fit = NULL,
  use_standardizedSolution = FALSE
)

```

Arguments

object	A <code>lavaan</code> object or the output of <code>lavaan::parameterEstimates()</code> or <code>lavaan::standardizedSolution()</code> .
...	Optional arguments to be passed to <code>lavaan::parameterEstimates()</code> . Ignored if object is an output of <code>lavaan::parameterEstimates()</code> or <code>lavaan::standardizedSolution()</code> .
col_names	A vector of the column names in the parameter estimate tables to be included. Default is "est".
group_first	If TRUE, the columns will be grouped by groups first and then by columns in the parameter estimates tables. Default is TRUE.
group_labels	A character vector of group labels. Will be assigned to group id = 1, 2, 3, etc. If not provided, will try to be retrieved from object if it is a <code>lavaan::lavaan</code> object.
fit	Optional. A <code>lavaan::lavaan</code> object. If object is a parameter estimates table and <code>group_labels</code> is NULL, it will try to retrieve the group labels from <code>fit</code> if supplied.
use_standardizedSolution	If TRUE and object is not an estimates table, then <code>lavaan::standardizedSolution()</code> will be used to generate the table. If FALSE, the default, then <code>lavaan::parameterEstimates()</code> will be used if necessary.

Value

A data-frame-like object of the class `est_table`.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```
library(lavaan)
set.seed(5478374)
n <- 100
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
city <- sample(c("City Alpha", "City Beta"), 100,
              replace = TRUE)
dat <- data.frame(x = x, y = y, m = m, city = city)
model <-
'|
m ~ c(a1, a2)*x
y ~ c(b1, b2)*m
a1b1 := a1*b1
a2b2 := a2*b2
'|

fit <- sem(model, data = dat, fixed.x = FALSE,
           group = "city")
(est <- parameterEstimates(fit))
```

```

# Group them by groups
group_by_groups(fit)

# Can also work on a parameter estimates table
# To have group labels, need to supply the fit object
group_by_groups(est, fit = fit)

# Can be used with some other functions in semhelpinghands
# when used on a parameter estimates table
group_by_groups(filter_by(est, op = "~"), fit = fit)

# Also support piping
est |> filter_by(op = "~") |>
  group_by_groups(fit = fit)

```

group_by_models

Group Estimates By Models

Description

Groups parameter estimates or other information such as p -values into a table with models as columns.

Usage

```

group_by_models(
  object_list,
  ...,
  col_names = "est",
  group_first = FALSE,
  model_first = TRUE,
  use_standardizedSolution = FALSE
)

```

Arguments

object_list	A named list of lavaan objects, a named list of the output of <code>lavaan::parameterEstimates()</code> , or a named list of the output of <code>lavaan::standardizedSolution()</code> .
...	Optional arguments to be passed to <code>lavaan::parameterEstimates()</code> . Ignored if the elements in <code>object_list</code> are the results of <code>lavaan::parameterEstimates()</code> or <code>lavaan::standardizedSolution()</code> .
col_names	A vector of the column names in the parameter estimate tables to be included. Default is "est", or "est.std" if <code>use_standardizedSolution</code> is TRUE.
group_first	If TRUE, the rows will be grouped by groups first and then by parameters. Ignored if the model has only one group. Default is FALSE.
model_first	If TRUE, the columns will be grouped by models first and then by columns in the parameter estimates tables. Default is TRUE.

use_standardizedSolution

If TRUE and object_list is not a list of estimates tables, then `lavaan::standardizedSolution()` will be used to generate the table. If FALSE, the default, then `lavaan::parameterEstimates()` will be used if necessary.

Value

A data-frame-like object of the class `est_table`.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448> Inspired by the proposal Rönkkö posted in a GitHub <https://github.com/simsem/semTools/issues/24#issue-235172313> for `semTools`. I want something simple for a quick overview and so I wrote this function.

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model1 <-
'
m ~ a*x
y ~ b*m
ab := a*b
'

fit1 <- sem(model1, data = dat, fixed.x = FALSE)
model2 <-
'
m ~ a*x
y ~ b*m + x
ab := a*b
'

fit2 <- sem(model2, data = dat, fixed.x = FALSE)
parameterEstimates(fit1)
parameterEstimates(fit2)
group_by_models(list(no_direct = fit1,
                    direct = fit2),
                col_names = c("est", "pvalue"))
# Can also be used with some other functions in
# semhelpinghands
group_by_models(list(no_direct = fit1,
                    direct = fit2),
                col_names = c("est", "pvalue")) |>
  filter_by(op = "~")
```


Description

Groups parameter estimates or other information such as p-values into a table with dependent variables as columns and independent variables as rows, or a transpose of this table.

Usage

```
group_by_dvs(
  object,
  ...,
  col_name = "est",
  add_prefix = TRUE,
  group_first = FALSE,
  use_standardizedSolution = FALSE
)
```

```
group_by_ivs(
  object,
  ...,
  col_name = "est",
  add_prefix = TRUE,
  group_first = FALSE,
  use_standardizedSolution = FALSE
)
```

Arguments

object	A <code>lavaan</code> object or the output of <code>lavaan::parameterEstimates()</code> or <code>lavaan::standardizedSolution()</code> .
...	Optional arguments to be passed to <code>lavaan::parameterEstimates()</code> . Ignored if object is an output of <code>lavaan::parameterEstimates()</code> or <code>lavaan::standardizedSolution()</code> .
col_name	The column name of information to be grouped. Default is "est". It accepts only one name.
add_prefix	If TRUE, the default, col_name will be added as prefix to the column names of the output.
group_first	If TRUE, the rows will be grouped by groups first and then by independent variables Ignored if the model has only one group. Default is FALSE.
use_standardizedSolution	If TRUE and object is not an estimates table, then <code>lavaan::standardizedSolution()</code> will be used to generate the table. If FALSE, the default, then <code>lavaan::parameterEstimates()</code> will be used if necessary.

Details

It gets a `lavaan` object or the output of `lavaan::parameterEstimates()` or `lavaan::standardizedSolution()` and group selected columns by "dependent" variables `group_by_dvs()` or by "independent" variables `group_by_ivs()`.

"Dependent" variables are defined as variables on the left hand side of the operator `~`.

"Independent" variables are defined as variables on the right hand side of the operator `~`.

Note that a variable can both be a "dependent" variable and an "independent" variable in a model.

Value

A data-frame-like object of the class `est_table`.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model <-
'
m ~ a*x
y ~ b*m
ab := a*b
'

fit <- sem(model, data = dat, fixed.x = FALSE)
parameterEstimates(fit)

# Group by DVs
group_by_dvs(fit)

# Group by IVs
group_by_ivs(fit)
```

Description

Plots for examining the distribution of bootstrap estimates in a model fitted by lavaan.

Usage

```
plot_boot(
  object,
  param,
  standardized = NULL,
  nclass = NULL,
  hist_color = "lightgrey",
  hist_linewidth = 1,
  density_line_type = "solid",
  density_line_color = "blue",
  density_line_linewidth = 2,
  est_line_type = "dotted",
  est_line_color = "red",
  est_line_linewidth = 2,
  qq_dot_size = 2,
  qq_dot_color = "black",
  qq_dot_pch = 16,
  qq_line_linewidth = 2,
  qq_line_color = "black",
  qq_line_linetype = "solid"
)
```

Arguments

object	A <code>lavaan::lavaan</code> object with bootstrap estimates stored. For standardized solution and user-defined parameters, the estimates need to be stored by <code>store_boot_est_std()</code> or <code>store_boot_def()</code> .
param	String. The name of the parameter to be plotted, which should be the name as appeared in a call to <code>coef()</code> .
standardized	Logical. Whether the estimates from the standardized solution are to be plotted. Default is <code>NULL</code> . This is a required parameter and users need to explicitly set it to <code>TRUE</code> or <code>FALSE</code> .
nclass	The number of breaks. This argument will be passed to <code>hist()</code> . Default is <code>NULL</code> .
hist_color	String. The color of the bars in the histogram. It will be passed to <code>hist()</code> for the argument <code>col</code> . Default is <code>"lightgrey"</code> .
hist_linewidth	The width of the borders of the bars in the histogram. Default is 1.

density_line_type	String. The type of the line of the density curve in the histogram. It will be passed to <code>lines()</code> for the argument <code>lty</code> . Default is "solid".
density_line_color	String. The color of the density curve in the histogram. It will be passed to <code>lines()</code> for the argument <code>col</code> . Default is "blue".
density_line_linewidth	The width of the density curve in the histogram. It will be passed to <code>lines()</code> for the argument <code>lwd</code> . Default is 2.
est_line_type	String. The type of the vertical line in the histogram showing the point estimate of the parameter. It will be passed to <code>abline()</code> for the argument <code>lty</code> . Default is "dotted",
est_line_color	String. The color of the vertical line showing the point estimate in the histogram. It will be passed to <code>abline()</code> for the argument <code>col</code> . Default is "red".
est_line_linewidth	The width of the vertical line showing the point estimate in the histogram. It will be passed to <code>hist()</code> for the argument <code>lwd</code> . Default is 2.
qq_dot_size	The size of the points in the normal QQ-plot. It will be passed to <code>qqnorm()</code> for the argument <code>cex</code> . Default is 2.
qq_dot_color	String. The color of the points in the normal QQ-plot. It will be passed to <code>qqnorm()</code> for the argument <code>col</code> . Default is "black".
qq_dot_pch	Numeric. The shape of the points in the normal QQ-plot. It will be passed to <code>qqnorm()</code> for the argument <code>pch</code> . Default is 16.
qq_line_linewidth	The width of the diagonal line to be drawn in the normal QQ-plot. It will be passed to <code>qqline()</code> for the argument <code>lwd</code> . Default is 2.
qq_line_color	String. The color of the diagonal line to be drawn in the normal QQ-plot. It will be passed to <code>qqline()</code> for the argument <code>col</code> . Default is "black".
qq_line_linetype	The type of the diagonal line to be drawn in the normal QQ-plot. Default is "solid".

Details

Rousselet, Pernet, and Wilcox (2021) argued that when using bootstrapping, it is necessary to examine the distribution of bootstrap estimates. This can be done when `boot::boot()` is used because it has a `plot` method for its output. This cannot be easily done in model fitted by `lavaan::lavaan()`.

The function `plot_boot()` is used for plotting the distribution of bootstrap estimates for a model fitted by `lavaan` in a format similar to that of the output of `boot::boot()`, with a histogram on the left and a normal QQ-plot on the right.

For free parameters in a model (unstandardized), it can be called directly on the output of `lavaan` and retrieves the stored estimates.

For estimates of user-defined parameters, call `store_boot_def()` first to compute and store the bootstrap estimates first.

For estimates in standardized solution, for both free and user-defined parameters, call `store_boot_est_std()` first to compute and store the bootstrap estimates in the standardized solution.

Value

Return the original `lavaan::lavaan` object invisibly. Called for its side-effect (plotting the graphs).

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

References

Rousselet, G. A., Pernet, C. R., & Wilcox, R. R. (2021). The percentile bootstrap: A primer with step-by-step instructions in R. *Advances in Methods and Practices in Psychological Science*, 4(1), 1–10. doi:10.1177/2515245920911881

See Also

`lavaan::fitMeasures()`, `store_boot_est_std()`, and `store_boot_def()`.

Examples

```
library(lavaan)

data(simple_mediation)
mod <-
"
m ~ a * x
y ~ b * m + x
ab := a * b
"
fit <- sem(mod, simple_mediation,
          se = "bootstrap",
          bootstrap = 50,
          iseed = 985714)

# Can plot bootstrap estimates for
# free parameters directly
# Note that 'standardized' must be always be set to
# either TRUE or FALSE. No default value.
plot_boot(fit, "a", standardized = FALSE)

# For estimates of user-defined parameters,
# call store_boot_def() first.
fit <- store_boot_def(fit)
plot_boot(fit, "ab", standardized = FALSE)

# For estimates in standardized solution,
# call store_boot_est_std() first.
fit <- store_boot_est_std(fit)
plot_boot(fit, "a", standardized = TRUE)
plot_boot(fit, "ab", standardized = TRUE)
```

plot_models_fm *Plot Models on a Chi-Squares-vs-Dfs Graph*

Description

Plot models on a graph with model chi-square against model the degrees of freedom, with lines for equal fit measures.

Usage

```
plot_models_fm(
  ...,
  fit_measure = c("cfi", "tli", "rmsea"),
  fit_values,
  line_size = 1,
  label_size = 8,
  point_size = 5,
  position_dodge = 0.5,
  include_model_values = FALSE,
  include_baseline = FALSE
)
```

Arguments

...	The lavaan::lavaan objects to be plotted. Can also be a named list of the lavaan::lavaan objects. If it is as list, it must be named and the names will be used in the plot.
fit_measure	A length-one character vector of the fit measures to use to plot the lines. Only supports "cfi" (the default), "tli", and "rmsea".
fit_values	A numeric vector of the values of the fit measure used to plot the lines. The default values are c(.90, .95) for "cfi" and "tli", and c(.00, .02, .05, .08) for "rmsea".
line_size	The size of the lines. Default is 1.
label_size	The size of the model names. Default is 8.
point_size	The size of the point representing a model. Default is 2.
position_dodge	Offsetting the label of a model from the point. Default is .5. Used by ggrepel::geom_label_repel() .
include_model_values	If TRUE, the values of the models on fit_measure will be added to fit_values. Default is FALSE.
include_baseline	If TRUE, the baseline model is included in the plot. Default is FALSE.

Details

This function plots models based on their model chi-squares and model degrees of freedoms. It can also add lines for chi-square-df combination with equal values on selected fit measures. Currently supports CFI, TLI, and RMSEA.

Value

Return a `ggplot2::ggplot()` output that can be further modified.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

See Also

`lavaan::fitMeasures()`

Examples

```
library(lavaan)

# From the help page of modificationIndices

HS.model <- '
visual  =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed   =~ x7 + x8 + x9
'

fit <- cfa(HS.model, data = HolzingerSwineford1939)
modindices(fit, sort = TRUE, op = "=")

fit2 <- update(fit, add = "visual =~ x9")
fit3 <- update(fit, add = "textual =~ x3\nvisual =~ x7")

models <- list(Initial = fit,
              Model_2 = fit2,
              Model_3 = fit3)
fit_cfi <- sapply(models, fitMeasures, fit.measures = "cfi")
fit_tli <- sapply(models, fitMeasures, fit.measures = "tli")
fit_rmsea <- sapply(models, fitMeasures, fit.measures = "rmsea")

# Supply the models as arguments
plot_models_fm(fit, fit2, fit3)

# Plot lines for selected values on a fit measure (CFI by default)
plot_models_fm(fit, fit2, fit3, fit_values = c(.90, .925, .95, fit_cfi))

# Plot the models' values on the fit measures
plot_models_fm(fit, fit2, fit3, include_model_values = TRUE)

# Supply the models as a named list
plot_models_fm(list(A = fit, B = fit2, C = fit3),
              fit_values = c(.90, .925, .95))

# Plot the models, fit measure set to TLI
```

```

plot_models_fm(fit, fit2, fit3, fit_measure = "tli")
plot_models_fm(fit, fit2, fit3, fit_measure = "tli",
               fit_values = c(.90, .925, .95, fit_tli))
plot_models_fm(fit, fit2, fit3, fit_measure = "tli",
               include_model_values = TRUE)

# Plot the models, fit measure set to RMSEA
plot_models_fm(fit, fit2, fit3, fit_measure = "rmsea")
plot_models_fm(fit, fit2, fit3, fit_measure = "rmsea",
               include_model_values = TRUE)

```

print.est_table	<i>Print an 'est_table' Object</i>
-----------------	------------------------------------

Description

Print method for an 'est_table' object

Usage

```

## S3 method for class 'est_table'
print(x, ..., nd = 3, empty_cells = "--", group_first = FALSE)

```

Arguments

x	Object of the class est_table.
...	Optional arguments to be passed to <code>print()</code> methods.
nd	The number of digits to be printed. Default is 3. (Scientific notation will never be used.)
empty_cells	String to be printed for empty cells or cells with no values. Default is "--".
group_first	Not used.

Value

x is returned invisibly. Called for its side effect.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

```
print.fit_by_models
```

Print a 'fit_by_models' Object

Description

Print method for a 'fit_by_models' object

Usage

```
## S3 method for class 'fit_by_models'
print(
  x,
  ...,
  nd = 3,
  type = c("compact"),
  remove_all_na = TRUE,
  measures_compact = c("npar", "chisq", "chisq.scaled", "df", "df.scaled", "pvalue",
    "pvalue.scaled", "chisq.scaling.factor", "cfi", "cfi.robust", "tli", "tli.robust",
    "aic", "bic", "bic2", "rmsea", "rmsea.ci.level", "rmsea.ci.lower", "rmsea.ci.upper",
    "rmsea.close.h0", "rmsea.pvalue", "rmsea.robust", "rmsea.ci.lower.robust",
    "rmsea.ci.upper.robust", "rmsea.pvalue.robust", "srmr", "srmr_omean")
)
```

Arguments

x	Object of the class fit_by_models.
...	Optional arguments to be passed to <code>print()</code> methods.
nd	The number of digits to be printed. Default is 3. (Scientific notation will never be used.)
type	String. The type of the output. Currently only supports one type, "compact".
remove_all_na	Logical. Whether rows with NA in all columns will be removed. Default is TRUE.
measures_compact	If output type is "compact", the character vector of fit measures to be printed. The names should be the names of the output of <code>lavaan::fitMeasures()</code> , in vector form.

Details

This function is intended to print the fit measures of one or more groups in a simple and compact table for quick preview. For a well-organized layout, call `lavaan::fitMeasures()` and set output to "text". For comparing the models with notations on models with the best fit on each measures, use `semTools::compareFit()`.

Value

x is returned invisibly. Called for its side effect.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

See Also

[fitMeasures_by_models\(\)](#)

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model1 <-
'
m ~ a*x
y ~ b*m
ab := a*b
'

fit1 <- sem(model1, data = dat, fixed.x = FALSE)
model2 <-
'
m ~ a*x
y ~ b*m + x
ab := a*b
'

fit2 <- sem(model2, data = dat, fixed.x = FALSE)

out <- fitMeasures_by_models(list(no_direct = fit1,
                                direct = fit2))

out

print(out, nd = 4, measures_compact = c("chisq", "cfi", "rmsea"))
```

print.std_solution_boot

Print an 'std_solution_boot' Object

Description

Print method for an 'std_solution_boot' object, which is the output of [standardizedSolution_boot_ci\(\)](#).

Usage

```
## S3 method for class 'std_solution_boot'
print(x, ..., nd = 3, output = c("table", "text"), standardized_only = TRUE)
```

Arguments

x Object of the class `std_solution_boot`.

... Optional arguments to be passed to `print()` methods.

nd The number of digits after the decimal place. Default is 3.

output String. How the results are printed. Default is "table" and the results are printed in a table format similar to that of `lavaan::standardizedSolution()`. If "text", the results will be printed in a text format similar to the printout of the output of `summary()` of a `lavaan` object.

standardized_only Logical. If TRUE, the default, only the results for the standardized solution will be printed. If FALSE, then the standardized solution is printed alongside the unstandardized solution, as in the printout of the output of `summary()` of a `lavaan` object.

Details

The default format of the printout is that of `lavaan::standardizedSolution()`, which is compact but not easy to read. Users can request a format similar to that of the printout of the summary of a `lavaan` output by setting `output` to "text".

For the "text" format, users can also select whether only the standardized solution is printed (the default) or whether the standardized solution is appended to the right of the printout.

Value

`x` is returned invisibly. Called for its side effect.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

See Also

[standardizedSolution_boot_ci\(\)](#)

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
```

```

model <-
'
m ~ a*x
y ~ b*m
ab := a*b
'

# Should set bootstrap to at least 2000 in real studies
fit <- sem(model, data = dat, fixed.x = FALSE,
           se = "boot",
           bootstrap = 50)
std_out <- standardizedSolution_boot_ci(fit)
std_out
print(std_out, output = "text")
print(std_out, output = "text", standardized_only = FALSE)

```

ptable_to_syntax

Convert a 'lavaan' Parameter Table to a 'lavaan' Model Syntax

Description

It tries to generate a 'lavaan' model syntax from a lavaan parameter table.

Usage

```
ptable_to_syntax(object, allow_incomplete = FALSE)
```

```
compare_ptables(object1, object2)
```

Arguments

object	If set to a lavaan object, such as the output of <code>lavaan::sem()</code> or <code>lavaan::cfa()</code> , the parameter table will be extracted from it by <code>lavaan::parameterTable()</code> . If set to a parameter table, it will be used to generate the model syntax. It can also be the output of <code>lavaan::lavParseModelString()</code> with <code>as.data.frame. = TRUE</code> , if <code>allow_incomplete</code> is set to <code>TRUE</code> . Note that <code>allow_incomplete</code> is set to <code>FALSE</code> by default because <code>lavaan::lavParseModelString()</code> only parses the model syntax and there is no guarantee that the model defined is valid.
allow_incomplete	Whether incomplete parameter table formed by <code>lavaan::lavParseModelString()</code> with <code>as.data.frame. = TRUE</code> is allowed. Default if <code>FALSE</code> .
object1	The first lavaan parameter table, to be compared with <code>object2</code> . If it is set to a lavaan object (e.g., the output of <code>lavaan::sem()</code> or <code>lavaan::cfa()</code>), then the parameter table will be extracted from it.
object2	The second lavaan parameter table, to be compared with <code>object1</code> . If it is set to a lavaan object (e.g., the output of <code>lavaan::sem()</code> or <code>lavaan::cfa()</code>), then the parameter table will be extracted from it.

Details

This function tries to convert a lavaan parameter table to a text representation of the lavaan model specified in model syntax.

When users call `lavaan::sem()`, in addition to the model syntax, other arguments not stored in the syntax are also used to produce the final model (e.g., `meanstructure`, `fixed.x`, and `std.lv`). To produce exactly the same model, these arguments are also needed to be specified, which is difficult to generate using only the parameter table.

Therefore, the model syntax produced will state all aspects of a model explicitly, even for those aspects that usually can be omitted due to the default values of these arguments. This approach requires users to call `lavaan::lavaan()` directly, instead of its wrappers (e.g., `lavaan::sem()`), to produce the same parameter table.

The model syntax produced this way is more difficult to read. However, it ensures that original model can be reproduced, without the need to know the arguments to set.

Due to the nearly unlimited possibilities in the form of a model, it is recommended to compare the model generated by the model syntax with the original parameter table using `compare_ptables()`. It only compares the forms of the two models, including user starting values, if any. It does not compare parameter estimates and standard errors.

Raw Specification From `lavaan::lavParseModelString()`:

There may be cases in which the parameter table is the "incomplete" table generated by `lavaan::lavParseModelString()` with `as.data.frame. = TRUE`. This table is "incomplete" because it is formed merely by parsing the model syntax. There is no guarantee that the model is valid.

The function `ptable_to_syntax()` has basic support for this kind of tables but it is disabled by default. To process an incomplete parameter table formed by `lavaan::lavParseModelString()`, set `allow_incomplete` to `TRUE`.

Limitations:

The function `ptable_to_syntax()` does not yet support the following models:

- Multiple-group models.
- Multilevel models.
- Models with categorical variables.
- Models with user-specified lower or upper bounds.
- Models with the operator `<~`.
- Models with constraints imposed by `equal()`.
- Models with labels having spaces.
- Models with labels having syntax operators (e.g., `~`, `=~`, etc.).

Value

`ptable_to_syntax()` returns a length-one character vector that stores the generated lavaan model syntax.

`compare_ptables()` returns a length-one logical vector. `TRUE` if the two models are identical in form. `FALSE` if they are not identical.

Functions

- `ptable_to_syntax()`: Convert a lavaan parameter a lavaan model syntax.
- `compare_ptables()`: Compare two lavaan parameter tables.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>. This function is inspired by a discussion at the Google Group <https://groups.google.com/g/lavaan/c/1ueFiue9qLM/m/cJhxDoqeBAAJ>.

See Also

[lavaan::lavaan\(\)](#), [lavaan::parameterTable\(\)](#)

Examples

```
library(lavaan)

mod <-
"
visual =~ x3 + x1 + x2
textual =~ x4 + x6 + x5
speed =~ x7 + x8 + x9 + start(0.1) * x6
visual ~ a*textual
speed ~ b*visual
ab: = a * b
"

fit <- sem(mod, data = HolzingerSwineford1939)

mod_chk <- ptable_to_syntax(fit)
cat(mod_chk, sep = "\n")
# Need to call lavaan() directly
fit_chk <- lavaan(mod_chk, data = HolzingerSwineford1939)
fit_chk
fit
# Compare the parameter table:
(ptable1 <- parameterTable(fit))
(ptable2 <- parameterTable(fit_chk))
compare_ptables(ptable1, ptable2)
```

Description

Record the minimization history when a model is fitted by `lavaan::lavaan()` or its wrappers (e.g., `lavaan::sem()` or `lavaan::cfa()`).

Usage

```
record_history(object)

## S3 method for class 'fit_history'
plot(x, params, last_n = -1, orientation = c("horizontal", "vertical"), ...)

## S3 method for class 'fit_history'
print(x, n_iterations = 10, digits = 3, ...)
```

Arguments

object	A lavaan object.
x	A <code>fit_history</code> class object, the output of record_history() .
params	A character vector of the names of parameters to be plotted. Must be the names of one or more columns in x.
last_n	The last n iterations to be plotted. Default is -1, plotting all iterations.
orientation	The orientation of the plot. Either "horizontal" (the default) or "vertical".
...	Optional arguments. To be passed to the print method of data frame.
n_iterations	The number of iterations to print. Default is 10, printing the first 10 iterations (or all iterations, if the number of iterations is less than 10).
digits	The number of digits to be displayed. Default is 3.

Details

It records the minimization history when a model is fitted by [lavaan::lavaan\(\)](#) or its wrappers (e.g., [lavaan::sem\(\)](#) or [lavaan::cfa\(\)](#)). The recorded history can then be plotted or displayed, for visualizing how the estimates of free parameters is found.

It will refit the model by the update method of [lavaan::lavaan](#), setting `se = "none"` and `test = "standard"` because they have no impact on the minimization process.

This and related functions are adapted from the package `semunpack`. The version in this package will be revised to be an advanced version intended for diagnostic purpose in real studies.

Value

A `fit_history`-class object with a plot method (`plot.fit_history()`).

Functions

- `plot(fit_history)`: The plot method for the output of [record_history\(\)](#).
- `print(fit_history)`: The print method for the output of [record_history\(\)](#).

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```

# Adapted from the example for CFA in lavaan::cfa().
# Using only two of the factors
library(lavaan)
HS.model <-
'
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
'

fit <- cfa(HS.model, data = HolzingerSwineford1939)

# Refit the model with the history recorded
fit_h <- record_history(fit)
fit_h

# Plot the history for selected parameters
plot(fit_h, params = c("visual=~x2", "visual=~x3",
                      "visual~~textual"),
      last_n = 10)
plot(fit_h, params = c("visual=~x2", "visual=~x3",
                      "visual~~textual"),
      last_n = 10,
      orientation = "vertical")

```

show_cfi

Visualize How CFI and TLI Are Computed

Description

Show how CFI and TLI are computed using a graph of model chi-square vs. model degrees of freedom.

Usage

```
show_cfi(fit, ...)
```

```
show_tli(fit, ...)
```

```
show_ifi(fit, fit_measures = c("cfi", "tli"), test = c("standard"))
```

Arguments

fit An output of `lavaan::lavaan()` or its wrappers (e.g., `lavaan::cfa()` and `lavaan::sem()`).

... Arguments to be passed to `show_ifi()`.

fit_measures The fit measures to be plotted. Acceptable values are "cfi" and "tli".

`test` The type of model chi-square test. It corresponds to the `test` argument of `lavaan::lavaan()` or its wrappers. Only "standard" is supported for now.

Details

This function receives an output of `lavaan::lavaan()` or its wrappers (e.g., `lavaan::cfa()` and `lavaan::sem()`) and illustrates how CFI is computed.

Value

An output of `ggplot2::ggplot()` that can be further modified.

Functions

- `show_cfi()`: A wrapper of `show_ifi()` with `fit_measures = "cfi"`.
- `show_tli()`: A wrapper of `show_ifi()` with `fit_measures = "tli"`.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```
library(lavaan)

# From the help page of lavaan::cfa().

HS.model <- '
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed   =~ x7 + x8 + x9
'

fit <- cfa(HS.model, data = HolzingerSwineford1939)

# By default show how CFI is computed
show_ifi(fit)

# Show how TLI is computed
show_ifi(fit, fit_measures = "tli")

# Wrappers
show_cfi(fit)
show_tli(fit)
```

show_more_options *Show More Major Options in an Output of 'lavaan'*

Description

Display the values of more major options in a model fitted by `lavaan::lavaan()` or its wrappers (e.g., `lavaan::sem` or `lavaan::cfa()`).

Usage

```
show_more_options(fit)

## S3 method for class 'show_more_options'
print(x, ...)
```

Arguments

<code>fit</code>	An output of <code>lavaan::lavaan()</code> or its wrappers (e.g., <code>lavaan::cfa()</code> and <code>lavaan::sem()</code>)
<code>x</code>	The output of <code>show_more_options()</code> .
<code>...</code>	Additional arguments. Ignored.

Details

It extracts the values of major options in the output of `lavaan::lavaan()` or its wrappers (e.g., `lavaan::sem` or `lavaan::cfa()`). Most of the values are also reported in the summary of a `lavaan` object. This function is used to show the values in one single table for a quick overview.

It checks the actual values, not the call used. This is useful for understanding how a prepackaged estimator such as ML, MLM, and MLR set other options. It supports the following options:

- Estimator (`estimator`)
- Standard error (`se`)
- Model chi-square test(s) (`test`)
- Missing data method (`missing`)
- Information matrix used for computing standard errors (`information`)
- Information matrix used for computing model chi-square (`information`)
- Whether the mean structure is included.

It is named `show_more_options()` to differentiate it from `show_options()`, originally in the `semunpack` package, which is intended for new users of `lavaan`. The code is adapted from `show_options` with more advanced options added.

Value

A `show_more_options`-class object with a `print` method that formats the output.

Methods (by generic)

- `print(show_more_options)`: The print method of the output of `show_more_options()`.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```
library(lavaan)

# From the help page of lavaan::cfa().

HS.model <- '
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed   =~ x7 + x8 + x9
'

fit <- cfa(HS.model, data = HolzingerSwineford1939)

tmp <- show_more_options(fit)
tmp

fit <- cfa(HS.model, data = HolzingerSwineford1939, estimator = "MLR")
show_more_options(fit)
fit <- cfa(HS.model, data = HolzingerSwineford1939, estimator = "MLM")
show_more_options(fit)
```

show_options

Show Major Options in an Output of 'lavaan'

Description

Display the values of major options in a model fitted by `lavaan::lavaan()` or its wrappers (e.g., `lavaan::sem()` or `lavaan::cfa()`).

Usage

```
show_options(fit)

## S3 method for class 'show_options'
print(x, ...)
```

Arguments

<code>fit</code>	An output of <code>lavaan::lavaan()</code> or its wrappers (e.g., <code>lavaan::cfa()</code> and <code>lavaan::sem()</code>)
<code>x</code>	The output of <code>show_options()</code> .
<code>...</code>	Additional arguments. Ignored.

Details

It extracts the values of major options in the output of `lavaan::lavaan()` or its wrappers (e.g., `lavaan::sem()` or `lavaan::cfa()`).

It checks the actual values, not the call used. This is useful for understanding how a prepackaged estimator such as ML, MLM, and MLR set other options. It supports the following options:

- Estimator (estimator)
- Standard error (se)
- Model chi-square test(s) (test)
- Missing data method (missing)
- Information matrix used for computing standard errors (information)
- Information matrix used for computing model chi-square (information)
- Whether the mean structure is included.

Value

A `show_options`-class object with a `print` method that formats the output.

Methods (by generic)

- `print(show_options)`: The `print` method of the output of `show_options()`.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```
library(lavaan)

# From the help page of lavaan::cfa().

HS.model <- '
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9
'

fit <- cfa(HS.model, data = HolzingerSwineford1939)

tmp <- show_options(fit)
tmp

fit <- cfa(HS.model, data = HolzingerSwineford1939, estimator = "MLR")
show_options(fit)
fit <- cfa(HS.model, data = HolzingerSwineford1939, estimator = "MLM")
show_options(fit)
```

simple_mediation *Sample Dataset: Simple Mediation*

Description

A simple mediation model.

Usage

```
simple_mediation
```

Format

A data frame with 100 rows and 5 variables:

x Predictor. Numeric.

m Mediator. Numeric.

y Outcome variable. Numeric.

city Group variable: "City A" or "City B". String.

Examples

```
library(lavaan)
data(simple_mediation)
mod <-
"
m ~ a * x
y ~ b * m + x
ab := a * b
"

fit <- sem(mod, simple_mediation, fixed.x = FALSE)
parameterEstimates(fit)
mod_gp <-
"
m ~ c(a1, a2) * x
y ~ c(b1, b2) * m + x
a1b1 := a1 * b1
a2b2 := a2 * b2
ab_diff := a1b1 - a2b2
"

fit_gp <- sem(mod_gp, simple_mediation, fixed.x = FALSE, group = "city")
parameterEstimates(fit_gp)
```

 sort_by

Sort a Parameter Estimates Table

Description

Sort a parameter estimates table or a similar table in `lavaan` by common fields such as `op` (operator) and `lhs` (left- hand side).

Usage

```
sort_by(
  object,
  by = c("op", "lhs", "rhs"),
  op_priority = c("=~", "~", "~~", ":", "=~1", "|", "~*~"),
  number_rows = TRUE
)
```

Arguments

<code>object</code>	The output of <code>lavaan::parameterEstimates()</code> , <code>lavaan::standardizedSolution()</code> , or a <code>lavaan.data.frame</code> object. May also work on an <code>est_table</code> -class object returned by functions like <code>group_by_dvs()</code> but there is no guarantee.
<code>by</code>	A character vector of the columns for filtering. Default is <code>c("op", "lhs", "rhs")</code> .
<code>op_priority</code>	How rows are sorted by <code>op</code> . Default is <code>c("=~", "~", "~~", ":", "=~1", " ", "~*~")</code> . Can set only a few of the operators, e.g., <code>c("~", "~~")</code> . Other operators will be placed to the end with orders not changed.
<code>number_rows</code>	Whether the row names will be set to row numbers after sorting <i>if</i> the row names of <code>object</code> is equal to row numbers. Default is <code>TRUE</code> .

Details

This functions accepts the output of `lavaan::parameterEstimates()` and `lavaan::standardizedSolution()` and filter the rows by commonly used field.

Value

The sorted version of the input object.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

Examples

```

library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model1 <-
'
m ~ a*x
y ~ b*m
ab := a*b
'
fit1 <- sem(model1, data = dat, fixed.x = FALSE)
model2 <-
'
m ~ a*x
y ~ b*m + x
ab := a*b
'
fit2 <- sem(model2, data = dat, fixed.x = FALSE)
parameterEstimates(fit1)
parameterEstimates(fit2)
out <- group_by_models(list(no_direct = fit1,
                           direct = fit2),
                       col_names = c("est", "pvalue"))

out
sort_by(out)
sort_by(out, op_priority = c("~", ":="))
sort_by(out, by = c("op", "rhs"))

```

standardizedSolution_boot_ci

Bootstrap CIs for Standardized Solution

Description

Functions for forming bootstrap confidence intervals for the standardized solution.

Usage

```

standardizedSolution_boot_ci(
  object,
  level = 0.95,
  type = "std.all",

```

```

    save_boot_est_std = TRUE,
    force_run = FALSE,
    boot_delta_ratio = FALSE,
    ...
)

store_boot_est_std(object, type = "std.all", force_run = FALSE, ...)

get_boot_est_std(object)

```

Arguments

<code>object</code>	A <code>lavaan</code> object, fitted with <code>'se = "boot"'</code> .
<code>level</code>	The level of confidence of the confidence intervals. Default is <code>.95</code> .
<code>type</code>	The type of standard estimates. The same argument of <code>lavaan::standardizedSolution()</code> , and support all values supported by <code>lavaan::standardizedSolution()</code> . Default is <code>"std.all"</code> .
<code>save_boot_est_std</code>	Whether the bootstrap estimates of the standardized solution are saved. If saved, they will be stored in the attribute <code>boot_est_std</code> . Default is <code>TRUE</code> .
<code>force_run</code>	If <code>TRUE</code> , will skip checks and run models without checking the estimates. For internal use. Default is <code>FALSE</code> .
<code>boot_delta_ratio</code>	The ratio of (a) the distance of the bootstrap confidence limit from the point estimate to (b) the distance of the delta-method limit from the point estimate. Default is <code>FALSE</code> .
<code>...</code>	Other arguments to be passed to <code>lavaan::standardizedSolution()</code> .

Details

`standardizedSolution_boot_ci()` receives a `lavaan::lavaan` object fitted with bootstrapping standard errors requested and forms the confidence intervals for the standardized solution.

It works by calling `lavaan::standardizedSolution()` with the bootstrap estimates of free parameters in each bootstrap sample to compute the standardized estimates in each sample.

A more reliable way is to use function like `lavaan::bootstrapLavaan()`. Nevertheless, this simple function is good enough for some simple scenarios, and does not require repeating the bootstrapping step.

`store_boot_est_std()` computes the standardized solution for each bootstrap sample, stores them the `lavaan::lavaan` object, and returns it. These estimates can be used by other functions, such as `plot_boot()`, to examine the estimates, without the need to repeat the computation.

`get_boot_est_std()` retrieves the bootstrap estimates of the standardized solution stored by `store_boot_est_std()`.

Value

The output of `lavaan::standardizedSolution()`, with bootstrap confidence intervals appended to the right, with class set to `std_solution_boot` (since version 0.1.8.4). It has a print method

(`print.std_solution_boot()`) that can be used to print the standardized solution in a format similar to that of the printout of the `summary()` of a `lavaan::lavaan` object.

`store_boot_est_std()` returns the fit object set to object, with the bootstrap values of standardized solution in the bootstrap samples, as a matrix, stored in the slot `external` under the name `shh_boot_est_std`.

`get_boot_est_std()` returns a matrix of the stored bootstrap estimates of standardized solution. If none is stored, NULL is returned.

`store_boot_est_std()` is usually used with diagnostic functions such as `plot_boot()`.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>. Originally proposed in an issue at GitHub <https://github.com/simsem/semTools/issues/101#issue-1021974657>, inspired by a discussion at the Google group for lavaan <https://groups.google.com/g/lavaan/c/qQBXSz5cd0o/m/R8YT5HxNAgAJ>. `boot::boot.ci()` is used to form the percentile confidence intervals in this version.

See Also

`lavaan::standardizedSolution()`, `plot_boot()`

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model <-
'
m ~ a*x
y ~ b*m
ab := a*b
'

# Should set bootstrap to at least 2000 in real studies
fit <- sem(model, data = dat, fixed.x = FALSE,
           se = "boot",
           bootstrap = 100)
summary(fit)

std <- standardizedSolution_boot_ci(fit)
std

# Print in a friendly format with only standardized solution
print(std, output = "text")
```

```

# Print in a friendly format with both unstandardized
# and standardized solution
print(std, output = "text", standardized_only = FALSE)

# store_boot_est_std() is usually used with plot_boot()
# First, store the bootstrap estimates of the
# standardized solution
fit_with_boot_std <- store_boot_est_std(fit)
# Second, plot the distribution of the bootstrap estimates of
# standardized 'ab'
plot_boot(fit_with_boot_std, "ab", standardized = TRUE)

```

store_boot_def

Store Bootstrap Estimates of User-Defined Parameters

Description

It receives a [lavaan::lavaan](#) object fitted with bootstrapping standard errors requested, computes the user-defined parameters in each bootstrap samples, and returns a [lavaan::lavaan](#) object with the estimates stored.

Usage

```

store_boot_def(object, force_run = FALSE)

get_boot_def(object)

```

Arguments

object	A lavaan object, fitted with 'se = "boot"'.
force_run	If TRUE, will skip checks and run models without checking the estimates. For internal use. Default is FALSE.

Details

[lavaan::lavaan\(\)](#) and its wrappers, such as [lavaan::sem\(\)](#) and [lavaan::cfa\(\)](#), stores the estimates of free parameters in each bootstrap sample if bootstrapping is requested. However, if a model has user-defined parameters, their values in each bootstrap sample are not stored. [store_boot_def\(\)](#) computes the retrieves the stored bootstrap estimates and computes the values of user-defined parameters. The values are then stored in the slot `external` of the object, in the element `shh_boot_def`. The bootstrap estimates can then be used by other functions for diagnostics purposes.

[get_boot_def\(\)](#) extracts the bootstrap estimates of user-defined parameters from a [lavaan](#) object. If none is stored, NULL is returned.

[store_boot_def\(\)](#) is usually used with diagnostic functions such as [plot_boot\(\)](#).

Value

`store_boot_def()` returns the fit object set to object, with the bootstrap values of user-defined parameters in the bootstrap samples, as a matrix, stored in the slot `external` of object under the name `shh_boot_def`.

`get_boot_def()` returns a matrix of the stored bootstrap estimates of user-defined parameters

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>.

See Also

`plot_boot()`

Examples

```
library(lavaan)
set.seed(5478374)
n <- 50
x <- runif(n) - .5
m <- .40 * x + rnorm(n, 0, sqrt(1 - .40))
y <- .30 * m + rnorm(n, 0, sqrt(1 - .30))
dat <- data.frame(x = x, y = y, m = m)
model <-
'
m ~ a*x
y ~ b*m
ab := a*b
'

# Should set bootstrap to at least 2000 in real studies
fit <- sem(model, data = dat, fixed.x = FALSE,
           se = "boot",
           bootstrap = 100)
summary(fit)

# store_boot_def() is usually used with plot_boot()
# First, store the bootstrap estimates of user-defined
# parameters
fit_with_boot_def <- store_boot_def(fit)
# Second, plot the distribution of the bootstrap estimates of
# 'ab'
plot_boot(fit_with_boot_def, "ab", standardized = FALSE)
```

`vec_rsquare`*Wrapper Functions to Extract Information as a Vector*

Description

A set of wrapper functions to extract information from a lavaan-class object and return a named vector.

Usage`vec_rsquare(object)``vec_sample_vcov(object)``vec_sample_var(object)``vec_est_var(object)``vec_est_se(object)``vec_def_var(object)``vec_def_se(object)`

```
vec_lavTestLRT(  
  object,  
  ...,  
  method = "default",  
  A.method = "delta",  
  scaled.shifted = TRUE,  
  H1 = TRUE,  
  model.names = NULL  
)
```

```
vec_lavTestScore(  
  object,  
  add = NULL,  
  release = NULL,  
  univariate = TRUE,  
  information = "expected"  
)
```

`vec_lavTestWald(object, constraints = NULL, prefix = NULL)``vec_compRelSEM(object, ...)`

Arguments

object	A lavaan-class object.
...	Additional arguments to be passed to the original function.
method	An argument to be passed to <code>lavaan::lavTestLRT()</code> . Please refer to the help page of <code>lavaan::lavTestLRT()</code> .
A.method	An argument to be passed <code>lavaan::lavTestLRT()</code> . Please refer to the help page of <code>lavaan::lavTestLRT()</code> .
scaled.shifted	An argument to be passed to <code>lavaan::lavTestLRT()</code> . Please refer to the help page of <code>lavaan::lavTestLRT()</code> .
H1	An argument to be passed to <code>lavaan::lavTestLRT()</code> . Please refer to the help page of <code>lavaan::lavTestLRT()</code> .
model.names	An argument to be passed to <code>lavaan::lavTestLRT()</code> . Please refer to the help page of <code>lavaan::lavTestLRT()</code> . Unlike <code>lavaan::lavTestLRT()</code> , this argument is required, for the sake of naming the vector to be returned.
add	An argument to be passed to <code>lavaan::lavTestScore()</code> . Please refer to the help page of <code>lavaan::lavTestScore()</code> .
release	An argument to be passed to <code>lavaan::lavTestScore()</code> . Please refer to the help page of <code>lavaan::lavTestScore()</code> .
univariate	An argument to be passed to <code>lavaan::lavTestScore()</code> . Please refer to the help page of <code>lavaan::lavTestScore()</code> .
information	An argument to be passed to <code>lavaan::lavTestScore()</code> . Please refer to the help page of <code>lavaan::lavTestScore()</code> .
constraints	An argument to be passed to <code>lavaan::lavTestWald()</code> . Please refer to the help page of <code>lavaan::lavTestWald()</code> .
prefix	Optional. A character string to be added as a prefix to names of the output. Default is NULL.

Details

This set of wrapper functions are for functions like `lavaan::bootstrapLavaan()` that require users to supply a function that receives a lavaan-class object and returns a vector of values.

All wrappers functions are designed to have the same form of output: a named numeric vector.

Many of the tasks of this set of wrappers can be performed by writing our own functions. The wrapper functions are developed just to save the coding time for some commonly requested information.

The wrapper functions are designed to be as simple to use as possible, with as few arguments as possible. If advanced control is needed, users are recommended to write their own wrappers.

Value

All of them return a named numeric vector.

Functions

- `vec_rsquare()`: Get R-squares in a model.
- `vec_sample_vcov()`: Get sample variances and covariances.
- `vec_sample_var()`: Get sample variances.
- `vec_est_var()`: Sampling variances of free parameters.
- `vec_est_se()`: Standard errors of free parameters.
- `vec_def_var()`: Sampling variances of user-defined parameters.
- `vec_def_se()`: Standard errors of user-defined parameters.
- `vec_lavTestLRT()`: Get sample variances.
- `vec_lavTestScore()`: Do score tests.
- `vec_lavTestWald()`: Do a Wald test.
- `vec_compRelSEM()`: Composite reliability.

Author(s)

Shu Fai Cheung <https://orcid.org/0000-0002-9871-9448>

See Also

[lavaan::lavInspect\(\)](#)

Examples

```
# From the help page of lavaan::cfa().

library(lavaan)
HS.model <- '
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9
'

fit <- cfa(HS.model, data = HolzingerSwineford1939)

vec_rsquare(fit)
vec_sample_vcov(fit)
vec_sample_var(fit)
vec_est_var(fit)
vec_est_se(fit)

HS.model.sem1 <- '
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9
textual ~ a * visual
speed ~ b * textual
ab := a * b
'
```

```
fit_sem1 <- sem(HS.model.sem1, data = HolzingerSwineford1939)

HS.model.sem2 <- '
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9
textual ~ a * visual
speed ~ b * textual + cp * visual
ab := a * b
'

fit_sem2 <- sem(HS.model.sem2, data = HolzingerSwineford1939)

vec_def_var(fit_sem1)
vec_def_se(fit_sem1)

vec_lavTestLRT(fit_sem1, fit_sem2,
               model.names = c("No Direct", "Direct"))

vec_lavTestScore(fit_sem1,
                 add = "speed ~ visual")

vec_lavTestWald(fit_sem2,
                constraints = "cp == 0")

if (requireNamespace("semTools")) {
  vec_compRelSEM(fit)
}
```

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