

# Package ‘tidySEM’

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**Title** Tidy Structural Equation Modeling

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**Description** A tidy workflow for generating, estimating, reporting, and plotting structural equation models using 'lavaan', 'OpenMx', or 'Mplus'. Throughout this workflow, elements of syntax, results, and graphs are represented as 'tidy' data, making them easy to customize.

**License** GPL (>= 3)

**URL** <https://cjvanlissa.github.io/tidySEM/>

**BugReports** <https://github.com/cjvanlissa/tidySEM/issues>

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---

`add_paths`*Add paths to an object of class 'tidy\_sem'*

---

## Description

Add paths to an object of class `tidy_sem`, or replace existing paths. The paths must be specified as [model.syntax](#), and separated by commas.

## Usage

```
add_paths(model, ...)
```

## Arguments

<code>model</code>	An object of class <code>tidy_sem</code> .
<code>...</code>	Paths to add or substitute, specified in <a href="#">lavaan{model.syntax}</a> , and separated by commas.

## Details

Currently, only the [lavaan{lavaan}](#) commands `~`, `~~`, `=~`, and `~1` are parsed.

This function relies on [lavaan model.syntax](#) to convert syntax strings to lavaan parameter tables. By default, it uses the arguments `int.ov.free = TRUE`, `int.lv.free = FALSE`, `auto.fix.first = TRUE`, `auto.fix.single = TRUE`, `auto.var = TRUE`, `auto.cov.lv.x = TRUE`, `auto.efa = TRUE`, `auto.th = TRUE`, `auto.delta = TRUE`, `auto.cov.y = TRUE`, `meanstructure = TRUE`, in a similar way to [sem](#) and [cfa](#).

## Value

An object of class `tidy_sem`.

## See Also

[model.syntax](#)

## Examples

```
library(lavaan)
df <- iris[, 1:4]
names(df) <- paste0("x_", 1:4)
model <- tidy_sem(df)
model <- measurement(model)
model <- add_paths(model, x =~ a*x_1 + b*x_2 + a*x_3 + b*x_4)
res <- estimate_lavaan(model)
summary(res)
```

---

as_lavaan	<i>Convert tidy_sem to 'lavaan' syntax</i>
-----------	--

---

**Description**

Final stage in the 'tidySEM' workflow for syntax generation: Convert the tidy\_sem object to lavaan syntax in tabular format (see [model.syntax](#)).

**Usage**

```
as_lavaan(x, ...)
```

**Arguments**

x	An object of class tidy_sem
...	Additional parameters to be passed to and from functions.

**Value**

Character vector.

**Examples**

```
mod <- list(syntax = structure(list(lhs = "x", op = "~", rhs = "y",
                                free = TRUE, value = "", label = "",
                                category = "", aspect = ""),
                              class = "data.frame", row.names = c(NA, -1L)))
class(mod) <- "tidy_sem"
as_lavaan(mod)
```

---

as_mplus	<i>Convert tidy_sem to 'Mplus' syntax</i>
----------	---

---

**Description**

Final stage in the 'tidySEM' workflow for syntax generation: Convert the tidy\_sem object to 'Mplus' syntax.

**Usage**

```
as_mplus(x, ...)
```

**Arguments**

x	An object of class tidy_sem.
...	Additional parameters to be passed to and from functions.

**Value**

Character vector.

**Examples**

```
mod <- list(syntax = structure(list(lhs = "x", op = "~", rhs = "y",
                                free = TRUE, value = "", label = "",
                                category = "", aspect = ""),
                              class = "data.frame", row.names = c(NA, -1L)))
class(mod) <- "tidy_sem"
as_mplus(mod)
```

---

as\_ram

---

*Convert lavaan syntax to RAM specification*


---

**Description**

Converts SEM models to RAM models for OpenMx.

**Usage**

```
as_ram(x, ...)
```

**Arguments**

**x** An object for which a method exists, such as a `tidy_sem` object, or character vector describing the user-specified model using the lavaan model syntax.

**...** Parameters passed on to other functions.

**Details**

For models specified using lavaan syntax, the procedure is as follows:

1. Apply [lavaanify](#) to the model. The default arguments to [lavaanify](#) correspond to those of the [sem](#) function.
2. Convert each row of the resulting lavaan parameter table to a [mxPath](#).
3. Apply [mxModel](#) to the `mxPaths` to create an OpenMx model using RAM specification

**Value**

Returns an [mxModel](#).

**Examples**

```
as_ram("y ~ x")
```

**Description**

Estimate an auxiliary model based on a latent classification by means of mixture modeling (see [mx\\_mixture](#)).

The auxiliary model is treated as a multi-group model. All cases are used in all groups, but they are weighted by group-specific BCH weights as described in Bolck, Croon, & Hagnaars, 2004.

**Usage**

```
BCH(x, model, data, ...)
```

**Arguments**

x	An object for which a method exists.
model	An object that can be converted to an OpenMx model using <a href="#">as_ram</a> .
data	A data.frame on which the auxiliary model can be evaluated.
...	further arguments to be passed to or from other methods.

**Value**

An MxModel.

**References**

Bolck, A., Croon, M., & Hagnaars, J. (2004). Estimating latent structure models with categorical variables: One-step versus three-step estimators. *Political Analysis*, 12(1), 3–27. <doi:10.2307/25791751>

**Examples**

```
dat <- data.frame(x = iris$Petal.Length)
mixmod <- mx_profiles(dat,
                      classes = 2)
res <- BCH(mixmod, "y ~ 1", data = data.frame(y = iris$Sepal.Length))
```

---

 BLRT

*Conduct Bootstrapped Likelihood Ratio Test*


---

**Description**

Conduct Bootstrapped Likelihood Ratio Test to compare two mixture models.

**Usage**

```
BLRT(x, ...)
```

**Arguments**

`x` An object for which a method exists.  
`...` further arguments to be passed to or from other methods.

**Value**

A data.frame.

**Examples**

```
## Not run:
df <- iris[, 1, drop = FALSE]
names(df) <- "x"
res <- mx_mixture(model = "x ~ m{C}*1
                        x ~~ v{C}*x", classes = 1:2, data = df)
BLRT(res, replications = 4)

## End(Not run)
```

---

 class\_prob

*Obtain latent class probabilities*


---

**Description**

Obtain latent class probabilities for an object for which a method exists. See Details.

**Usage**

```
class_prob(
  x,
  type = c("sum.posterior", "sum.mostlikely", "mostlikely.class", "avg.mostlikely",
           "individual"),
  ...
)
```

**Arguments**

x	An object for which a method exists.
type	Character vector, indicating which types of probabilities to extract. See Details.
...	Further arguments to be passed to or from other methods.

**Details**

The following types are available:

- "sum.posterior" A summary table of the posterior class probabilities; this indicates what proportion of your data contributes to each class.
- "sum.mostlikely" A summary table of the most likely class membership, based on the highest posterior class probability. Note that this is subject to measurement error.
- "mostlikely.class" If C is the true class of an observation, and N is the most likely class based on the model, then this table shows the probability  $P(N=i|C=j)$ . The diagonal represents the probability that observations in each class will be correctly classified.
- "avg.mostlikely" Average posterior probabilities for each class, for the subset of observations with most likely class of 1:k, where k is the number of classes.
- "individual" The posterior probability matrix, with dimensions n (number of cases in the data) x k (number of classes).

**Value**

A data.frame.

**Examples**

```
## Not run:
df <- iris[, 1, drop = FALSE]
names(df) <- "x"
res <- mx_mixture(model = "x ~ m{C}*1
                        x ~~ v{C}*x", classes = 1, data = df)
class_prob(res)

## End(Not run)
```

---

conf\_int

*Format confidence intervals*

---

**Description**

Creates 'APA'-formatted confidence intervals, either from an object for which a method exists, or from the arguments lb and ub. When argument x is a numeric vector, it is also possible to construct a confidence interval using the standard error (se) and a percentile interval (ci).



**Usage**

```
conf_int(x, digits = 2, se = NULL, lb = NULL, ub = NULL, ci = 95)
```

**Arguments**

<code>x</code>	Optional. An object for which a method exists.
<code>digits</code>	Integer. The number of digits to round the confidence boundaries to.
<code>se</code>	Optional, numeric. Standard error of the parameters.
<code>lb</code>	Optional, numeric. Lower boundary of confidence intervals.
<code>ub</code>	Optional, numeric. Upper boundary of confidence intervals.
<code>ci</code>	Optional, numeric. What percentage CI to use (only used when computing CI from a numeric vector <code>x</code> , and the standard error <code>se</code> , based on a normal distribution).

**Value**

A character vector of formatted confidence intervals.

**Author(s)**

Caspar J. van Lissa

**See Also**

`table_results` `est_sig`

Other Reporting tools: [est\\_sig\(\)](#), [table\\_fit\(\)](#), [table\\_results\(\)](#)

**Examples**

```
conf_int(x = c(1.325, 2.432), se = c(.05336, .00325))
```

---

`cors`

*Generate syntax for correlations*

---

**Description**

Generate syntax for correlations between variables.

**Usage**

```
cors(x, ...)
```

**Arguments**

- `x` Object for which a method exists. If `x` is an object of class `tidy_sem`, then correlations between all observed and latent variables in the data dictionary of that object are computed, by default. If `x` is a character vector, all elements of the vector are used.
- `...` Optional additional character vectors of variables to be correlated. If `x` is an object of class `tidy_sem`, then up to two vectors can be provided. If `x` is a vector, then one more optional vector can be provided. When no additional vectors of variable names are provided, only the correlations among the elements of `x` are returned.

**Value**

An object of class `tidy_sem`.

**Examples**

```
dict <- tidy_sem(c("bfi_1", "bfi_2", "bfi_3", "bfi_4", "bfi_5"))
cors(dict, c("bfi_1", "bfi_2"))
```

---

create\_scales

*Create scale scores from observed variables*

---

**Description**

This function calculates mean or sum scores from a `data.frame` and a named list describing the items in each scale. It returns the scores, a scale descriptive table, and a scale correlation table. It relies on several functions from the `psych` package.

**Usage**

```
create_scales(
  x,
  keys.list,
  missing = TRUE,
  impute = "none",
  omega = NULL,
  digits = 2,
  ...
)

## S3 method for class 'tidy_sem'
create_scales(
  x,
  keys.list,
  missing = TRUE,
  impute = "none",
```

```

    omega = NULL,
    digits = 2,
    ...
  )

```

### Arguments

<code>x</code>	A data.frame containing all variables referenced in the <code>keys.list</code> , or an object of class <code>tidy_sem</code> .
<code>keys.list</code>	A named list, indicating which variables belong to which scale.
<code>missing</code>	Whether to use rows with partially missing values. Default: TRUE.
<code>impute</code>	Method for handling missing values, Default: 'none'. This default method uses all available data to calculate scale scores, which is acceptable for mean scales, but not for sum scales.
<code>omega</code>	Which of McDonald's <a href="#">omega</a> coefficients to report. Default: NULL; valid options include: "omega_h", "omega_lim", "alpha", "omega_tot", "G6".
<code>digits</code>	Number of digits for rounding, Default: 2
<code>...</code>	Additional parameters to pass to and from functions.

### Details

For scales with less than 3 items, Cronbach's alpha might not be suitable as an estimate of reliability. For such scales, the Spearman-Brown reliability coefficient for two-item scales is computed, as described in Eisinga, R., Grotenhuis, M. te, & Pelzer, B. (2012). The reliability of a two-item scale: Pearson, Cronbach, or Spearman-Brown? *International Journal of Public Health*, 58(4), 637–642. <doi:10.1007/s00038-012-0416-3>. These coefficients are marked with "(sb)".

### Value

List with elements: `$descriptives`, `$correlations`, and `$scores`.

### Examples

```

out <- create_scales(iris, keys.list = list(scalename =
  c("Sepal.Length", "Sepal.Width", "Petal.Length", "Petal.Width")))
out$descriptives
dict <- tidy_sem(iris, split = "\\.")
create_scales(dict)

```

---

curry\_mac

*Simulated MAC data*

---

### Description

This simulated dataset, based on Curry et al., 2019, contains data on moral relevance and judgment across the seven domains of the Morality As Cooperation scale.

**Usage**

```
data(curry_mac)
```

**Format**

A data.frame with 1392 rows and 42 variables.

**Details**

<b>sex</b>	factor	Self-identified sex of participants, Male, Female, or Transgendered.
<b>age_years</b>	numeric	Participants' age in years.
<b>KinshipR</b>	numeric	Mean score of moral relevance, kinship subscale.
<b>MutualismR</b>	numeric	Mean score of moral relevance, mutualism subscale.
<b>ExchangeR</b>	numeric	Mean score of moral relevance, exchange subscale.
<b>HawkR</b>	numeric	Mean score of moral relevance, hawk subscale.
<b>DoveR</b>	numeric	Mean score of moral relevance, dove subscale.
<b>DivisionR</b>	numeric	Mean score of moral relevance, division subscale.
<b>PossessionR</b>	numeric	Mean score of moral relevance, possession subscale.
<b>KinshipJ</b>	numeric	Mean score of moral judgment, kinship subscale.
<b>MutualismJ</b>	numeric	Mean score of moral judgment, mutualism subscale.
<b>ExchangeJ</b>	numeric	Mean score of moral judgment, exchange subscale.
<b>HawkJ</b>	numeric	Mean score of moral judgment, hawk subscale.
<b>DoveJ</b>	numeric	Mean score of moral judgment, dove subscale.
<b>DivisionJ</b>	numeric	Mean score of moral judgment, division subscale.
<b>PossessionJ</b>	numeric	Mean score of moral judgment, possession subscale.

**References**

Curry, O. S., Jones Chesters, M., & Van Lissa, C. J. (2019). Mapping morality with a compass: Testing the theory of 'morality-as-cooperation' with a new questionnaire. *Journal of Research in Personality*, 78, 106–124. doi: [10.1016/j.jrp.2018.10.008](https://doi.org/10.1016/j.jrp.2018.10.008)

---

 descriptives

*Describe a dataset*


---

**Description**

Provide descriptive statistics for a dataset.

**Usage**

```
descriptives(x, ...)
```

**Arguments**

x                    An object for which a method exists.  
...                   Additional arguments.

**Value**

A data.frame with descriptive statistics for x.

**Examples**

```
descriptives(iris)
```

---

dictionary	<i>Extract dictionary from tidy_sem</i>
------------	---

---

**Description**

Provides access to the dictionary element of a tidy\_sem object. This can be used to return or assign to the dictionary element.

**Usage**

```
dictionary(x)  
dictionary(x) <- value
```

**Arguments**

x                    Object of class tidy\_sem.  
value                A valid value for dictionary(x).

**Value**

data.frame

**Examples**

```
dict <- tidy_sem(iris, split = "\\.")  
dictionary(dict)
```

---

edges	<i>Extract edges from sem_graph</i>
-------	-------------------------------------

---

**Description**

Provides access to the edges element of a sem\_graph object. This can be used to return or assign to the edges element.

**Usage**

```
edges(x)
```

```
edges(x) <- value
```

**Arguments**

x	Object of class sem_graph.
value	A valid value for edges(x).

**Value**

data.frame

**Examples**

```
edg <- data.frame(from = "x", to = "y")
p <- prepare_graph(edges = edg, layout = get_layout("x", "y", rows = 1))
edges(p)
```

---

edit_graph	<i>Edit graph elements</i>
------------	----------------------------

---

**Description**

Evaluate an R expression within the environment of the elements of a sem\_graph object, and return the modified sem\_graph.

**Usage**

```
edit_graph(x, expr, element = c("edges", "nodes"), ...)
```

```
edit_nodes(x, expr, ...)
```

```
edit_edges(x, expr, ...)
```

**Arguments**

<code>x</code>	An object of class <code>sem_graph</code> .
<code>expr</code>	expression to evaluate.
<code>element</code>	Character. The element of the <code>sem_graph</code> to edit, defaults to <code>c("edges", "nodes")</code> .
<code>...</code>	Arguments passed on to <code>within</code> .

**Value**

An object of class `sem_graph`.

**Examples**

```
p <- prepare_graph(layout = get_layout("x", rows = 1))
p <- edit_graph(p, {colour = "blue"}, element = "nodes")
plot(p)
```

---

empathy	<i>Simulated empathy data</i>
---------	-------------------------------

---

**Description**

This simulated dataset, based on Van Lissa et al., 2014, contains six annual assessments of adolescents' mean scores on the empathic concern and perspective taking subscales of the Interpersonal Reactivity Index (Davis, 1983). The first measurement wave occurred when adolescents were, on average, 13 years old, and the last one when they were 18 years old.

**Usage**

```
data(empathy)
```

**Format**

A data frame with 467 rows and 13 variables.

**Details**

<b>ec1</b>	numeric	Mean score of empathic concern in wave 1
<b>ec2</b>	numeric	Mean score of empathic concern in wave 2
<b>ec3</b>	numeric	Mean score of empathic concern in wave 3
<b>ec4</b>	numeric	Mean score of empathic concern in wave 4
<b>ec5</b>	numeric	Mean score of empathic concern in wave 5
<b>ec6</b>	numeric	Mean score of empathic concern in wave 6
<b>pt1</b>	numeric	Mean score of perspective taking in wave 1
<b>pt2</b>	numeric	Mean score of perspective taking in wave 2
<b>pt3</b>	numeric	Mean score of perspective taking in wave 3
<b>pt4</b>	numeric	Mean score of perspective taking in wave 4

<b>pt5</b>	numeric	Mean score of perspective taking in wave 5
<b>pt6</b>	numeric	Mean score of perspective taking in wave 6
<b>sex</b>	factor	Adolescent sex; M = male, F = female.

## References

Van Lissa, C. J., Hawk, S. T., Branje, S. J., Koot, H. M., Van Lier, P. A., & Meeus, W. H. (2014). Divergence Between Adolescent and Parental Perceptions of Conflict in Relationship to Adolescent Empathy Development. *Journal of Youth and Adolescence*, (Journal Article), 1–14. doi: [10.1007/s1096401401525](https://doi.org/10.1007/s1096401401525)

---

estimate_lavaan	<i>Estimate tidy_sem using 'lavaan'</i>
-----------------	---

---

## Description

This function is a wrapper for the [lavaan](#) estimating functions. By default, the wrapper uses [sem](#), but users can also specify [lavaan](#), [cfa](#), or [growth](#).

## Usage

```
estimate_lavaan(x, func = "sem", ...)
```

## Arguments

x	An object of class <code>tidy_sem</code> .
func	The <a href="#">lavaan</a> modeling function to invoke, Default: 'sem'.
...	Additional parameters passed to the estimating function.

## Value

An object of class `lavaan`.

## Examples

```
library(lavaan)
model <- tidy_sem(iris, "\\.")
model <- measurement(model)
res <- estimate_lavaan(model)
summary(res)
```



---

estimate_mplus	<i>Estimate tidy_sem using 'Mplus'</i>
----------------	--

---

**Description**

This function is a wrapper for the functions `mplusObject` and `mplusModeler`. Using this function requires 'Mplus' to be installed.

**Usage**

```
estimate_mplus(x, ...)
```

**Arguments**

x	An object of class <code>tidy_sem</code> .
...	Additional parameters passed to <code>mplusObject</code> and <code>mplusModeler</code> . These arguments are matched to the correct function by name. The arguments <code>rdata</code> , and <code>MODEL</code> cannot be edited, as they are determined from the <code>tidy_sem</code> object.

**Details**

The arguments `dataout`, `modelout`, and `run` are optional. If these are not specified, the model will be run in `tempdir`.

**Value**

An object of class `mplusObject`.

**Examples**

```
library(MplusAutomation)
model <- tidy_sem(iris, "\\.")
model <- measurement(model)
if(mplusAvailable() == 0){
  estimate_mplus(model)
}
```

---

est_sig	<i>Add significance asterisks to object</i>
---------	---

---

**Description**

Takes an object, and adds significance asterisks.

**Usage**

```
est_sig(x, digits = 2, sig = NULL)
```

**Arguments**

x	An object for which a method exists. This will be treated as numeric by the default method.
digits	Integer. The number of digits to round the estimate column to.
sig	Optional, a vector of p-values for the default method.

**Value**

A character vector of formatted estimates.

**Author(s)**

Caspar J. van Lissa

**See Also**

table\_results

Other Reporting tools: [conf\\_int\(\)](#), [table\\_fit\(\)](#), [table\\_results\(\)](#)

**Examples**

```
est_sig(c(.222, .3333), sig = c(.054, .045))
```

---

get\_data

*Extract data from tidy\_sem*

---

**Description**

Provides access to the data element of a tidy\_sem object. This can be used to return or assign to the data element.

**Usage**

```
get_data(x)
```

```
get_data(x) <- value
```

**Arguments**

x	Object of class tidy_sem.
value	A valid value for get_data(x).

**Value**

data.frame

**Examples**

```
dict <- tidy_sem(iris, split = "\\.")
get_data(dict)
```

---

get\_edges

*Extract edges from a SEM model object*


---

**Description**

Attempts to extract edges from a SEM model object, where edges are defined as regression paths and covariances between variables (nodes).

**Usage**

```
get_edges(x, label = "est_sig", ...)
```

**Arguments**

x	A model object of class <code>mplusObject</code> or <code>lavaan</code> .
label	Either a character, indicating which column to use for edge labels, or an expression. See Details. Defaults to "est_sig", which labels edges with the estimated value with significance asterisks, as obtained from <code>table_results</code> . See Details and examples for more information.
...	Additional parameters passed to <code>table_results</code> . For example, users can pass the <code>digits</code> argument to control the number of digits in the edge label, or pass the <code>columns</code> argument to retain auxiliary columns in the <code>tidy_edges</code> data.frame for further processing (see Examples).

**Details**

The function `get_edges` identifies all regression paths, latent variable definitions, and covariances in the model as edges. The output of `table_results` for those paths is used to label the edges.

**## Custom labels**

One way to create custom edge labels is by passing an expression to `label`. When an expression is passed to `label`, it is evaluated in the context of a `data.frame` containing the results of a call to `table_results` on the `x` argument.

Another way to create custom labels is by requesting auxiliary variables using the `columns` argument (which is passed to `table_results`), and then using these columns to construct a new label. See examples.

**Value**

An object of class 'tidy\_edges'

**Examples**

```

# Standard use
library(lavaan)
res <- sem("dist ~ speed", cars)
get_edges(res)

# Pass an expression to the 'label' argument for custom labels
get_edges(res, label = paste(est_sig, confint))

# Pass the argument 'columns' to table_results through '...' to retain
# auxiliary columns for further processing
edg <- get_edges(res, columns = c("est_sig", "confint"))
edg
edg <- within(edg, {label <- paste(est_sig, confint)})
edg

```

---

get\_fit

*Get fit indices from objects*


---

**Description**

Get fit indices from objects for which a method exists.

**Usage**

```
get_fit(x, ...)
```

**Arguments**

`x` An object for which a method exists.  
`...` further arguments to be passed to or from other methods.

**Value**

A data.frame.

**Examples**

```

## Not run:
df <- iris[, 1, drop = FALSE]
names(df) <- "x"
res <- mx_mixture(model = "x ~ m{C}*1
                        x ~~ v{C}*x", classes = 1, data = df)

table_fit(res)

## End(Not run)

```

---

get_layout.lavaan	<i>Generate graph layout</i>
-------------------	------------------------------

---

### Description

Generate a tidy\_layout for a SEM graph.

### Usage

```
## S3 method for class 'lavaan'
get_layout(x, ..., layout_algorithm = "layout_as_tree")

get_layout(x, ...)

## Default S3 method:
get_layout(x, ..., rows = NULL)
```

### Arguments

x	An object for which a method exists; currently, methods exist for character, lavaan, and mplus.model objects.
...	Character arguments corresponding to layout elements. Use node names, empty strings (""), or NA values.
layout_algorithm	Optional argument for fit model objects. Character string, indicating which igraph layout algorithm to apply to position the nodes. Defaults to "layout_as_tree"; see details for more options.
rows	Numeric, indicating the number of rows of the graph.

### Details

There are three ways to generate a layout:

1. Specify the layout in the call to get\_layout() by providing node names and the number of rows to create a layout matrix. Empty strings ("") or NA can be used for empty cells. See Example 1.
2. Call get\_layout() on a model object or tidy\_results object. It will use the function [layout\\_as\\_tree](#), or any other layout function from the igraph package, to generate a rudimentary layout. See Example 2.
3. Instead of using get\_layout(), just use a matrix or data.frame with your layout. For example, specify the layout in a spreadsheet program, and load it into R (see Example 3). Or, copy the layout to the clipboard from your spreadsheet program, and load it from the clipboard (see Example 4)

The layout algorithms imported from igraph are: c("layout\_as\_star", "layout\_as\_tree", "layout\_in\_circle", "layo... These can be used by specifying the optional argument layout\_algorithm = "".

**Value**

Object of class 'tidy\_layout'

**Examples**

```
# Example 1
get_layout("c", NA, "d",
           NA, "e", NA, rows = 2)

# Example 2
library(lavaan)
fit <- cfa(' visual =~ x1 + x2 + x3 ',
          data = HolzingerSwineford1939[1:50, ])
get_layout(fit)

## Not run:
# Example 3
# Here, we first write the layout to .csv, but you could create it in a
# spreadsheet program, and save the spreadsheet to .csv:
write.csv(matrix(c("c", "", "d", "", "e", ""), nrow = 2, byrow = TRUE),
          file = file.path(tempdir(), "example3.csv"), row.names = FALSE)
# Now, we load the .csv:
read.csv(file.path(tempdir(), "example3.csv"))

# Example 4
# For this example, make your layout in a spreadsheet program, select it, and
# copy to clipboard. Reading from the clipboard works differently in Windows
# and Mac. For this example, I used Microsoft Excel.
# On Windows, run:
read.table("clipboard", sep = "\t")
# On Mac, run:
read.table(pipe("pbpaste"), sep="\t")

## End(Not run)
```

---

get\_nodes

*Extract nodes from a SEM model object*

---

**Description**

Attempts to extract nodes from a SEM model object, where nodes are defined as observed or latent variables.

**Usage**

```
get_nodes(x, label = paste2(name, est_sig, sep = "\n"), ...)
```

**Arguments**

x	A model object of class <code>mplusObject</code> or <code>lavaan</code> .
label	Either a character, indicating which column to use for node labels, or an expression. See Details. Defaults to <code>paste(name, est_sig, sep = "\n"</code> , which gives the node name followed by the estimated value with significance asterisks.
...	Additional parameters passed to <code>table_results</code> . For example, users can pass the <code>digits</code> argument to control the number of digits in the node label, or pass the <code>columns</code> argument to retain auxiliary columns in the <code>tidy_nodes</code> <code>data.frame</code> for further processing (see Examples).

**Details**

The function `get_nodes` identifies all dependent and independent variables in the model as nodes. If a mean structure / intercepts are included in the model, the output of `table_results` for those means / intercepts is used to label the nodes.

## Custom labels

One way to create custom node labels is by passing an expression to `label`, as in the default value of the argument. When an expression is passed to `label`, it is evaluated in the context of a `data.frame` containing the results of a call to `table_results` on the `x` argument, with an additional column labeled `name`, which contains the node names.

Another way to create custom labels is by requesting auxiliary variables using the `columns` argument (which is passed to `table_results`), and then using these columns to construct a new label. See examples.

**Value**

An object of class `'tidy_nodes'`

**Examples**

```
# Standard use extracts node names and shape
# (rect for observed, oval for latent)
library(lavaan)
res <- sem("dist ~ speed", cars)
get_nodes(res)

# To label nodes with mean values, include meanstructure in the model
# Note that it is possible to pass the argument 'digits' to table_results
# through '...'
res <- sem("dist ~ speed", cars, meanstructure = TRUE)
get_nodes(res, digits = 3)

# Pass an expression to the 'label' argument for custom labels
get_nodes(res, label = paste0(name, " ", est_sig, "\n", confint))

# Pass the argument 'columns' to table_results through '...' to retain
# auxiliary columns for further processing
nod <- get_nodes(res, columns = c("est_sig", "confint"))
nod
```

```
nod <- within(nod, {label <- paste0(name, " ", est_sig, "\n", confint)})
nod
```

---

graph\_sem

*Render a graph*


---

## Description

Render a graph based on a layout, and either nodes and edges, or a model object.

## Usage

```
graph_sem(...)

## Default S3 method:
graph_sem(
  edges = NULL,
  layout = NULL,
  nodes = NULL,
  rect_width = 1.2,
  rect_height = 0.8,
  ellipses_width = 1,
  ellipses_height = 1,
  variance_diameter = 0.8,
  spacing_x = 2,
  spacing_y = 2,
  text_size = 4,
  curvature = 60,
  angle = NULL,
  fix_coord = FALSE,
  ...
)

## S3 method for class 'lavaan'
graph_sem(model, edges = NULL, layout = NULL, nodes = NULL, ...)

## S3 method for class 'MxModel'
graph_sem(model, edges = NULL, layout = NULL, nodes = NULL, ...)

## S3 method for class 'mplus.model'
graph_sem(model, edges = NULL, layout = NULL, nodes = NULL, ...)

## S3 method for class 'character'
graph_sem(...)

## S3 method for class 'mplusObject'
graph_sem(model, edges = NULL, layout = NULL, nodes = NULL, ...)
```



**Arguments**

...	Additional arguments passed to and from functions.
edges	Object of class 'tidy_edges', or a data.frame with (at least) the columns c("from", "to"), and optionally, c("arrow", "label", "connect_from", "connect_to", "curvature").
layout	A matrix (or data.frame) that describes the layout; see <a href="#">get_layout</a> .
nodes	Optional, object of class 'tidy_nodes', created with the <a href="#">get_nodes</a> function, or a data.frame with (at least) the column c("name"), and optionally, c("shape", "label"). If set to NULL (the default), nodes are inferred from the layout and edges arguments.
rect_width	Width of rectangles (used to display observed variables), Default: 1.2
rect_height	Height of rectangles (used to display observed variables), Default: 0.8
ellipses_width	Width of ellipses (used to display latent variables), Default: 1
ellipses_height	Height of ellipses (used to display latent variables), Default: 1
variance_diameter	Diameter of variance circles, Default: .8
spacing_x	Spacing between columns of the graph, Default: 1
spacing_y	Spacing between rows of the graph, Default: 1
text_size	Point size of text, Default: 4
curvature	Curvature of curved edges. The curve is a circle segment originating in a point that forms a triangle with the two connected points, with angles at the two connected points equal to curvature. To flip a curved edge, use a negative value for curvature. Default: 60
angle	Angle used to connect nodes by the top and bottom. Defaults to NULL, which means Euclidean distance is used to determine the shortest distance between node sides. A numeric value between 0-180 can be provided, where 0 means that only nodes with the same x-coordinates are connected top-to-bottom, and 180 means that all nodes are connected top-to-bottom.
fix_coord	Whether or not to fix the aspect ratio of the graph. Does not work with multi-group or multilevel models. Default: FALSE.
model	Instead of the edges argument, it is also possible to use the model argument and pass an object for which a method exists (e.g., <code>mplus.model</code> or <code>lavaan</code> ).

**Details**

The default interface simply Runs the functions [prepare\\_graph](#) and `plot`. The alternative interface first runs [get\\_nodes](#) and [get\\_edges](#) on the `model` argument.

**Value**

Object of class 'sem\_graph'

**Examples**

```
library(lavaan)
res <- sem("dist ~ speed", cars)
graph_sem(res)
```

---

if\_edit

*Conditionally edit a sem\_graph object*


---

**Description**

This function allows users to conditionally manipulate the edges and nodes of a `sem_graph` object. The generic function `if_edit` applies the expression `expr` to all rows of the nodes and edges data.frames for which `condition` is TRUE.

The wrapper functions documented in the Usage section have a hard-coded `expr` and `condition`; for example, `color_sig(color = "green")` colors all nodes and edges with `pval < .05` green. If no column exists for the assigned aesthetic (e.g., `color`), the wrappers assign the default argument (in this case, `color = "black"`) to all other nodes and edges.

**Usage**

```
if_edit(data, condition, expr, ...)

if_edges(data, condition, expr, ...)

if_nodes(data, condition, expr, ...)

## S3 method for class 'sem_graph'
if_edit(data, condition, expr, element = c("edges", "nodes"), ...)

all_sig(data, expr, ...)

hide_sig(data, ...)

show_sig(data, ...)

colour_sig(data, colour = "black", ...)

color_sig(data, color = "black", ...)

linetype_sig(data, linetype = 1, ...)

size_sig(data, size = 1, ...)

alpha_sig(data, alpha = 1, ...)

fill_sig(data, fill = "white", ...)
```

```
label_colour_sig(data, label_colour = "black", ...)
label_color_sig(data, label_color = "black", ...)
label_fill_sig(data, label_fill = "white", ...)
label_size_sig(data, label_size = 4, ...)
label_alpha_sig(data, label_alpha = 1, ...)
label_family_sig(data, label_family = "sans", ...)
label_fontface_sig(data, label_fontface = "plain", ...)
label_hjust_sig(data, label_hjust = "center", ...)
label_vjust_sig(data, label_vjust = "middle", ...)
label_lineheight_sig(data, label_lineheight = 1, ...)
label_location_sig(data, label_location = 0.5, ...)
all_nonsig(data, expr, ...)
hide_nonsig(data, ...)
show_nonsig(data, ...)
colour_nonsig(data, colour = "black", ...)
color_nonsig(data, color = "black", ...)
linetype_nonsig(data, linetype = 1, ...)
size_nonsig(data, size = 1, ...)
alpha_nonsig(data, alpha = 1, ...)
fill_nonsig(data, fill = "white", ...)
label_colour_nonsig(data, label_colour = "black", ...)
label_color_nonsig(data, label_color = "black", ...)
label_fill_nonsig(data, label_fill = "white", ...)
label_size_nonsig(data, label_size = 4, ...)
```

```
label_alpha_nonsig(data, label_alpha = 1, ...)  
label_family_nonsig(data, label_family = "sans", ...)  
label_fontface_nonsig(data, label_fontface = "plain", ...)  
label_hjust_nonsig(data, label_hjust = "center", ...)  
label_vjust_nonsig(data, label_vjust = "middle", ...)  
label_lineheight_nonsig(data, label_lineheight = 1, ...)  
label_location_nonsig(data, label_location = 0.5, ...)  
all_fixed(data, expr, ...)  
hide_fixed(data, ...)  
show_fixed(data, ...)  
colour_fixed(data, colour = "black", ...)  
color_fixed(data, color = "black", ...)  
linetype_fixed(data, linetype = 1, ...)  
size_fixed(data, size = 1, ...)  
alpha_fixed(data, alpha = 1, ...)  
fill_fixed(data, fill = "white", ...)  
label_colour_fixed(data, label_colour = "black", ...)  
label_color_fixed(data, label_color = "black", ...)  
label_fill_fixed(data, label_fill = "white", ...)  
label_size_fixed(data, label_size = 4, ...)  
label_alpha_fixed(data, label_alpha = 1, ...)  
label_family_fixed(data, label_family = "sans", ...)  
label_fontface_fixed(data, label_fontface = "plain", ...)  
label_hjust_fixed(data, label_hjust = "center", ...)
```

```
label_vjust_fixed(data, label_vjust = "middle", ...)  
label_lineheight_fixed(data, label_lineheight = 1, ...)  
label_location_fixed(data, label_location = 0.5, ...)  
all_pos(data, expr, ...)  
hide_pos(data, ...)  
show_pos(data, ...)  
colour_pos(data, colour = "black", ...)  
color_pos(data, color = "black", ...)  
linetype_pos(data, linetype = 1, ...)  
size_pos(data, size = 1, ...)  
alpha_pos(data, alpha = 1, ...)  
fill_pos(data, fill = "white", ...)  
label_colour_pos(data, label_colour = "black", ...)  
label_color_pos(data, label_color = "black", ...)  
label_fill_pos(data, label_fill = "white", ...)  
label_size_pos(data, label_size = 4, ...)  
label_alpha_pos(data, label_alpha = 1, ...)  
label_family_pos(data, label_family = "sans", ...)  
label_fontface_pos(data, label_fontface = "plain", ...)  
label_hjust_pos(data, label_hjust = "center", ...)  
label_vjust_pos(data, label_vjust = "middle", ...)  
label_lineheight_pos(data, label_lineheight = 1, ...)  
label_location_pos(data, label_location = 0.5, ...)  
all_neg(data, expr, ...)
```

```
hide_neg(data, ...)  
show_neg(data, ...)  
colour_neg(data, colour = "black", ...)  
color_neg(data, color = "black", ...)  
linetype_neg(data, linetype = 1, ...)  
size_neg(data, size = 1, ...)  
alpha_neg(data, alpha = 1, ...)  
fill_neg(data, fill = "white", ...)  
label_colour_neg(data, label_colour = "black", ...)  
label_color_neg(data, label_color = "black", ...)  
label_fill_neg(data, label_fill = "white", ...)  
label_size_neg(data, label_size = 4, ...)  
label_alpha_neg(data, label_alpha = 1, ...)  
label_family_neg(data, label_family = "sans", ...)  
label_fontface_neg(data, label_fontface = "plain", ...)  
label_hjust_neg(data, label_hjust = "center", ...)  
label_vjust_neg(data, label_vjust = "middle", ...)  
label_lineheight_neg(data, label_lineheight = 1, ...)  
label_location_neg(data, label_location = 0.5, ...)  
all_var(data, expr, ...)  
hide_var(data, ...)  
show_var(data, ...)  
colour_var(data, colour = "black", ...)  
color_var(data, color = "black", ...)
```

```
linetype_var(data, linetype = 1, ...)  
size_var(data, size = 1, ...)  
alpha_var(data, alpha = 1, ...)  
label_colour_var(data, label_colour = "black", ...)  
label_color_var(data, label_color = "black", ...)  
label_fill_var(data, label_fill = "white", ...)  
label_size_var(data, label_size = 4, ...)  
label_alpha_var(data, label_alpha = 1, ...)  
label_family_var(data, label_family = "sans", ...)  
label_fontface_var(data, label_fontface = "plain", ...)  
label_hjust_var(data, label_hjust = "center", ...)  
label_vjust_var(data, label_vjust = "middle", ...)  
label_lineheight_var(data, label_lineheight = 1, ...)  
all_cov(data, expr, ...)  
hide_cov(data, ...)  
show_cov(data, ...)  
colour_cov(data, colour = "black", ...)  
color_cov(data, color = "black", ...)  
linetype_cov(data, linetype = 1, ...)  
size_cov(data, size = 1, ...)  
alpha_cov(data, alpha = 1, ...)  
label_colour_cov(data, label_colour = "black", ...)  
label_color_cov(data, label_color = "black", ...)  
label_fill_cov(data, label_fill = "white", ...)
```

```
label_size_cov(data, label_size = 4, ...)  
label_alpha_cov(data, label_alpha = 1, ...)  
label_family_cov(data, label_family = "sans", ...)  
label_fontface_cov(data, label_fontface = "plain", ...)  
label_hjust_cov(data, label_hjust = "center", ...)  
label_vjust_cov(data, label_vjust = "middle", ...)  
label_lineheight_cov(data, label_lineheight = 1, ...)  
label_location_cov(data, label_location = 0.5, ...)  
all_reg(data, expr, ...)  
hide_reg(data, ...)  
show_reg(data, ...)  
colour_reg(data, colour = "black", ...)  
color_reg(data, color = "black", ...)  
linetype_reg(data, linetype = 1, ...)  
size_reg(data, size = 1, ...)  
alpha_reg(data, alpha = 1, ...)  
label_colour_reg(data, label_colour = "black", ...)  
label_color_reg(data, label_color = "black", ...)  
label_fill_reg(data, label_fill = "white", ...)  
label_size_reg(data, label_size = 4, ...)  
label_alpha_reg(data, label_alpha = 1, ...)  
label_family_reg(data, label_family = "sans", ...)  
label_fontface_reg(data, label_fontface = "plain", ...)  
label_hjust_reg(data, label_hjust = "center", ...)
```



```
label_vjust_reg(data, label_vjust = "middle", ...)  
label_lineheight_reg(data, label_lineheight = 1, ...)  
label_location_reg(data, label_location = 0.5, ...)  
all_load(data, expr, ...)  
hide_load(data, ...)  
show_load(data, ...)  
colour_load(data, colour = "black", ...)  
color_load(data, color = "black", ...)  
linetype_load(data, linetype = 1, ...)  
size_load(data, size = 1, ...)  
alpha_load(data, alpha = 1, ...)  
label_colour_load(data, label_colour = "black", ...)  
label_color_load(data, label_color = "black", ...)  
label_fill_load(data, label_fill = "white", ...)  
label_size_load(data, label_size = 4, ...)  
label_alpha_load(data, label_alpha = 1, ...)  
label_family_load(data, label_family = "sans", ...)  
label_fontface_load(data, label_fontface = "plain", ...)  
label_hjust_load(data, label_hjust = "center", ...)  
label_vjust_load(data, label_vjust = "middle", ...)  
label_lineheight_load(data, label_lineheight = 1, ...)  
label_location_load(data, label_location = 0.5, ...)  
all_obs(data, expr, ...)  
hide_obs(data, ...)
```

```
show_obs(data, ...)  
colour_obs(data, colour = "black", ...)  
color_obs(data, color = "black", ...)  
linetype_obs(data, linetype = 1, ...)  
size_obs(data, size = 1, ...)  
alpha_obs(data, alpha = 1, ...)  
fill_obs(data, fill = "white", ...)  
label_colour_obs(data, label_colour = "black", ...)  
label_color_obs(data, label_color = "black", ...)  
label_fill_obs(data, label_fill = "white", ...)  
label_size_obs(data, label_size = 4, ...)  
label_alpha_obs(data, label_alpha = 1, ...)  
label_family_obs(data, label_family = "sans", ...)  
label_fontface_obs(data, label_fontface = "plain", ...)  
label_hjust_obs(data, label_hjust = "center", ...)  
label_vjust_obs(data, label_vjust = "middle", ...)  
label_lineheight_obs(data, label_lineheight = 1, ...)  
all_latent(data, expr, ...)  
hide_latent(data, ...)  
show_latent(data, ...)  
colour_latent(data, colour = "black", ...)  
color_latent(data, color = "black", ...)  
linetype_latent(data, linetype = 1, ...)  
size_latent(data, size = 1, ...)
```

```
alpha_latent(data, alpha = 1, ...)
fill_latent(data, fill = "white", ...)
label_colour_latent(data, label_colour = "black", ...)
label_color_latent(data, label_color = "black", ...)
label_fill_latent(data, label_fill = "white", ...)
label_size_latent(data, label_size = 4, ...)
label_alpha_latent(data, label_alpha = 1, ...)
label_family_latent(data, label_family = "sans", ...)
label_fontface_latent(data, label_fontface = "plain", ...)
label_hjust_latent(data, label_hjust = "center", ...)
label_vjust_latent(data, label_vjust = "middle", ...)
label_lineheight_latent(data, label_lineheight = 1, ...)
all_sig_nodes(data, expr, ...)
hide_sig_nodes(data, ...)
show_sig_nodes(data, ...)
colour_sig_nodes(data, colour = "black", ...)
color_sig_nodes(data, color = "black", ...)
linetype_sig_nodes(data, linetype = 1, ...)
size_sig_nodes(data, size = 1, ...)
alpha_sig_nodes(data, alpha = 1, ...)
label_colour_sig_nodes(data, label_colour = "black", ...)
label_color_sig_nodes(data, label_color = "black", ...)
label_fill_sig_nodes(data, label_fill = "white", ...)
label_size_sig_nodes(data, label_size = 4, ...)
```

```
label_alpha_sig_nodes(data, label_alpha = 1, ...)
label_family_sig_nodes(data, label_family = "sans", ...)
label_fontface_sig_nodes(data, label_fontface = "plain", ...)
label_hjust_sig_nodes(data, label_hjust = "center", ...)
label_vjust_sig_nodes(data, label_vjust = "middle", ...)
label_lineheight_sig_nodes(data, label_lineheight = 1, ...)
all_nonsig_nodes(data, expr, ...)
hide_nonsig_nodes(data, ...)
show_nonsig_nodes(data, ...)
colour_nonsig_nodes(data, colour = "black", ...)
color_nonsig_nodes(data, color = "black", ...)
linetype_nonsig_nodes(data, linetype = 1, ...)
size_nonsig_nodes(data, size = 1, ...)
alpha_nonsig_nodes(data, alpha = 1, ...)
label_colour_nonsig_nodes(data, label_colour = "black", ...)
label_color_nonsig_nodes(data, label_color = "black", ...)
label_fill_nonsig_nodes(data, label_fill = "white", ...)
label_size_nonsig_nodes(data, label_size = 4, ...)
label_alpha_nonsig_nodes(data, label_alpha = 1, ...)
label_family_nonsig_nodes(data, label_family = "sans", ...)
label_fontface_nonsig_nodes(data, label_fontface = "plain", ...)
label_hjust_nonsig_nodes(data, label_hjust = "center", ...)
label_vjust_nonsig_nodes(data, label_vjust = "middle", ...)
label_lineheight_nonsig_nodes(data, label_lineheight = 1, ...)
```

```
all_fixed_nodes(data, expr, ...)
hide_fixed_nodes(data, ...)
show_fixed_nodes(data, ...)
colour_fixed_nodes(data, colour = "black", ...)
color_fixed_nodes(data, color = "black", ...)
linetype_fixed_nodes(data, linetype = 1, ...)
size_fixed_nodes(data, size = 1, ...)
alpha_fixed_nodes(data, alpha = 1, ...)
label_colour_fixed_nodes(data, label_colour = "black", ...)
label_color_fixed_nodes(data, label_color = "black", ...)
label_fill_fixed_nodes(data, label_fill = "white", ...)
label_size_fixed_nodes(data, label_size = 4, ...)
label_alpha_fixed_nodes(data, label_alpha = 1, ...)
label_family_fixed_nodes(data, label_family = "sans", ...)
label_fontface_fixed_nodes(data, label_fontface = "plain", ...)
label_hjust_fixed_nodes(data, label_hjust = "center", ...)
label_vjust_fixed_nodes(data, label_vjust = "middle", ...)
label_lineheight_fixed_nodes(data, label_lineheight = 1, ...)
all_pos_nodes(data, expr, ...)
hide_pos_nodes(data, ...)
show_pos_nodes(data, ...)
colour_pos_nodes(data, colour = "black", ...)
color_pos_nodes(data, color = "black", ...)
linetype_pos_nodes(data, linetype = 1, ...)
```

```
size_pos_nodes(data, size = 1, ...)
alpha_pos_nodes(data, alpha = 1, ...)
label_colour_pos_nodes(data, label_colour = "black", ...)
label_color_pos_nodes(data, label_color = "black", ...)
label_fill_pos_nodes(data, label_fill = "white", ...)
label_size_pos_nodes(data, label_size = 4, ...)
label_alpha_pos_nodes(data, label_alpha = 1, ...)
label_family_pos_nodes(data, label_family = "sans", ...)
label_fontface_pos_nodes(data, label_fontface = "plain", ...)
label_hjust_pos_nodes(data, label_hjust = "center", ...)
label_vjust_pos_nodes(data, label_vjust = "middle", ...)
label_lineheight_pos_nodes(data, label_lineheight = 1, ...)
all_neg_nodes(data, expr, ...)
hide_neg_nodes(data, ...)
show_neg_nodes(data, ...)
colour_neg_nodes(data, colour = "black", ...)
color_neg_nodes(data, color = "black", ...)
linetype_neg_nodes(data, linetype = 1, ...)
size_neg_nodes(data, size = 1, ...)
alpha_neg_nodes(data, alpha = 1, ...)
label_colour_neg_nodes(data, label_colour = "black", ...)
label_color_neg_nodes(data, label_color = "black", ...)
label_fill_neg_nodes(data, label_fill = "white", ...)
label_size_neg_nodes(data, label_size = 4, ...)
```

```
label_alpha_neg_nodes(data, label_alpha = 1, ...)
label_family_neg_nodes(data, label_family = "sans", ...)
label_fontface_neg_nodes(data, label_fontface = "plain", ...)
label_hjust_neg_nodes(data, label_hjust = "center", ...)
label_vjust_neg_nodes(data, label_vjust = "middle", ...)
label_lineheight_neg_nodes(data, label_lineheight = 1, ...)
all_sig_edges(data, expr, ...)
hide_sig_edges(data, ...)
show_sig_edges(data, ...)
colour_sig_edges(data, colour = "black", ...)
color_sig_edges(data, color = "black", ...)
linetype_sig_edges(data, linetype = 1, ...)
size_sig_edges(data, size = 1, ...)
alpha_sig_edges(data, alpha = 1, ...)
label_colour_sig_edges(data, label_colour = "black", ...)
label_color_sig_edges(data, label_color = "black", ...)
label_fill_sig_edges(data, label_fill = "white", ...)
label_size_sig_edges(data, label_size = 4, ...)
label_alpha_sig_edges(data, label_alpha = 1, ...)
label_family_sig_edges(data, label_family = "sans", ...)
label_fontface_sig_edges(data, label_fontface = "plain", ...)
label_hjust_sig_edges(data, label_hjust = "center", ...)
label_vjust_sig_edges(data, label_vjust = "middle", ...)
label_lineheight_sig_edges(data, label_lineheight = 1, ...)
```

```
all_nonsig_edges(data, expr, ...)
hide_nonsig_edges(data, ...)
show_nonsig_edges(data, ...)
colour_nonsig_edges(data, colour = "black", ...)
color_nonsig_edges(data, color = "black", ...)
linetype_nonsig_edges(data, linetype = 1, ...)
size_nonsig_edges(data, size = 1, ...)
alpha_nonsig_edges(data, alpha = 1, ...)
label_colour_nonsig_edges(data, label_colour = "black", ...)
label_color_nonsig_edges(data, label_color = "black", ...)
label_fill_nonsig_edges(data, label_fill = "white", ...)
label_size_nonsig_edges(data, label_size = 4, ...)
label_alpha_nonsig_edges(data, label_alpha = 1, ...)
label_family_nonsig_edges(data, label_family = "sans", ...)
label_fontface_nonsig_edges(data, label_fontface = "plain", ...)
label_hjust_nonsig_edges(data, label_hjust = "center", ...)
label_vjust_nonsig_edges(data, label_vjust = "middle", ...)
label_lineheight_nonsig_edges(data, label_lineheight = 1, ...)
all_fixed_edges(data, expr, ...)
hide_fixed_edges(data, ...)
show_fixed_edges(data, ...)
colour_fixed_edges(data, colour = "black", ...)
color_fixed_edges(data, color = "black", ...)
linetype_fixed_edges(data, linetype = 1, ...)
```



```
size_fixed_edges(data, size = 1, ...)
alpha_fixed_edges(data, alpha = 1, ...)
label_colour_fixed_edges(data, label_colour = "black", ...)
label_color_fixed_edges(data, label_color = "black", ...)
label_fill_fixed_edges(data, label_fill = "white", ...)
label_size_fixed_edges(data, label_size = 4, ...)
label_alpha_fixed_edges(data, label_alpha = 1, ...)
label_family_fixed_edges(data, label_family = "sans", ...)
label_fontface_fixed_edges(data, label_fontface = "plain", ...)
label_hjust_fixed_edges(data, label_hjust = "center", ...)
label_vjust_fixed_edges(data, label_vjust = "middle", ...)
label_lineheight_fixed_edges(data, label_lineheight = 1, ...)
all_pos_edges(data, expr, ...)
hide_pos_edges(data, ...)
show_pos_edges(data, ...)
colour_pos_edges(data, colour = "black", ...)
color_pos_edges(data, color = "black", ...)
linetype_pos_edges(data, linetype = 1, ...)
size_pos_edges(data, size = 1, ...)
alpha_pos_edges(data, alpha = 1, ...)
label_colour_pos_edges(data, label_colour = "black", ...)
label_color_pos_edges(data, label_color = "black", ...)
label_fill_pos_edges(data, label_fill = "white", ...)
label_size_pos_edges(data, label_size = 4, ...)
```

```
label_alpha_pos_edges(data, label_alpha = 1, ...)
label_family_pos_edges(data, label_family = "sans", ...)
label_fontface_pos_edges(data, label_fontface = "plain", ...)
label_hjust_pos_edges(data, label_hjust = "center", ...)
label_vjust_pos_edges(data, label_vjust = "middle", ...)
label_lineheight_pos_edges(data, label_lineheight = 1, ...)
all_neg_edges(data, expr, ...)
hide_neg_edges(data, ...)
show_neg_edges(data, ...)
colour_neg_edges(data, colour = "black", ...)
color_neg_edges(data, color = "black", ...)
linetype_neg_edges(data, linetype = 1, ...)
size_neg_edges(data, size = 1, ...)
alpha_neg_edges(data, alpha = 1, ...)
label_colour_neg_edges(data, label_colour = "black", ...)
label_color_neg_edges(data, label_color = "black", ...)
label_fill_neg_edges(data, label_fill = "white", ...)
label_size_neg_edges(data, label_size = 4, ...)
label_alpha_neg_edges(data, label_alpha = 1, ...)
label_family_neg_edges(data, label_family = "sans", ...)
label_fontface_neg_edges(data, label_fontface = "plain", ...)
label_hjust_neg_edges(data, label_hjust = "center", ...)
label_vjust_neg_edges(data, label_vjust = "middle", ...)
label_lineheight_neg_edges(data, label_lineheight = 1, ...)
```

**Arguments**

data	Object to manipulate.
condition	Expression that returns a logical vector when evaluated in the environment of data.
expr	Expression to perform on elements of data for which condition == TRUE.
...	Additional arguments passed to and from functions.
element	Character vector. The elements of the sem_graph to edit, defaults to c("edges", "nodes").
colour	Atomic character vector, indicating which colour to assign to the selected elements.
color	Atomic character vector, indicating which color to assign to the selected elements.
linetype	Atomic character vector, indicating which linetype to assign to the selected elements.
size	Atomic character vector, indicating which size to assign to the selected elements.
alpha	Atomic character vector, indicating which alpha to assign to the selected elements.
fill	Atomic character vector, indicating which fill to assign to the selected elements.
label_colour	Atomic character vector, indicating which label_colour to assign to the selected elements.
label_color	Atomic character vector, indicating which label_color to assign to the selected elements.
label_fill	Atomic character vector, indicating which label_fill to assign to the selected elements.
label_size	Atomic character vector, indicating which label_size to assign to the selected elements.
label_alpha	Atomic character vector, indicating which label_alpha to assign to the selected elements.
label_family	Atomic character vector, indicating which label_family to assign to the selected elements.
label_fontface	Atomic character vector, indicating which label_fontface to assign to the selected elements.
label_hjust	Atomic character vector, indicating which label_hjust to assign to the selected elements.
label_vjust	Atomic character vector, indicating which label_vjust to assign to the selected elements.
label_lineheight	Atomic character vector, indicating which label_lineheight to assign to the selected elements.
label_location	Atomic character vector, indicating which label_location to assign to the selected elements.

**Value**

Object of the same class as data.

**Examples**

```

library(lavaan)
res <- sem("dist ~ speed", cars, meanstructure = TRUE)
p <- prepare_graph(res)
out <- if_edit(p, condition = {pval < .05}, expr = {label = "sig"})
out <- if_edges(p, condition = {pval < .05}, expr = {label = "sig"})
out <- if_nodes(p, condition = {pval < .05}, expr = {label = "sig"})
out <- all_sig(p, expr = {label = "sig"})
out <- hide_sig(p)
out <- show_sig(p)
out <- colour_sig(p, { colour = "black" })
out <- color_sig(p, { color = "black" })
out <- linetype_sig(p, { linetype = 1 })
out <- size_sig(p, { size = 1 })
out <- alpha_sig(p, { alpha = 1 })
out <- fill_sig(p, { fill = "white" })
out <- label_colour_sig(p, { label_colour = "black" })
out <- label_color_sig(p, { label_color = "black" })
out <- label_fill_sig(p, { label_fill = "white" })
out <- label_size_sig(p, { label_size = 4 })
out <- label_alpha_sig(p, { label_alpha = 1 })
out <- label_family_sig(p, { label_family = "sans" })
out <- label_fontface_sig(p, { label_fontface = "plain" })
out <- label_hjust_sig(p, { label_hjust = "center" })
out <- label_vjust_sig(p, { label_vjust = "middle" })
out <- label_lineheight_sig(p, { label_lineheight = 1 })
out <- label_location_sig(p, { label_location = .5 })
out <- all_nonsig(p, expr = {label = "sig"})
out <- hide_nonsig(p)
out <- show_nonsig(p)
out <- colour_nonsig(p, { colour = "black" })
out <- color_nonsig(p, { color = "black" })
out <- linetype_nonsig(p, { linetype = 1 })
out <- size_nonsig(p, { size = 1 })
out <- alpha_nonsig(p, { alpha = 1 })
out <- fill_nonsig(p, { fill = "white" })
out <- label_colour_nonsig(p, { label_colour = "black" })
out <- label_color_nonsig(p, { label_color = "black" })
out <- label_fill_nonsig(p, { label_fill = "white" })
out <- label_size_nonsig(p, { label_size = 4 })
out <- label_alpha_nonsig(p, { label_alpha = 1 })
out <- label_family_nonsig(p, { label_family = "sans" })
out <- label_fontface_nonsig(p, { label_fontface = "plain" })
out <- label_hjust_nonsig(p, { label_hjust = "center" })
out <- label_vjust_nonsig(p, { label_vjust = "middle" })
out <- label_lineheight_nonsig(p, { label_lineheight = 1 })
out <- label_location_nonsig(p, { label_location = .5 })
out <- all_fixed(p, expr = {label = "sig"})

```

```
out <- hide_fixed(p)
out <- show_fixed(p)
out <- colour_fixed(p, { colour = "black" })
out <- color_fixed(p, { color = "black" })
out <- linetype_fixed(p, { linetype = 1 })
out <- size_fixed(p, { size = 1 })
out <- alpha_fixed(p, { alpha = 1 })
out <- fill_fixed(p, { fill = "white" })
out <- label_colour_fixed(p, { label_colour = "black" })
out <- label_color_fixed(p, { label_color = "black" })
out <- label_fill_fixed(p, { label_fill = "white" })
out <- label_size_fixed(p, { label_size = 4 })
out <- label_alpha_fixed(p, { label_alpha = 1 })
out <- label_family_fixed(p, { label_family = "sans" })
out <- label_fontface_fixed(p, { label_fontface = "plain" })
out <- label_hjust_fixed(p, { label_hjust = "center" })
out <- label_vjust_fixed(p, { label_vjust = "middle" })
out <- label_lineheight_fixed(p, { label_lineheight = 1 })
out <- label_location_fixed(p, { label_location = .5 })
out <- all_pos(p, expr = {label = "sig"})
out <- hide_pos(p)
out <- show_pos(p)
out <- colour_pos(p, { colour = "black" })
out <- color_pos(p, { color = "black" })
out <- linetype_pos(p, { linetype = 1 })
out <- size_pos(p, { size = 1 })
out <- alpha_pos(p, { alpha = 1 })
out <- fill_pos(p, { fill = "white" })
out <- label_colour_pos(p, { label_colour = "black" })
out <- label_color_pos(p, { label_color = "black" })
out <- label_fill_pos(p, { label_fill = "white" })
out <- label_size_pos(p, { label_size = 4 })
out <- label_alpha_pos(p, { label_alpha = 1 })
out <- label_family_pos(p, { label_family = "sans" })
out <- label_fontface_pos(p, { label_fontface = "plain" })
out <- label_hjust_pos(p, { label_hjust = "center" })
out <- label_vjust_pos(p, { label_vjust = "middle" })
out <- label_lineheight_pos(p, { label_lineheight = 1 })
out <- label_location_pos(p, { label_location = .5 })
out <- all_neg(p, expr = {label = "sig"})
out <- hide_neg(p)
out <- show_neg(p)
out <- colour_neg(p, { colour = "black" })
out <- color_neg(p, { color = "black" })
out <- linetype_neg(p, { linetype = 1 })
out <- size_neg(p, { size = 1 })
out <- alpha_neg(p, { alpha = 1 })
out <- fill_neg(p, { fill = "white" })
out <- label_colour_neg(p, { label_colour = "black" })
out <- label_color_neg(p, { label_color = "black" })
out <- label_fill_neg(p, { label_fill = "white" })
out <- label_size_neg(p, { label_size = 4 })
out <- label_alpha_neg(p, { label_alpha = 1 })
```

```

out <- label_family_neg(p, { label_family = "sans" })
out <- label_fontface_neg(p, { label_fontface = "plain" })
out <- label_hjust_neg(p, { label_hjust = "center" })
out <- label_vjust_neg(p, { label_vjust = "middle" })
out <- label_lineheight_neg(p, { label_lineheight = 1 })
out <- label_location_neg(p, { label_location = .5 })
out <- all_var(p, expr = {label = "sig"})
out <- hide_var(p)
out <- show_var(p)
out <- colour_var(p, { colour = "black" })
out <- color_var(p, { color = "black" })
out <- linetype_var(p, { linetype = 1 })
out <- size_var(p, { size = 1 })
out <- alpha_var(p, { alpha = 1 })
out <- label_colour_var(p, { label_colour = "black" })
out <- label_color_var(p, { label_color = "black" })
out <- label_fill_var(p, { label_fill = "white" })
out <- label_size_var(p, { label_size = 4 })
out <- label_alpha_var(p, { label_alpha = 1 })
out <- label_family_var(p, { label_family = "sans" })
out <- label_fontface_var(p, { label_fontface = "plain" })
out <- label_hjust_var(p, { label_hjust = "center" })
out <- label_vjust_var(p, { label_vjust = "middle" })
out <- label_lineheight_var(p, { label_lineheight = 1 })
out <- all_cov(p, expr = {label = "sig"})
out <- hide_cov(p)
out <- show_cov(p)
out <- colour_cov(p, { colour = "black" })
out <- color_cov(p, { color = "black" })
out <- linetype_cov(p, { linetype = 1 })
out <- size_cov(p, { size = 1 })
out <- alpha_cov(p, { alpha = 1 })
out <- label_colour_cov(p, { label_colour = "black" })
out <- label_color_cov(p, { label_color = "black" })
out <- label_fill_cov(p, { label_fill = "white" })
out <- label_size_cov(p, { label_size = 4 })
out <- label_alpha_cov(p, { label_alpha = 1 })
out <- label_family_cov(p, { label_family = "sans" })
out <- label_fontface_cov(p, { label_fontface = "plain" })
out <- label_hjust_cov(p, { label_hjust = "center" })
out <- label_vjust_cov(p, { label_vjust = "middle" })
out <- label_lineheight_cov(p, { label_lineheight = 1 })
out <- label_location_cov(p, { label_location = .5 })
out <- all_reg(p, expr = {label = "sig"})
out <- hide_reg(p)
out <- show_reg(p)
out <- colour_reg(p, { colour = "black" })
out <- color_reg(p, { color = "black" })
out <- linetype_reg(p, { linetype = 1 })
out <- size_reg(p, { size = 1 })
out <- alpha_reg(p, { alpha = 1 })
out <- label_colour_reg(p, { label_colour = "black" })
out <- label_color_reg(p, { label_color = "black" })

```

```

out <- label_fill_reg(p, { label_fill = "white" })
out <- label_size_reg(p, { label_size = 4 })
out <- label_alpha_reg(p, { label_alpha = 1 })
out <- label_family_reg(p, { label_family = "sans" })
out <- label_fontface_reg(p, { label_fontface = "plain" })
out <- label_hjust_reg(p, { label_hjust = "center" })
out <- label_vjust_reg(p, { label_vjust = "middle" })
out <- label_lineheight_reg(p, { label_lineheight = 1 })
out <- label_location_reg(p, { label_location = .5 })
out <- all_load(p, expr = {label = "sig"})
out <- hide_load(p)
out <- show_load(p)
out <- colour_load(p, { colour = "black" })
out <- color_load(p, { color = "black" })
out <- linetype_load(p, { linetype = 1 })
out <- size_load(p, { size = 1 })
out <- alpha_load(p, { alpha = 1 })
out <- label_colour_load(p, { label_colour = "black" })
out <- label_color_load(p, { label_color = "black" })
out <- label_fill_load(p, { label_fill = "white" })
out <- label_size_load(p, { label_size = 4 })
out <- label_alpha_load(p, { label_alpha = 1 })
out <- label_family_load(p, { label_family = "sans" })
out <- label_fontface_load(p, { label_fontface = "plain" })
out <- label_hjust_load(p, { label_hjust = "center" })
out <- label_vjust_load(p, { label_vjust = "middle" })
out <- label_lineheight_load(p, { label_lineheight = 1 })
out <- label_location_load(p, { label_location = .5 })
out <- all_obs(p, expr = {label = "sig"})
out <- hide_obs(p)
out <- show_obs(p)
out <- colour_obs(p, { colour = "black" })
out <- color_obs(p, { color = "black" })
out <- linetype_obs(p, { linetype = 1 })
out <- size_obs(p, { size = 1 })
out <- alpha_obs(p, { alpha = 1 })
out <- fill_obs(p, { fill = "white" })
out <- label_colour_obs(p, { label_colour = "black" })
out <- label_color_obs(p, { label_color = "black" })
out <- label_fill_obs(p, { label_fill = "white" })
out <- label_size_obs(p, { label_size = 4 })
out <- label_alpha_obs(p, { label_alpha = 1 })
out <- label_family_obs(p, { label_family = "sans" })
out <- label_fontface_obs(p, { label_fontface = "plain" })
out <- label_hjust_obs(p, { label_hjust = "center" })
out <- label_vjust_obs(p, { label_vjust = "middle" })
out <- label_lineheight_obs(p, { label_lineheight = 1 })
out <- all_latent(p, expr = {label = "sig"})
out <- hide_latent(p)
out <- show_latent(p)
out <- colour_latent(p, { colour = "black" })
out <- color_latent(p, { color = "black" })
out <- linetype_latent(p, { linetype = 1 })

```

```

out <- size_latent(p, { size = 1 })
out <- alpha_latent(p, { alpha = 1 })
out <- fill_latent(p, { fill = "white" })
out <- label_colour_latent(p, { label_colour = "black" })
out <- label_color_latent(p, { label_color = "black" })
out <- label_fill_latent(p, { label_fill = "white" })
out <- label_size_latent(p, { label_size = 4 })
out <- label_alpha_latent(p, { label_alpha = 1 })
out <- label_family_latent(p, { label_family = "sans" })
out <- label_fontface_latent(p, { label_fontface = "plain" })
out <- label_hjust_latent(p, { label_hjust = "center" })
out <- label_vjust_latent(p, { label_vjust = "middle" })
out <- label_lineheight_latent(p, { label_lineheight = 1 })
out <- all_sig(p, expr = {label = "sig"})
out <- hide_sig(p)
out <- show_sig(p)
out <- colour_sig(p, { colour = "black" })
out <- color_sig(p, { color = "black" })
out <- linetype_sig(p, { linetype = 1 })
out <- size_sig(p, { size = 1 })
out <- alpha_sig(p, { alpha = 1 })
out <- label_colour_sig(p, { label_colour = "black" })
out <- label_color_sig(p, { label_color = "black" })
out <- label_fill_sig(p, { label_fill = "white" })
out <- label_size_sig(p, { label_size = 4 })
out <- label_alpha_sig(p, { label_alpha = 1 })
out <- label_family_sig(p, { label_family = "sans" })
out <- label_fontface_sig(p, { label_fontface = "plain" })
out <- label_hjust_sig(p, { label_hjust = "center" })
out <- label_vjust_sig(p, { label_vjust = "middle" })
out <- label_lineheight_sig(p, { label_lineheight = 1 })
out <- all_nonsig(p, expr = {label = "sig"})
out <- hide_nonsig(p)
out <- show_nonsig(p)
out <- colour_nonsig(p, { colour = "black" })
out <- color_nonsig(p, { color = "black" })
out <- linetype_nonsig(p, { linetype = 1 })
out <- size_nonsig(p, { size = 1 })
out <- alpha_nonsig(p, { alpha = 1 })
out <- label_colour_nonsig(p, { label_colour = "black" })
out <- label_color_nonsig(p, { label_color = "black" })
out <- label_fill_nonsig(p, { label_fill = "white" })
out <- label_size_nonsig(p, { label_size = 4 })
out <- label_alpha_nonsig(p, { label_alpha = 1 })
out <- label_family_nonsig(p, { label_family = "sans" })
out <- label_fontface_nonsig(p, { label_fontface = "plain" })
out <- label_hjust_nonsig(p, { label_hjust = "center" })
out <- label_vjust_nonsig(p, { label_vjust = "middle" })
out <- label_lineheight_nonsig(p, { label_lineheight = 1 })
out <- all_fixed(p, expr = {label = "sig"})
out <- hide_fixed(p)
out <- show_fixed(p)
out <- colour_fixed(p, { colour = "black" })

```



```
out <- color_fixed(p, { color = "black" })
out <- linetype_fixed(p, { linetype = 1 })
out <- size_fixed(p, { size = 1 })
out <- alpha_fixed(p, { alpha = 1 })
out <- label_colour_fixed(p, { label_colour = "black" })
out <- label_color_fixed(p, { label_color = "black" })
out <- label_fill_fixed(p, { label_fill = "white" })
out <- label_size_fixed(p, { label_size = 4 })
out <- label_alpha_fixed(p, { label_alpha = 1 })
out <- label_family_fixed(p, { label_family = "sans" })
out <- label_fontface_fixed(p, { label_fontface = "plain" })
out <- label_hjust_fixed(p, { label_hjust = "center" })
out <- label_vjust_fixed(p, { label_vjust = "middle" })
out <- label_lineheight_fixed(p, { label_lineheight = 1 })
out <- all_pos(p, expr = {label = "sig"})
out <- hide_pos(p)
out <- show_pos(p)
out <- colour_pos(p, { colour = "black" })
out <- color_pos(p, { color = "black" })
out <- linetype_pos(p, { linetype = 1 })
out <- size_pos(p, { size = 1 })
out <- alpha_pos(p, { alpha = 1 })
out <- label_colour_pos(p, { label_colour = "black" })
out <- label_color_pos(p, { label_color = "black" })
out <- label_fill_pos(p, { label_fill = "white" })
out <- label_size_pos(p, { label_size = 4 })
out <- label_alpha_pos(p, { label_alpha = 1 })
out <- label_family_pos(p, { label_family = "sans" })
out <- label_fontface_pos(p, { label_fontface = "plain" })
out <- label_hjust_pos(p, { label_hjust = "center" })
out <- label_vjust_pos(p, { label_vjust = "middle" })
out <- label_lineheight_pos(p, { label_lineheight = 1 })
out <- all_neg(p, expr = {label = "sig"})
out <- hide_neg(p)
out <- show_neg(p)
out <- colour_neg(p, { colour = "black" })
out <- color_neg(p, { color = "black" })
out <- linetype_neg(p, { linetype = 1 })
out <- size_neg(p, { size = 1 })
out <- alpha_neg(p, { alpha = 1 })
out <- label_colour_neg(p, { label_colour = "black" })
out <- label_color_neg(p, { label_color = "black" })
out <- label_fill_neg(p, { label_fill = "white" })
out <- label_size_neg(p, { label_size = 4 })
out <- label_alpha_neg(p, { label_alpha = 1 })
out <- label_family_neg(p, { label_family = "sans" })
out <- label_fontface_neg(p, { label_fontface = "plain" })
out <- label_hjust_neg(p, { label_hjust = "center" })
out <- label_vjust_neg(p, { label_vjust = "middle" })
out <- label_lineheight_neg(p, { label_lineheight = 1 })
out <- all_sig(p, expr = {label = "sig"})
out <- hide_sig(p)
out <- show_sig(p)
```

```

out <- colour_sig(p, { colour = "black" })
out <- color_sig(p, { color = "black" })
out <- linetype_sig(p, { linetype = 1 })
out <- size_sig(p, { size = 1 })
out <- alpha_sig(p, { alpha = 1 })
out <- label_colour_sig(p, { label_colour = "black" })
out <- label_color_sig(p, { label_color = "black" })
out <- label_fill_sig(p, { label_fill = "white" })
out <- label_size_sig(p, { label_size = 4 })
out <- label_alpha_sig(p, { label_alpha = 1 })
out <- label_family_sig(p, { label_family = "sans" })
out <- label_fontface_sig(p, { label_fontface = "plain" })
out <- label_hjust_sig(p, { label_hjust = "center" })
out <- label_vjust_sig(p, { label_vjust = "middle" })
out <- label_lineheight_sig(p, { label_lineheight = 1 })
out <- all_nonsig(p, expr = {label = "sig"})
out <- hide_nonsig(p)
out <- show_nonsig(p)
out <- colour_nonsig(p, { colour = "black" })
out <- color_nonsig(p, { color = "black" })
out <- linetype_nonsig(p, { linetype = 1 })
out <- size_nonsig(p, { size = 1 })
out <- alpha_nonsig(p, { alpha = 1 })
out <- label_colour_nonsig(p, { label_colour = "black" })
out <- label_color_nonsig(p, { label_color = "black" })
out <- label_fill_nonsig(p, { label_fill = "white" })
out <- label_size_nonsig(p, { label_size = 4 })
out <- label_alpha_nonsig(p, { label_alpha = 1 })
out <- label_family_nonsig(p, { label_family = "sans" })
out <- label_fontface_nonsig(p, { label_fontface = "plain" })
out <- label_hjust_nonsig(p, { label_hjust = "center" })
out <- label_vjust_nonsig(p, { label_vjust = "middle" })
out <- label_lineheight_nonsig(p, { label_lineheight = 1 })
out <- all_fixed(p, expr = {label = "sig"})
out <- hide_fixed(p)
out <- show_fixed(p)
out <- colour_fixed(p, { colour = "black" })
out <- color_fixed(p, { color = "black" })
out <- linetype_fixed(p, { linetype = 1 })
out <- size_fixed(p, { size = 1 })
out <- alpha_fixed(p, { alpha = 1 })
out <- label_colour_fixed(p, { label_colour = "black" })
out <- label_color_fixed(p, { label_color = "black" })
out <- label_fill_fixed(p, { label_fill = "white" })
out <- label_size_fixed(p, { label_size = 4 })
out <- label_alpha_fixed(p, { label_alpha = 1 })
out <- label_family_fixed(p, { label_family = "sans" })
out <- label_fontface_fixed(p, { label_fontface = "plain" })
out <- label_hjust_fixed(p, { label_hjust = "center" })
out <- label_vjust_fixed(p, { label_vjust = "middle" })
out <- label_lineheight_fixed(p, { label_lineheight = 1 })
out <- all_pos(p, expr = {label = "sig"})
out <- hide_pos(p)

```

```

out <- show_pos(p)
out <- colour_pos(p, { colour = "black" })
out <- color_pos(p, { color = "black" })
out <- linetype_pos(p, { linetype = 1 })
out <- size_pos(p, { size = 1 })
out <- alpha_pos(p, { alpha = 1 })
out <- label_colour_pos(p, { label_colour = "black" })
out <- label_color_pos(p, { label_color = "black" })
out <- label_fill_pos(p, { label_fill = "white" })
out <- label_size_pos(p, { label_size = 4 })
out <- label_alpha_pos(p, { label_alpha = 1 })
out <- label_family_pos(p, { label_family = "sans" })
out <- label_fontface_pos(p, { label_fontface = "plain" })
out <- label_hjust_pos(p, { label_hjust = "center" })
out <- label_vjust_pos(p, { label_vjust = "middle" })
out <- label_lineheight_pos(p, { label_lineheight = 1 })
out <- all_neg(p, expr = {label = "sig"})
out <- hide_neg(p)
out <- show_neg(p)
out <- colour_neg(p, { colour = "black" })
out <- color_neg(p, { color = "black" })
out <- linetype_neg(p, { linetype = 1 })
out <- size_neg(p, { size = 1 })
out <- alpha_neg(p, { alpha = 1 })
out <- label_colour_neg(p, { label_colour = "black" })
out <- label_color_neg(p, { label_color = "black" })
out <- label_fill_neg(p, { label_fill = "white" })
out <- label_size_neg(p, { label_size = 4 })
out <- label_alpha_neg(p, { label_alpha = 1 })
out <- label_family_neg(p, { label_family = "sans" })
out <- label_fontface_neg(p, { label_fontface = "plain" })
out <- label_hjust_neg(p, { label_hjust = "center" })
out <- label_vjust_neg(p, { label_vjust = "middle" })
out <- label_lineheight_neg(p, { label_lineheight = 1 })

```

---

lsub

*Apply pattern replacement over a vector*


---

## Description

lsub returns a list of the same length as replacement, each element of which is the result of applying `gsub` to `x` using `lapply`.

## Usage

```
lsub(x, replacement = NULL, pattern = "{C}", fixed = TRUE, ...)
```

**Arguments**

<code>x</code>	A character vector where matches are sought.
<code>replacement</code>	a character vector of length 1 or more. Each element is applied to <code>x</code> in turn. Default: <code>NULL</code>
<code>pattern</code>	A character string containing a regular expression (or character string when <code>fixed = TRUE</code> ). Default: <code>'{C}'</code> .
<code>fixed</code>	logical. If <code>TRUE</code> , <code>pattern</code> is a string to be matched as is. Default: <code>TRUE</code>
<code>...</code>	Parameters passed on to <code>gsub</code> .

**Value**

A list of results returned by `gsub`.

**Examples**

```
lsub("a{C}", 1:3)
```

---

measurement

*Generate syntax for a measurement model*

---

**Description**

Generate syntax for a measurement model for latent variables. This function relies on `add_paths` to generate syntax.

**Usage**

```
measurement(x, ...)
```

**Arguments**

<code>x</code>	An object for which a method exists, including <code>tidy_sem</code> (generated using <code>dictionary</code> , or <code>data.frame</code> (for which <code>dictionary</code> will be run first).
<code>...</code>	Additional parameters passed to <code>add_paths</code> .

**Value**

An object of class `tidy_sem`.

**Examples**

```
dict <- tidy_sem(c("bfi_1", "bfi_2", "bfi_3", "bfi_4", "bfi_5"))
measurement(dict)
```

---

mixture_starts	<i>Automatically set starting values for an OpenMx mixture model</i>
----------------	--

---

### Description

Automatically set starting values for an OpenMx mixture model. This function was designed to work with mixture models created using tidySEM functions like `mx_mixture`, and may not work with other `mxModels`.

### Usage

```
mixture_starts(model, splits, ...)
```

### Arguments

model	A mixture model of class <code>mxModel</code> .
splits	Optional. A numeric vector of length equal to the number of rows in the <code>mxData</code> used in the <code>model</code> object. The data will be split by this vector. See Details for the default setting and possible alternatives.
...	Additional arguments, passed to functions.

### Details

Starting values are derived by the following procedure:

1. The mixture model is converted to a multi-group model.
2. The data are split along `splits`, and assigned to the corresponding groups of the multi-group model.
3. The multi-group model is run, and the final values of each group are assigned to the corresponding mixture component as starting values.
4. The mixture model is returned with these starting values.

If the argument `splits` is not provided, the function will call `kmeans(x = data, centers = classes)$cluster`, where `data` is extracted from the `model` argument.

Sensible ways to split the data include:

- Using Hierarchical clustering: `cutree(hclust(dist(data)), k = classes)`
- Using K-means clustering: `kmeans(x = data, centers = classes)$cluster`
- Using agglomerative hierarchical clustering: `hclass(hc(data = data), G = classes)[, 1]`
- Using a random split: `sample.int(n = classes, size = nrow(data), replace = TRUE)`

### Value

Returns an `mxModel` with starting values.

## References

Shireman, E., Steinley, D. & Brusco, M.J. Examining the effect of initialization strategies on the performance of Gaussian mixture modeling. *Behav Res* 49, 282–293 (2017). <doi:10.3758/s13428-015-0697-6>

## Examples

```
## Not run:
df <- iris[, 1, drop = FALSE]
names(df) <- "x"
mod <- mx_mixture(model = "x ~ m{C}*1
                        x ~~ v{C}*x",
                  classes = 2,
                  data = df,
                  run = FALSE)

mod <- mixture_starts(mod)

## End(Not run)
```

---

mplus\_expand\_names      *Expand abbreviated Mplus variable names*

---

## Description

Expand the Mplus syntax for abbreviating lists of variable names.

## Usage

```
mplus_expand_names(x)
```

## Arguments

x                      Atomic character string containing the variable names section of an Mplus syntax file.

## Value

Character vector of names.

## Examples

```
mplus_expand_names("test1-test12")
```



---

 mx\_mixture

*Estimate mixture models using OpenMx*


---

### Description

Dynamically creates a batch of mixture models, with intelligent defaults. See Details for more information.

### Usage

```
mx_mixture(model, classes = 1L, data = NULL, run = TRUE, ...)
```

### Arguments

model	Syntax for the model; either a character string, or a list of character strings, or a list of mxModel objects. See Details.
classes	A vector of integers, indicating which class solutions to generate. Defaults to 1L. E.g., classes = 1:6, classes = c(1:4, 6:8).
data	The data.frame to be used for model fitting.
run	Logical, whether or not to run the model. If run = TRUE, the function calls <a href="#">mixture_starts</a> and <a href="#">run_mx</a> .
...	Additional arguments, passed to functions.

### Details

Model syntax can be specified in three ways, for ease of use and flexibility:

1. An atomic character string with lavaan syntax. Within this syntax, the character string {C} is dynamically substituted with the correct class number using [lsub](#), for example to set unique parameter labels for each class, or to specify equality constraints. E.g.,  $x \sim m\{C\}*1$  will be expanded to  $x \sim m1*1$  and  $x \sim m2*1$  when classes = 2. The resulting syntax for each class will be converted to an mxModel using [as\\_ram](#).
2. A list of character strings with lavaan syntax. Each item of the list will be converted to a class-specific mxModel using [as\\_ram](#).
3. A list of mxModel objects, specified by the user.

### Value

Returns an [mxModel](#).



**Examples**

```
## Not run:
# Example 1: Dynamic model generation using {C}
df <- iris[, 1, drop = FALSE]
names(df) <- "x"
mx_mixture(model = "x ~ m{C}*1
                 x ~~ v{C}*x", classes = 1, data = df)
# Example 2: Manually specified class-specific models
df <- iris[1:2]
names(df) <- c("x", "y")
mx_mixture(model = list("y ~ a*x",
                       "y ~ b*x"),
           meanstructure = TRUE,
           data = df) -> res

# Example 3: Latent growth model
df <- empathy[1:6]
mx_mixture(model = "i =~ 1*ec1 + 1*ec2 + 1*ec3 + 1*ec4 + 1*ec5 + 1*ec6
                  s =~ 0*ec1 + 1*ec2 + 2*ec3 + 3*ec4 + 4*ec5 + 5*ec6",
           classes = 2,
           data = df) -> res

## End(Not run)
```

mx\_profiles

*Estimate latent profile analyses using OpenMx***Description**

This function is a wrapper around `mx_mixture` to simplify the specification of latent profile models, also known as finite mixture models. By default, the function estimates free means for all observed variables across classes.

**Usage**

```
mx_profiles(
  data = NULL,
  classes = 1L,
  variances = "equal",
  covariances = "zero",
  run = TRUE,
  ...
)
```

**Arguments**

`data` The data.frame to be used for model fitting.

`classes` A vector of integers, indicating which class solutions to generate. Defaults to 1L. E.g., `classes = 1:6`,

variances	Character vector. Specifies which variance components to estimate. Defaults to "equal" (constrain variances across classes); the other option is "varying" (estimate variances freely across classes). Each element of this vector refers to one of the models you wish to run.
covariances	Character vector. Specifies which covariance components to estimate. Defaults to "zero" (covariances constrained to zero; this corresponds to an assumption of conditional independence of the indicators); other options are "equal" (covariances between items constrained to be equal across classes), and "varying" (free covariances across classes). <code>classes = c(1:4, 6:8)</code> .
run	Logical, whether or not to run the model. If <code>run = TRUE</code> , the function calls <code>mixture_starts</code> and <code>run_mx</code> .
...	Additional arguments, passed to functions.

**Value**

Returns an `mxModel`.

**Examples**

```
## Not run:
data("empathy")
df <- empathy[1:6]
mx_profiles(data = df,
            classes = 2) -> res

## End(Not run)
```

---

nodes *Extract nodes from sem\_graph*

---

**Description**

Provides access to the nodes element of a `sem_graph` object. This can be used to return or assign to the nodes element.

**Usage**

```
nodes(x)
```

```
nodes(x) <- value
```

**Arguments**

`x` Object of class `sem_graph`.

`value` A valid value for `nodes(x)`.

**Value**

data.frame

**Examples**

```
edg <- data.frame(from = "x", to = "y")
p <- prepare_graph(edges = edg, layout = get_layout("x", "y", rows = 1))
nodes(p)
```

---

paste2

*Concatenate Strings while omitting NA*

---

**Description**

Concatenate vectors after converting to character and removing NA values. See [paste](#).

**Usage**

```
paste2(..., sep = " ", collapse = NULL, na.rm = TRUE)
```

**Arguments**

...	one or more R objects, to be converted to character vectors.
sep	a character string to separate the terms. Not NA_character_.
collapse	an optional character string to separate the results. Not NA_character_.
na.rm	logical, indicating whether NA values should be stripped before concatenation. Not NA_character_.

**Value**

A character vector of the concatenated values.

**Examples**

```
paste2("word", NA)
```

---

plot\_bivariate      *Create correlation plots for a mixture model*

---

### Description

Creates a faceted plot of two-dimensional correlation plots and unidimensional density plots for a single mixture model.

### Usage

```
plot_bivariate(  
  x,  
  variables = NULL,  
  sd = TRUE,  
  cors = TRUE,  
  rawdata = TRUE,  
  bw = FALSE,  
  alpha_range = c(0, 0.1),  
  return_list = FALSE  
)
```

### Arguments

x	An object for which a method exists.
variables	Which variables to plot. If NULL, plots all variables that are present in the model.
sd	Logical. Whether to show the estimated standard deviations as lines emanating from the cluster centroid.
cors	Logical. Whether to show the estimated correlation (standardized covariance) as ellipses surrounding the cluster centroid.
rawdata	Logical. Whether to plot raw data, weighted by posterior class probability.
bw	Logical. Whether to make a black and white plot (for print) or a color plot. Defaults to FALSE, because these density plots are hard to read in black and white.
alpha_range	Numeric vector (0-1). Sets the transparency of geom_density and geom_point.
return_list	Logical. Whether to return a list of ggplot objects, or just the final plot. Defaults to FALSE.

### Value

An object of class 'ggplot'.

### Author(s)

Caspar J. van Lissa

**Examples**

```
iris_sample <- iris[, c("Sepal.Length", "Sepal.Width")]
names(iris_sample) <- c("x", "y")
res <- mx_profiles(iris_sample, classes = 2)
plot_bivariate(res)
```

---

plot_density	<i>Create density plots for mixture models</i>
--------------	--

---

**Description**

Creates mixture density plots. For each variable, a Total density plot will be shown, along with separate density plots for each latent class, where cases are weighted by the posterior probability of being assigned to that class.

**Usage**

```
plot_density(
  x,
  variables = NULL,
  bw = FALSE,
  conditional = FALSE,
  alpha = 0.2,
  facet_labels = NULL
)
```

**Arguments**

x	Object for which a method exists.
variables	Which variables to plot. If NULL, plots all variables that are present in all models.
bw	Logical. Whether to make a black and white plot (for print) or a color plot. Defaults to FALSE, because these density plots are hard to read in black and white.
conditional	Logical. Whether to show a conditional density plot (surface area is divided among the latent classes), or a classic density plot (surface area of the total density plot is equal to one, and is divided among the classes).
alpha	Numeric (0-1). Only used when bw and conditional are FALSE. Sets the transparency of geom_density, so that classes with a small number of cases remain visible.
facet_labels	Named character vector, the names of which should correspond to the facet labels one wishes to rename, and the values of which provide new names for these facets. For example, to rename variables, in the example with the 'iris' data below, one could specify: facet_labels = c("Pet_leng" = "Petal length").

**Value**

An object of class 'ggplot'.

**Author(s)**

Caspar J. van Lissa

**Examples**

```
## Not run:
dat <-
  iris[, c("Sepal.Length", "Sepal.Width", "Petal.Length", "Petal.Width")]
names(dat) <- paste0("x", 1:4)
res <- mx_profiles(dat, 1:3)
plot_density(res)

## End(Not run)
```

---

plot\_profiles

*Create latent profile plots*

---

**Description**

Creates a profile plot (ribbon plot) according to best practices, focusing on the visualization of classification uncertainty by showing:

1. Bars reflecting a confidence interval for the class centroids
2. Boxes reflecting the standard deviations within each class; a box encompasses +/- 64% of the observations in a normal distribution
3. Raw data, whose transparency is weighted by the posterior class probability, such that each observation is most clearly visible for the class it is most likely to be a member of.

**Usage**

```
plot_profiles(
  x,
  variables = NULL,
  ci = 0.95,
  sd = TRUE,
  add_line = FALSE,
  rawdata = TRUE,
  bw = FALSE,
  alpha_range = c(0, 0.1),
  ...
)

## Default S3 method:
```

```

plot_profiles(
  x,
  variables = NULL,
  ci = 0.95,
  sd = TRUE,
  add_line = FALSE,
  rawdata = TRUE,
  bw = FALSE,
  alpha_range = c(0, 0.1),
  ...
)

```

### Arguments

x	An object containing the results of a mixture model analysis.
variables	A character vectors with the names of the variables to be plotted (optional).
ci	Numeric. What confidence interval should the error bars span? Defaults to a 95% confidence interval. Set to NULL to remove error bars.
sd	Logical. Whether to display a box encompassing +/- 1SD Defaults to TRUE.
add_line	Logical. Whether to display a line, connecting cluster centroids belonging to the same latent class. Defaults to FALSE, as it is not recommended to imply connectivity between the different variables on the X-axis.
rawdata	Should raw data be plotted in the background? Setting this to TRUE might result in long plotting times.
bw	Logical. Should the plot be black and white (for print), or color?
alpha_range	The minimum and maximum values of alpha (transparency) for the raw data. Minimum should be 0; lower maximum values of alpha can help reduce overplotting.
...	Arguments passed to and from other functions.

### Value

An object of class 'ggplot'.

### Author(s)

Caspar J. van Lissa

### Examples

```

df_plot <- data.frame(Variable = "x1",
  Class = "class1",
  Classes = 1,
  Model = "equal var 1",
  Value = 3.48571428571429,
  se = 0.426092805342181,
  Value.Variances = 3.81265306156537,
  se.Variances = 1.17660769119959)

```

```
plot_profiles(list(df_plot = df_plot, df_raw = NULL),
ci = NULL, sd = FALSE, add_line = FALSE,
rawdata = FALSE, bw = FALSE)
```

---

```
prepare_graph
```

```
Prepare graph data
```

---

## Description

Prepare an object of class `sem_graph`, containing data objects that can be rendered into a SEM graph. Using this function allows users to manually change the default graph specification before plotting it. Input consists of (at least) a layout, and either nodes and edges, or a model object.

## Usage

```
prepare_graph(...)

## Default S3 method:
prepare_graph(
  edges = NULL,
  layout = NULL,
  nodes = NULL,
  rect_width = 1.2,
  rect_height = 0.8,
  ellipses_width = 1,
  ellipses_height = 1,
  variance_diameter = 0.8,
  spacing_x = 2,
  spacing_y = 2,
  text_size = 4,
  curvature = 60,
  angle = NULL,
  fix_coord = FALSE,
  ...
)

## S3 method for class 'lavaan'
prepare_graph(
  model,
  edges = get_edges(x = model),
  layout = get_layout(x = model),
  nodes = get_nodes(x = model),
  ...
)

## S3 method for class 'MxModel'
prepare_graph(
```



```

    model,
    edges = get_edges(x = model),
    layout = get_layout(x = model),
    nodes = get_nodes(x = model),
    ...
)

## S3 method for class 'character'
prepare_graph(...)

## S3 method for class 'mplus.model'
prepare_graph(
  model,
  edges = get_edges(x = model),
  layout = get_layout(x = model),
  nodes = get_nodes(x = model),
  ...
)

## S3 method for class 'mplusObject'
prepare_graph(
  model,
  edges = get_edges(x = model),
  layout = get_layout(x = model),
  nodes = get_nodes(x = model),
  ...
)

```

## Arguments

...	Additional arguments passed to and from functions.
edges	Object of class 'tidy_edges', or a data.frame with (at least) the columns c("from", "to"), and optionally, c("arrow", "label", "connect_from", "connect_to", "curvature").
layout	A matrix (or data.frame) that describes the layout; see <a href="#">get_layout</a> .
nodes	Optional, object of class 'tidy_nodes', created with the <a href="#">get_nodes</a> function, or a data.frame with (at least) the column c("name"), and optionally, c("shape", "label"). If set to NULL (the default), nodes are inferred from the layout and edges arguments.
rect_width	Width of rectangles (used to display observed variables), Default: 1.2
rect_height	Height of rectangles (used to display observed variables), Default: 0.8
ellipses_width	Width of ellipses (used to display latent variables), Default: 1
ellipses_height	Height of ellipses (used to display latent variables), Default: 1
variance_diameter	Diameter of variance circles, Default: .8
spacing_x	Spacing between columns of the graph, Default: 1

spacing_y	Spacing between rows of the graph, Default: 1
text_size	Point size of text, Default: 4
curvature	Curvature of curved edges. The curve is a circle segment originating in a point that forms a triangle with the two connected points, with angles at the two connected points equal to curvature. To flip a curved edge, use a negative value for curvature. Default: 60
angle	Angle used to connect nodes by the top and bottom. Defaults to NULL, which means Euclidean distance is used to determine the shortest distance between node sides. A numeric value between 0-180 can be provided, where 0 means that only nodes with the same x-coordinates are connected top-to-bottom, and 180 means that all nodes are connected top-to-bottom.
fix_coord	Whether or not to fix the aspect ratio of the graph. Does not work with multi-group or multilevel models. Default: FALSE.
model	Instead of the edges argument, it is also possible to use the model argument and pass an object for which a method exists (e.g., <code>mplus.model</code> or <code>lavaan</code> ).

**Value**

Object of class 'sem\_graph'

**Examples**

```
library(lavaan)
res <- sem("dist ~ speed", cars)
prepare_graph(res)
```

---

run\_lavaan

*Run as lavaan model*

---

**Description**

This convenience function runs objects for which a method exists using lavaan. It is intended for use with `tidySEM`, and passes the `$syntax` and `$data` elements of a `tidy_sem` object on to [lavaan](#).

**Usage**

```
run_lavaan(x, ...)
```

**Arguments**

`x` An object for which a method exists.  
`...` Parameters passed on to other functions.

**Value**

Returns a lavaan object.

**Examples**

```
df <- iris[1:3]
names(df) <- paste0("X_", 1:3)
run_lavaan(measurement(tidy_sem(df), meanstructure = TRUE))
```

run\_mx

*Run as OpenMx model with sensible defaults***Description**

This convenience function runs objects for which a method exists using OpenMx, with sensible defaults. It is intended for use with tidySEM. For instance, it will convert a tidySEM object to a mxModel and run it, and it will try to ensure convergence for mixture models created using [mx\\_mixture](#). Knowledgeable users may want to run models manually.

**Usage**

```
run_mx(x, ...)
```

**Arguments**

`x` An object for which a method exists.  
`...` Parameters passed on to other functions.

**Value**

Returns an [mxModel](#) with free parameters updated to their final values.

**Examples**

```
df <- iris[1:3]
names(df) <- paste0("X_", 1:3)
run_mx(measurement(tidy_sem(df), meanstructure = TRUE))
```

skew\_kurtosis

*Calculate skew and kurtosis***Description**

Calculate skew and kurtosis, standard errors for both, and the estimates divided by two times the standard error. If this latter quantity exceeds an absolute value of 1, the skew/kurtosis is significant. With very large sample sizes, significant skew/kurtosis is common.

**Usage**

```
skew_kurtosis(x, verbose = FALSE, se = FALSE, ...)
```

**Arguments**

<code>x</code>	An object for which a method exists.
<code>verbose</code>	Logical. Whether or not to print messages to the console, Default: FALSE
<code>se</code>	Whether or not to return the standard errors, Default: FALSE
<code>...</code>	Additional arguments to pass to and from functions.

**Value**

A matrix of skew and kurtosis statistics for `x`.

**Examples**

```
skew_kurtosis(datasets::anscombe)
```

---

syntax	<i>Extract syntax from tidy_sem</i>
--------	-------------------------------------

---

**Description**

Provides access to the `syntax` element of a `tidy_sem` object. This can be used to return or assign to the `syntax` element.

**Usage**

```
syntax(x)

syntax(x) <- value
```

**Arguments**

<code>x</code>	Object of class <code>tidy_sem</code> .
<code>value</code>	A valid value for <code>syntax(x)</code> .

**Value**

data.frame

**Examples**

```
dict <- tidy_sem(iris, split = "\\.")
dict <- add_paths(dict, Sepal.Width ~ Sepal.Length)
syntax(dict)
```

---

table_cors	<i>Extract correlation tables</i>
------------	-----------------------------------

---

### Description

Extracts a publication-ready covariance or correlation matrix from an object for which a method exists.

### Usage

```
table_cors(x, value_column = "est_sig_std", digits = 2, ...)
```

### Arguments

x	An object for which a method exists.
value_column	Character. Name of the column to use to propagate the matrix. Defaults to "est_sig_std", the standardized estimate with significance asterisks.
digits	Number of digits to round to when formatting values.
...	Additional arguments passed to and from methods.

### Value

A Matrix or a list of matrices (in case there are between/within correlation matrices).

### Author(s)

Caspar J. van Lissa

### Examples

```
library(lavaan)
HS.model <- ' visual =~ x1 + x2 + x3
             textual =~ x4 + x5 + x6
             speed  =~ x7 + x8 + x9 '
fit <- cfa(HS.model,
           data = HolzingerSwineford1939,
           group = "school")
table_cors(fit)
```

---

table_fit	<i>Print model fit table formatted for publication</i>
-----------	--

---

### Description

Takes a model object, extracts model fit information, and formats it as a publication-ready table.

### Usage

```
table_fit(x, ...)
```

### Arguments

`x`                    A model object for which a method exists.  
`...`                 Arguments passed to other functions.

### Value

A data.frame of formatted results.

### Author(s)

Caspar J. van Lissa

### See Also

Other Reporting tools: [conf\\_int\(\)](#), [est\\_sig\(\)](#), [table\\_results\(\)](#)

### Examples

```
library(lavaan)
HS.model <- ' visual =~ x1 + x2 + x3
            textual =~ x4 + x5 + x6
            speed  =~ x7 + x8 + x9 '
fit <- cfa(HS.model,
          data = HolzingerSwineford1939,
          group = "school")
table_fit(fit)
```

---

table_results	<i>Print results table formatted for publication</i>
---------------	--

---

### Description

Takes a model object, and formats it as a publication-ready table.

### Usage

```
table_results(
  x,
  columns = c("label", "est_sig", "se", "pval", "confint", "group", "level"),
  digits = 2,
  ...
)
```

### Arguments

x	A model object for which a method exists.
columns	A character vector of columns to retain from the results section. If this is set to NULL, all available columns are returned. Defaults to c("label", "est_sig", "se", "pval", "confint", "group", "level"). These correspond to 1) the parameter label, 2) estimate column with significance asterisks appended (* <.05, ** <.01, *** <.001); 3) standard error, 4) p-value, 5) a formatted confidence interval, 6) grouping variable (if available), 7) level variable for multilevel models, if available.
digits	Number of digits to round to when formatting numeric columns.
...	Logical expressions used to filter the rows of results returned.

### Value

A data.frame of formatted results.

### Author(s)

Caspar J. van Lissa

### See Also

Other Reporting tools: [conf\\_int\(\)](#), [est\\_sig\(\)](#), [table\\_fit\(\)](#)

### Examples

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

```

      data = HolzingerSwineford1939,
      group = "school")
table_results(fit)

```

---

tidy\_sem

*Create a tidy\_sem object*


---

## Description

Create an object of class `tidy_sem`, which has the following elements:

- `dictionary` An overview of the variables in the `tidy_sem` object, and their assignment to scale/latent variables.
- `data` Optionally, the `data.frame` containing the data referenced in `$dictionary`.
- `syntax` Optionally, syntax defining a SEM-model by reference to the variables contained in `$data`.

## Usage

```
tidy_sem(x, split = "_")
```

## Arguments

<code>x</code>	An object for which a method exists, e.g., a vector of variable names, or a <code>data.frame</code> .
<code>split</code>	Character. Defining the regular expression used by <code>strsplit</code> to separate variable names into 1) the name of the scale/construct and 2) the number (or name) of the item.

## Details

When `tidy_sem` is called on a character string or `data.frame`, it attempts to assign variables to superordinate scale/latent variables based on the variable name and the splitting character defined in the `split` argument. Thus, the function will assign the variable `"scale_01"` to a scale/latent variable called `"scale"` when `split = "_"`. Alternatively, if the variable name is `"construct.1"`, the split character `"\."` separates the `"construct"` name from item number `"1"`. The character `"."` is escaped with a double backslash, because it is a special character in regular expressions.

## Value

An object of class `"tidy_sem"`

## Author(s)

Caspar J. van Lissa



**Examples**

```
tidy_sem(c("bfi_1", "bfi_2", "bfi_3", "bfi_4", "bfi_5",
"macqj_1", "macqj_2", "macqj_3", "macqj_4", "macqj_5", "macqj_6",
"macqj_7", "macqj_8", "macqj_9", "macqj_10", "macqj_11",
"macqj_12", "macqj_13", "macqj_14", "macqj_15", "macqj_16",
"macqj_17", "macqj_18", "macqj_19", "macqj_20", "macqj_21",
"macqr_1", "macqr_2", "macqr_3", "macqr_4", "macqr_5", "macqr_6",
"macqr_7", "macqr_8", "macqr_9", "macqr_10", "macqr_11",
"macqr_12", "macqr_13", "macqr_14", "macqr_15", "macqr_16",
"macqr_17", "macqr_18", "macqr_19", "macqr_20", "macqr_21", "sex"))
tidy_sem(c("bfi_1", "bfi_2", "bfi_3", "bfi_4", "bfi_5",
"mac_q_j_1", "mac_q_j_2", "mac_q_j_3", "mac_q_j_4", "mac_q_j_5", "mac_q_j_6",
"mac_q_j_7", "mac_q_j_8", "mac_q_j_9", "mac_q_j_10", "mac_q_j_11",
"mac_q_j_12", "mac_q_j_13", "mac_q_j_14", "mac_q_j_15", "mac_q_j_16",
"mac_q_j_17", "mac_q_j_18", "mac_q_j_19", "mac_q_j_20", "mac_q_j_21",
"mac_q_r_1", "mac_q_r_2", "mac_q_r_3", "mac_q_r_4", "mac_q_r_5", "mac_q_r_6",
"mac_q_r_7", "mac_q_r_8", "mac_q_r_9", "mac_q_r_10", "mac_q_r_11",
"mac_q_r_12", "mac_q_r_13", "mac_q_r_14", "mac_q_r_15", "mac_q_r_16",
"mac_q_r_17", "mac_q_r_18", "mac_q_r_19", "mac_q_r_20", "mac_q_r_21"))
```

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