

Package ‘broom.helpers’

November 24, 2020

Title Helpers for Model Coefficients Tibbles

Version 1.1.0

Description Provides suite of functions to work with regression model 'broom::tidy()' tibbles. The suite includes functions to group regression model terms by variable, insert reference and header rows for categorical variables, add variable labels, and more.

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URL <https://larmarange.github.io/broom.helpers/>

BugReports <https://github.com/larmarange/broom.helpers/issues>

Imports broom, dplyr, emmeans, labelled, lifecycle, purrr, rlang, stats, stringr, tibble, tidyr, usethis

Suggests broom.mixed, covr, datasets, forcats, gam, gee, geepack, ggplot2, gt, gtsummary, knitr, lavaan, lme4, MASS, mice, nnet, ordinal, rmarkdown, survey, survival, testthat, spelling

VignetteBuilder knitr

RdMacros lifecycle

Encoding UTF-8

LazyData true

RoxygenNote 7.1.1

Language en-US

NeedsCompilation no

Author Joseph Larmarange [aut, cre] (<<https://orcid.org/0000-0001-7097-700X>>), Daniel D. Sjoberg [aut] (<<https://orcid.org/0000-0003-0862-2018>>)

Maintainer Joseph Larmarange <joseph@larmarange.net>

Repository CRAN

Date/Publication 2020-11-24 08:00:02 UTC

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.clean_backticks	<i>Remove backticks around variable names</i>
------------------	---

Description

Remove backticks around variable names

Usage

```
.clean_backticks(x, variable_names = x)
```

Arguments

x a character vector to be cleaned
variable_names list of variable names, could be obtained with [model_list_variables\(only_variable = TRUE\)](#) to properly take into account interaction only terms/variables

See Also

Other other_helpers: [.escape_regex\(\)](#)

.escape_regex	<i>Escapes any characters that would have special meaning in a regular expression</i>
---------------	---

Description

This functions has been adapted from `Hmisc::escapeRegex()`

Usage

```
.escape_regex(string)
```

Arguments

string a character vector

See Also

Other other_helpers: [.clean_backticks\(\)](#)

.formula_list_to_named_list	<i>Convert formula selector to a named list</i>
-----------------------------	---

Description

Functions takes a list of formulas, e.g. `list(starts_with("age") ~ "continuous")`, and returns a named list, e.g. `list(age = "continuous")`.

Usage

```
.formula_list_to_named_list(  
  x,  
  data = NULL,  
  var_info = NULL,  
  arg_name = NULL,  
  select_single = FALSE  
)
```

Arguments

<code>x</code>	list of selecting formulas
<code>data</code>	A data frame to select columns from. Default is NULL
<code>var_info</code>	A data frame of variable names and attributes. May also pass a character vector of variable names. Default is NULL
<code>arg_name</code>	Optional string indicating the source argument name. This helps in the error messaging. Default is NULL.
<code>select_single</code>	Logical indicating whether the result must be a single variable. Default is FALSE

<code>.generic_selector</code>	<i>Generate a custom selector function</i>
--------------------------------	--

Description

Generate a custom selector function

Usage

```
.generic_selector(variable_column, select_column, select_expr, fun_name)
```

Arguments

<code>variable_column</code>	string indicating column variable names are stored
<code>select_column</code>	character vector of columns used in the <code>select_expr=</code> argument
<code>select_expr</code>	unquoted predicate command to subset a data frame to select variables
<code>fun_name</code>	quoted name of function where <code>.generic_selector()</code> is being used. This helps with error messaging.

Value

custom selector functions

.select_to_varnames *Variable selector*

Description

Function takes `select()`-like inputs and converts the selector to a character vector of variable names. Functions accepts `tidyselect` syntax, and additional selector functions defined within the package

Usage

```
.select_to_varnames(  
  select,  
  data = NULL,  
  var_info = NULL,  
  arg_name = NULL,  
  select_single = FALSE  
)
```

Arguments

<code>select</code>	A single object selecting variables, e.g. <code>c(age, stage)</code> , <code>starts_with("age")</code>
<code>data</code>	A data frame to select columns from. Default is <code>NULL</code>
<code>var_info</code>	A data frame of variable names and attributes. May also pass a character vector of variable names. Default is <code>NULL</code>
<code>arg_name</code>	Optional string indicating the source argument name. This helps in the error messaging. Default is <code>NULL</code> .
<code>select_single</code>	Logical indicating whether the result must be a single variable. Default is <code>FALSE</code>

Value

A character vector of variable names

`model_get_coefficients_type`
Get coefficient type

Description

Indicate the type of coefficient among "generic", "logistic", "poisson" and "prop_hazard".

Usage

```
model_get_coefficients_type(model)

## Default S3 method:
model_get_coefficients_type(model)

## S3 method for class 'glm'
model_get_coefficients_type(model)

## S3 method for class 'negbin'
model_get_coefficients_type(model)

## S3 method for class 'geeglm'
model_get_coefficients_type(model)

## S3 method for class 'glmerMod'
model_get_coefficients_type(model)

## S3 method for class 'clogit'
model_get_coefficients_type(model)

## S3 method for class 'polr'
model_get_coefficients_type(model)

## S3 method for class 'multinom'
model_get_coefficients_type(model)

## S3 method for class 'svyolr'
model_get_coefficients_type(model)

## S3 method for class 'clm'
model_get_coefficients_type(model)

## S3 method for class 'clmm'
model_get_coefficients_type(model)

## S3 method for class 'coxph'
model_get_coefficients_type(model)
```

Arguments

model a model object

See Also

Other `model_helpers`: [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```
lm(hp ~ mpg + factor(cyl), mtcars) %>%
  model_get_coefficients_type()

Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes"))) %>%
  glm(Survived ~ Class + Age * Sex, data = ., weights = .$n, family = binomial) %>%
  model_get_coefficients_type()
```

model_get_contrasts *Get contrasts used in the model*

Description

Get contrasts used in the model

Usage

```
model_get_contrasts(model)

## Default S3 method:
model_get_contrasts(model)
```

Arguments

model a model object

See Also

Other model_helpers: [model_get_coefficients_type\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```
glm(
  am ~ mpg + factor(cyl),
  data = mtcars,
  family = binomial,
  contrasts = list(`factor(cyl)` = contr.sum)
) %>%
  model_get_contrasts()
```

model_get_model	<i>Get the model from model objects</i>
-----------------	---

Description

Most model objects are proper R model objects. There are, however, some model objects that store the proper object internally (e.g. mice models). This function extracts that model object in those cases.

Usage

```
model_get_model(model)

## Default S3 method:
model_get_model(model)

## S3 method for class 'mira'
model_get_model(model)
```

Arguments

model a model object

See Also

Other model_helpers: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```
lm(hp ~ mpg + factor(cyl), mtcars) %>%
  model_get_model()
```

model_get_model_frame	<i>Get the model frame of a model</i>
-----------------------	---------------------------------------

Description

The structure of the object returned by `stats::model.frame()` could slightly differ for certain types of models. `model_get_model_frame()` will always return an object with the same data structure or NULL if it is not possible to compute model frame from model.

Usage

```

model_get_model_frame(model)

## Default S3 method:
model_get_model_frame(model)

## S3 method for class 'coxph'
model_get_model_frame(model)

## S3 method for class 'survreg'
model_get_model_frame(model)

```

Arguments

model a model object

See Also

[stats::model.frame\(\)](#)

Other `model_helpers`: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```

lm(hp ~ mpg + factor(cyl), mtcars) %>%
  model_get_model_frame() %>%
  head()

```

model_get_model_matrix

Get the model matrix of a model

Description

The structure of the object returned by [stats::model.matrix\(\)](#) could slightly differ for certain types of models. `model_get_model_matrix()` will always return an object with the same structure as [stats::model.matrix.default\(\)](#).

Usage

```

model_get_model_matrix(model)

## Default S3 method:
model_get_model_matrix(model)

## S3 method for class 'multinom'

```

```

model_get_model_matrix(model)

## S3 method for class 'clm'
model_get_model_matrix(model)

```

Arguments

model a model object

See Also

[stats::model.matrix\(\)](#)

Other model_helpers: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```

lm(hp ~ mpg + factor(cyl), mtcars) %>%
  model_get_model_matrix() %>%
  head()

```

model_get_nlevels	<i>Get the number of levels for each factor used in xlevels</i>
-------------------	---

Description

Get the number of levels for each factor used in xlevels

Usage

```

model_get_nlevels(model)

## Default S3 method:
model_get_nlevels(model)

```

Arguments

model a model object

Value

a tibble with two columns: "variable" and "var_nlevels"

See Also

Other model_helpers: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```
lm(hp ~ mpg + factor(cyl), mtcars) %>%  
  model_get_nlevels()
```

model_get_xlevels	<i>Get xlevels used in the model</i>
-------------------	--------------------------------------

Description

Get xlevels used in the model

Usage

```
model_get_xlevels(model)  
  
## Default S3 method:  
model_get_xlevels(model)  
  
## S3 method for class 'lmerMod'  
model_get_xlevels(model)  
  
## S3 method for class 'glmerMod'  
model_get_xlevels(model)
```

Arguments

model a model object

See Also

Other model_helpers: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```
lm(hp ~ mpg + factor(cyl), mtcars) %>%  
  model_get_xlevels()
```

model_identify_variables

Identify for each coefficient of a model the corresponding variable

Description

It will also identify interaction terms and intercept(s).

Usage

```
model_identify_variables(model)

## Default S3 method:
model_identify_variables(model)

## S3 method for class 'lavaan'
model_identify_variables(model)

## S3 method for class 'aov'
model_identify_variables(model)

## S3 method for class 'clm'
model_identify_variables(model)

## S3 method for class 'clmm'
model_identify_variables(model)
```

Arguments

model a model object

Value

A tibble with four columns:

- term: coefficients of the model
- variable: the corresponding variable
- var_class: class of the variable (cf. `stats::MFclass()`)
- var_type: "continuous", "dichotomous" (categorical variable with 2 levels), "categorical" (categorical variable with 3 or more levels), "intercept" or "interaction"
- var_nlevels: number of original levels for categorical variables

See Also

[tidy_identify_variables\(\)](#)

Other model_helpers: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```
Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes"))) %>%
  glm(
    Survived ~ Class + Age * Sex,
    data = ., weights = .$n,
    family = binomial
  ) %>%
  model_identify_variables()

iris %>%
  lm(
    Sepal.Length ~ poly(Sepal.Width, 2) + Species,
    data = .,
    contrasts = list(Species = contr.sum)
  ) %>%
  model_identify_variables()
```

model_list_contrasts *List contrasts used by a model*

Description

List contrasts used by a model

Usage

```
model_list_contrasts(model)
```

```
## Default S3 method:
model_list_contrasts(model)
```

Arguments

model a model object

Value

A tibble with three columns:

- variable: variable name
- contrasts: contrasts used
- contrasts_type: type of contrasts ("treatment", "sum", "poly", "helmert" or "other")
- reference: for variables with treatment, SAS or sum contrasts, position of the reference level

See Also

Other `model_helpers`: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_terms_levels\(\)](#), [model_list_variables\(\)](#)

Examples

```
glm(
  am ~ mpg + factor(cyl),
  data = mtcars,
  family = binomial,
  contrasts = list(`factor(cyl)` = contr.sum)
) %>%
  model_list_contrasts()
```

`model_list_terms_levels`

List levels of categorical terms

Description

Only for categorical variables with treatment, SAS or sum contrasts.

Usage

```
model_list_terms_levels(
  model,
  label_pattern = "{level}",
  variable_labels = NULL
)
```

```
## Default S3 method:
model_list_terms_levels(
  model,
  label_pattern = "{level}",
  variable_labels = NULL
)
```

Arguments

`model` a model object

`label_pattern` a [glue pattern](#) for term labels (see examples)

`variable_labels` an optional named list or named vector of custom variable labels passed to [model_list_variables\(\)](#)

Value

A tibble with ten columns:

- variable: variable
- contrasts_type: type of contrasts ("sum" or "treatment")
- term: term name
- level: term level
- reference: logical indicating which term is the reference level
- reference_level: level of the reference term
- var_label: variable label obtained with [model_list_variables\(\)](#)
- var_nlevels: number of levels in this variable
- dichotomous: logical indicating if the variable is dichotomous
- label: term label (by default equal to term level) The first nine columns can be used in label_pattern.

See Also

Other model_helpers: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_variables\(\)](#)

Examples

```

glm(
  am ~ mpg + factor(cyl),
  data = mtcars,
  family = binomial,
  contrasts = list(`factor(cyl)` = contr.sum)
) %>%
  model_list_terms_levels()

df <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes")))

mod <- df %>%
  glm(
    Survived ~ Class + Age + Sex,
    data = ., weights = .$n, family = binomial,
    contrasts = list(Age = contr.sum, Class = "contr.helmert")
  )
mod %>% model_list_terms_levels()
mod %>% model_list_terms_levels("{level} vs {reference_level}")
mod %>% model_list_terms_levels("{variable} [{level} - {reference_level}]")
mod %>% model_list_terms_levels(
  "{ifelse(reference, level, paste(level, '-', reference_level))}"
)

```

model_list_variables *List all the variables used in a model*

Description

Including variables used only in an interaction.

Usage

```
model_list_variables(model, labels = NULL, only_variable = FALSE)

## Default S3 method:
model_list_variables(model, labels = NULL, only_variable = FALSE)

## S3 method for class 'lavaan'
model_list_variables(model, labels = NULL, only_variable = FALSE)
```

Arguments

model a model object
 labels an optional named list or named vector of custom variable labels
 only_variable if TRUE, will return only "variable" column

Value

A tibble with three columns:

- variable: the corresponding variable
- var_class: class of the variable (cf. [stats::MFclass\(\)](#))
- label_attr: variable label defined in the original data frame with the label attribute (cf. [labelled::var_label\(\)](#))
- var_label: a variable label (by priority, labels if defined, label_attr if available, otherwise variable)

See Also

Other model_helpers: [model_get_coefficients_type\(\)](#), [model_get_contrasts\(\)](#), [model_get_model_frame\(\)](#), [model_get_model_matrix\(\)](#), [model_get_model\(\)](#), [model_get_nlevels\(\)](#), [model_get_xlevels\(\)](#), [model_identify_variables\(\)](#), [model_list_contrasts\(\)](#), [model_list_terms_levels\(\)](#)

Examples

```
Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes"))) %>%
  glm(
    Survived ~ Class + Age : Sex,
```



```

      data = ., weights = .$n,
      family = binomial
    ) %>%
    model_list_variables()

iris %>%
  lm(
    Sepal.Length ~ poly(Sepal.Width, 2) + Species,
    data = .,
    contrasts = list(Species = contr.sum)
  ) %>%
  model_list_variables()

if (requireNamespace("gtsummary")) {
  glm(
    response ~ poly(age, 3) + stage + grade * trt,
    na.omit(gtsummary::trial),
    family = binomial,
  ) %>%
  model_list_variables()
}

```

select_helpers

Select helper functions

Description

Set of functions to supplement the tidyselect set of functions for selecting columns of data frames (and other items as well).

- `all_continuous()` selects continuous variables
- `all_categorical()` selects categorical (including "dichotomous") variables
- `all_dichotomous()` selects only type "dichotomous"
- `all_interaction()` selects interaction terms from a regression model
- `all_intercepts()` selects intercept terms from a regression model
- `all_contrasts()` selects variables in regression model based on their type of contrast

Usage

```
all_continuous()
```

```
all_dichotomous()
```

```
all_categorical(dichotomous = TRUE)
```

```
all_interaction()
```

```
all_intercepts()
```

```
all_contrasts(contrasts_type = NULL)
```

Arguments

`dichotomous` Logical indicating whether to include dichotomous variables. Default is TRUE

`contrasts_type` type of contrast to select. When NULL, all variables with a contrast will be selected. Default is NULL. Select among contrast types `c("treatment", "sum", "poly", "helmert", "other")`

Value

A character vector of column names selected

Examples

```
glm(response ~ age * trt + grade, gtsummary::trial, family = binomial) %>%
  tidy_plus_plus(exponentiate = TRUE, include = all_categorical())

glm(response ~ age + trt + grade + stage,
  gtsummary::trial,
  family = binomial,
  contrasts = list(trt = contr.SAS, grade = contr.sum, stage = contr.poly)) %>%
  tidy_plus_plus(exponentiate = TRUE,
    include = all_contrasts(c("treatment", "sum")))
```

`tidy_add_coefficients_type`

Add coefficients type and label as attributes

Description

Add the type of coefficients ("generic", "logistic", "poisson" or "prop_hazard") and the corresponding coefficient labels, as attributes to `x` (respectively named `coefficients_type` and `coefficients_label`).

Usage

```
tidy_add_coefficients_type(
  x,
  exponentiate = attr(x, "exponentiate"),
  model = tidy_get_model(x)
)
```

Arguments

`x` a tidy tibble

`exponentiate` logical indicating whether or not to exponentiate the coefficient estimates. It should be consistent with the original call to `broom::tidy()`

`model` the corresponding model, if not attached to `x`

See Also

Other tidy_helpers: [tidy_add_contrasts\(\)](#), [tidy_add_estimate_to_reference_rows\(\)](#), [tidy_add_header_rows\(\)](#), [tidy_add_reference_rows\(\)](#), [tidy_add_term_labels\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_attach_model\(\)](#), [tidy_identify_variables\(\)](#), [tidy_plus_plus\(\)](#), [tidy_remove_intercept\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
ex1 <- lm(hp ~ mpg + factor(cyl), mtcars) %>%
  tidy_and_attach() %>%
  tidy_add_coefficients_type()
attr(ex1, "coefficients_type")
attr(ex1, "coefficients_label")

ex2 <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes"))) %>%
  glm(Survived ~ Class + Age * Sex, data = ., weights = .$n, family = binomial) %>%
  tidy_and_attach(exponentiate = TRUE) %>%
  tidy_add_coefficients_type()
attr(ex2, "coefficients_type")
attr(ex2, "coefficients_label")
```

`tidy_add_contrasts` *Add contrasts type for categorical variables*

Description

Add a contrasts column corresponding to contrasts used for a categorical variable and a contrasts_type column equal to "treatment", "sum", "poly", "helmert" or "other".

Usage

```
tidy_add_contrasts(x, model = tidy_get_model(x))
```

Arguments

<code>x</code>	a tidy tibble
<code>model</code>	the corresponding model, if not attached to <code>x</code>

Details

If the variable column is not yet available in `x`, [tidy_identify_variables\(\)](#) will be automatically applied.

See Also

Other tidy_helpers: [tidy_add_coefficients_type\(\)](#), [tidy_add_estimate_to_reference_rows\(\)](#), [tidy_add_header_rows\(\)](#), [tidy_add_reference_rows\(\)](#), [tidy_add_term_labels\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_attach_model\(\)](#), [tidy_identify_variables\(\)](#), [tidy_plus_plus\(\)](#), [tidy_remove_intercept\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
df <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes")))

df %>%
  glm(
    Survived ~ Class + Age + Sex,
    data = ., weights = .$n, family = binomial,
    contrasts = list(Age = contr.sum, Class = "contr.helmert")
  ) %>%
  tidy_and_attach() %>%
  tidy_add_contrasts()
```

tidy_add_estimate_to_reference_rows

Add an estimate value to references rows for categorical variables

Description

For categorical variables with a treatment contrast (`stats::contr.treatment()`) or a SAS contrast (`stats::contr.SAS()`) will add an estimate equal to 0 (or 1 if `exponentiate = TRUE`) to the reference row.

Usage

```
tidy_add_estimate_to_reference_rows(
  x,
  exponentiate = attr(x, "exponentiate"),
  model = tidy_get_model(x),
  quiet = FALSE
)
```

Arguments

<code>x</code>	a tidy tibble
<code>exponentiate</code>	logical indicating whether or not to exponentiate the coefficient estimates. It should be consistent with the original call to <code>broom::tidy()</code>
<code>model</code>	the corresponding model, if not attached to <code>x</code>
<code>quiet</code>	logical argument whether <code>broom.helpers</code> should not return a message when requested output cannot be generated. Default is <code>FALSE</code>

Details

For categorical variables with a sum contrast (`stats::contr.sum()`), the estimate value of the reference row will be equal to the sum of all other coefficients multiplied by -1 (eventually exponentiated if `exponentiate = TRUE`), and obtained with `stats::dummy.coef()`. For sum contrasts, the model coefficient corresponds to the difference of each level with the grand mean.

For other variables, no change will be made.

If the `reference_row` column is not yet available in `x`, `tidy_add_reference_rows()` will be automatically applied.

See Also

Other tidy_helpers: `tidy_add_coefficients_type()`, `tidy_add_contrasts()`, `tidy_add_header_rows()`, `tidy_add_reference_rows()`, `tidy_add_term_labels()`, `tidy_add_variable_labels()`, `tidy_attach_model()`, `tidy_identify_variables()`, `tidy_plus_plus()`, `tidy_remove_intercept()`, `tidy_select_variables()`

Examples

```
df <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(dplyr::across(where(is.character), factor))

df %>%
  glm(
    Survived ~ Class + Age + Sex,
    data = ., weights = .$n, family = binomial,
    contrasts = list(Age = contr.sum, Class = "contr.SAS")
  ) %>%
  tidy_and_attach(exponentiate = TRUE) %>%
  tidy_add_reference_rows() %>%
  tidy_add_estimate_to_reference_rows()

if (requireNamespace("gtsummary")) {
  glm(
    response ~ stage + grade * trt,
    gtsummary::trial,
    family = binomial,
    contrasts = list(
      stage = contr.treatment(4, base = 3),
      grade = contr.treatment(3, base = 2),
      trt = contr.treatment(2, base = 2)
    )
  ) %>%
  tidy_and_attach() %>%
  tidy_add_reference_rows() %>%
  tidy_add_estimate_to_reference_rows()
}
```

`tidy_add_header_rows` *Add header rows variables with several terms*

Description

For variables with several terms (usually categorical variables but could also be the case of continuous variables with polynomial terms or splines), `tidy_add_header_rows()` will add an additional row per variable, where `label` will be equal to `var_label`. These additional rows could be identified with `header_row` column.

Usage

```
tidy_add_header_rows(
  x,
  show_single_row = NULL,
  model = tidy_get_model(x),
  quiet = FALSE,
  strict = FALSE
)
```

Arguments

<code>x</code>	a tidy tibble
<code>show_single_row</code>	a vector indicating the names of binary variables that should be displayed on a single row. Accepts <code>tidyselect</code> syntax. Default is <code>NULL</code> . See also all_dichotomous()
<code>model</code>	the corresponding model, if not attached to <code>x</code>
<code>quiet</code>	logical argument whether broom.helpers should not return a message when requested output cannot be generated. Default is <code>FALSE</code>
<code>strict</code>	logical argument whether broom.helpers should return an error when requested output cannot be generated. Default is <code>FALSE</code>

Details

The `show_single_row` argument allows to specify a list of dichotomous variables that should be displayed on a single row instead of two rows.

The added `header_row` column will be equal to:

- `TRUE` for an header row;
- `FALSE` for a normal row of a variable with an header row;
- `NA` for variables without an header row.

If the `label` column is not yet available in `x`, [tidy_add_term_labels\(\)](#) will be automatically applied.

See Also

Other tidy_helpers: [tidy_add_coefficients_type\(\)](#), [tidy_add_contrasts\(\)](#), [tidy_add_estimate_to_reference_rows\(\)](#), [tidy_add_reference_rows\(\)](#), [tidy_add_term_labels\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_attach_model\(\)](#), [tidy_identify_variables\(\)](#), [tidy_plus_plus\(\)](#), [tidy_remove_intercept\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
df <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes")))

res <- df %>%
  glm(
    Survived ~ Class + Age + Sex,
    data = ., weights = .$n, family = binomial,
    contrasts = list(Age = contr.sum, Class = "contr.SAS")
  ) %>%
  tidy_and_attach() %>%
  tidy_add_variable_labels(labels = list(Class = "Custom label for Class")) %>%
  tidy_add_reference_rows()
res %>% tidy_add_header_rows()
res %>% tidy_add_header_rows(show_single_row = all_dichotomous())

if (requireNamespace("gtsummary")) {
  glm(
    response ~ stage + grade * trt,
    gtsummary::trial,
    family = binomial,
    contrasts = list(
      stage = contr.treatment(4, base = 3),
      grade = contr.treatment(3, base = 2),
      trt = contr.treatment(2, base = 2)
    )
  ) %>%
  tidy_and_attach() %>%
  tidy_add_reference_rows() %>%
  tidy_add_header_rows()
}
```

tidy_add_reference_rows

Add references rows for categorical variables

Description

For categorical variables with a treatment contrast ([stats::contr.treatment\(\)](#)), a SAS contrast ([stats::contr.SAS\(\)](#)) or a sum contrast ([stats::contr.sum\(\)](#)), add a reference row.

Usage

```
tidy_add_reference_rows(
  x,
  no_reference_row = NULL,
  model = tidy_get_model(x),
  quiet = FALSE
)
```

Arguments

x	a tidy tibble
no_reference_row	a vector indicating the name of variables for those no reference row should be added. Accepts tidyselect syntax. Default is NULL. See also all_categorical() and all_dichotomous()
model	the corresponding model, if not attached to x
quiet	logical argument whether broom.helpers should not return a message when requested output cannot be generated. Default is FALSE

Details

The added reference_row column will be equal to:

- TRUE for a reference row;
- FALSE for a normal row of a variable with a reference row;
- NA for variables without a reference row.

If the contrasts column is not yet available in x, [tidy_add_contrasts\(\)](#) will be automatically applied.

`tidy_add_reference_rows()` will not populate the label of the reference term. It is therefore better to apply [tidy_add_term_labels\(\)](#) after `tidy_add_reference_rows()` rather than before.

See Also

Other tidy_helpers: [tidy_add_coefficients_type\(\)](#), [tidy_add_contrasts\(\)](#), [tidy_add_estimate_to_reference_row\(\)](#), [tidy_add_header_rows\(\)](#), [tidy_add_term_labels\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_attach_model\(\)](#), [tidy_identify_variables\(\)](#), [tidy_plus_plus\(\)](#), [tidy_remove_intercept\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
df <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes")))

res <- df %>%
  glm(
    Survived ~ Class + Age + Sex,
    data = ., weights = .$n, family = binomial,
```



```

      contrasts = list(Age = contr.sum, Class = "contr.SAS")
    ) %>%
    tidy_and_attach()
  res %>% tidy_add_reference_rows()
  res %>% tidy_add_reference_rows(no_reference_row = all_dichotomous())
  res %>% tidy_add_reference_rows(no_reference_row = "Class")

  if (requireNamespace("gtsummary")) {
    glm(
      response ~ stage + grade * trt,
      gtsummary::trial,
      family = binomial,
      contrasts = list(
        stage = contr.treatment(4, base = 3),
        grade = contr.treatment(3, base = 2),
        trt = contr.treatment(2, base = 2)
      )
    ) %>%
    tidy_and_attach() %>%
    tidy_add_reference_rows()
  }

```

tidy_add_term_labels *Add term labels*

Description

Will add term labels in a label column, based on:

1. labels provided in labels argument if provided;
2. factor levels for categorical variables coded with treatment, SAS or sum contrasts (the label could be customized with categorical_terms_pattern argument);
3. variable labels when there is only one term per variable;
4. term name otherwise.

Usage

```

tidy_add_term_labels(
  x,
  labels = NULL,
  interaction_sep = " * ",
  categorical_terms_pattern = "{level}",
  model = tidy_get_model(x),
  quiet = FALSE,
  strict = FALSE
)

```

Arguments

x	a tidy tibble
labels	an optional named list or named vector of custom term labels
interaction_sep	separator for interaction terms
categorical_terms_pattern	a glue pattern for labels of categorical terms with treatment or sum contrasts (see examples and model_list_terms_levels())
model	the corresponding model, if not attached to x
quiet	logical argument whether broom.helpers should not return a message when requested output cannot be generated. Default is FALSE
strict	logical argument whether broom.helpers should return an error when requested output cannot be generated. Default is FALSE

Details

If the variable_label column is not yet available in x, [tidy_add_variable_labels\(\)](#) will be automatically applied. If the contrasts column is not yet available in x, [tidy_add_contrasts\(\)](#) will be automatically applied.

It is possible to pass a custom label for any term in labels, including interaction terms.

See Also

Other tidy_helpers: [tidy_add_coefficients_type\(\)](#), [tidy_add_contrasts\(\)](#), [tidy_add_estimate_to_reference_row\(\)](#), [tidy_add_header_rows\(\)](#), [tidy_add_reference_rows\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_attach_model\(\)](#), [tidy_identify_variables\(\)](#), [tidy_plus_plus\(\)](#), [tidy_remove_intercept\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
df <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes"))) %>%
  labelled::set_variable_labels(
    Class = "Passenger's class",
    Sex = "Sex"
  )

mod <- df %>%
  glm(Survived ~ Class * Age * Sex, data = ., weights = .$n, family = binomial)
mod %>%
  tidy_and_attach() %>%
  tidy_add_term_labels()

mod %>%
  tidy_and_attach() %>%
  tidy_add_term_labels(
    interaction_sep = " x ",
    categorical_terms_pattern = "{level} / {reference_level}"
  )
```

tidy_add_variable_labels
Add variable labels

Description

Will add variable labels in a `var_label` column, based on:

1. labels provided in `labels` argument if provided;
2. variable labels defined in the original data frame with the `label` attribute (cf. `labelled::var_label()`);
3. variable name otherwise.

Usage

```
tidy_add_variable_labels(  
  x,  
  labels = NULL,  
  interaction_sep = " * ",  
  model = tidy_get_model(x),  
  quiet = FALSE,  
  strict = FALSE  
)
```

Arguments

<code>x</code>	a tidy tibble
<code>labels</code>	an optional named list or named vector of custom variable labels
<code>interaction_sep</code>	separator for interaction terms
<code>model</code>	the corresponding model, if not attached to <code>x</code>
<code>quiet</code>	logical argument whether broom.helpers should not return a message when requested output cannot be generated. Default is FALSE
<code>strict</code>	logical argument whether broom.helpers should return an error when requested output cannot be generated. Default is FALSE

Details

If the variable column is not yet available in `x`, `tidy_identify_variables()` will be automatically applied.

It is possible to pass a custom label for an interaction term in `labels` (see examples).

See Also

Other tidy_helpers: `tidy_add_coefficients_type()`, `tidy_add_contrasts()`, `tidy_add_estimate_to_reference_rows()`, `tidy_add_header_rows()`, `tidy_add_reference_rows()`, `tidy_add_term_labels()`, `tidy_attach_model()`, `tidy_identify_variables()`, `tidy_plus_plus()`, `tidy_remove_intercept()`, `tidy_select_variables()`

Examples

```
df <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes"))) %>%
  labelled::set_variable_labels(
    Class = "Passenger's class",
    Sex = "Sex"
  )

df %>%
  glm(Survived ~ Class * Age * Sex, data = ., weights = .$n, family = binomial) %>%
  tidy_and_attach() %>%
  tidy_add_variable_labels(
    labels = list(
      "(Intercept)" = "Custom intercept",
      Sex = "Gender",
      "Class:Age" = "Custom label"
    )
  )
```

tidy_attach_model	<i>Attach a full model to the tibble of model terms</i>
-------------------	---

Description

To facilitate the use of broom helpers with pipe, it is recommended to attach the original model as an attribute to the tibble of model terms generated by `broom::tidy()`.

Usage

```
tidy_attach_model(x, model, .attributes = NULL)

tidy_and_attach(model, tidy_fun = broom::tidy, exponentiate = FALSE, ...)

tidy_get_model(x)

tidy_detach_model(x)
```

Arguments

<code>x</code>	a tibble of model terms
<code>model</code>	a model to be attached/tidied
<code>.attributes</code>	named list of additional attributes to be attached to <code>x</code>
<code>tidy_fun</code>	option to specify a custom tidier function
<code>exponentiate</code>	logical indicating whether or not to exponentiate the coefficient estimates. This is typical for logistic, Poisson and Cox models, but a bad idea if there is no log or logit link; defaults to <code>FALSE</code>
<code>...</code>	other arguments passed to <code>tidy_fun()</code>

Details

`tidy_attach_model()` attach the model to a tibble already generated while `tidy_and_attach()` will apply `broom::tidy()` and attach the model.

Use `tidy_get_model()` to get the model attached to the tibble and `tidy_detach_model()` to remove the attribute containing the model.

See Also

Other tidy_helpers: [tidy_add_coefficients_type\(\)](#), [tidy_add_contrasts\(\)](#), [tidy_add_estimate_to_reference_rows\(\)](#), [tidy_add_header_rows\(\)](#), [tidy_add_reference_rows\(\)](#), [tidy_add_term_labels\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_identify_variables\(\)](#), [tidy_plus_plus\(\)](#), [tidy_remove_intercept\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
mod <- lm(Sepal.Length ~ Sepal.Width + Species, data = iris)
tt <- mod %>%
  tidy_and_attach(conf.int = TRUE)
tt
tidy_get_model(tt)
```

`tidy_identify_variables`

Identify the variable corresponding to each model coefficient

Description

`tidy_identify_variables()` will add to the tidy tibble three additional columns: `variable`, `var_class`, `var_type` and `var_nlevels`.

Usage

```
tidy_identify_variables(
  x,
  model = tidy_get_model(x),
  quiet = FALSE,
  strict = FALSE
)
```

Arguments

<code>x</code>	a tidy tibble
<code>model</code>	the corresponding model, if not attached to <code>x</code>
<code>quiet</code>	logical argument whether broom.helpers should not return a message when requested output cannot be generated. Default is FALSE
<code>strict</code>	logical argument whether broom.helpers should return an error when requested output cannot be generated. Default is FALSE

Details

It will also identify interaction terms and intercept(s).

var_type could be:

- "continuous",
- "dichotomous" (categorical variable with 2 levels),
- "categorical" (categorical variable with 3 levels or more),
- "intercept"
- "interaction"
- "unknown" in the rare cases where tidy_identify_variables() will fail to identify the list of variables

For dichotomous and categorical variables, var_nlevels corresponds to the number of original levels in the corresponding variables.

See Also

[model_identify_variables\(\)](#)

Other tidy_helpers: [tidy_add_coefficients_type\(\)](#), [tidy_add_contrasts\(\)](#), [tidy_add_estimate_to_reference_row\(\)](#), [tidy_add_header_rows\(\)](#), [tidy_add_reference_rows\(\)](#), [tidy_add_term_labels\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_attach_model\(\)](#), [tidy_plus_plus\(\)](#), [tidy_remove_intercept\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived, c("No", "Yes"))) %>%
  glm(Survived ~ Class + Age * Sex, data = ., weights = .$n, family = binomial) %>%
  tidy_and_attach() %>%
  tidy_identify_variables()
```

```
lm(
  Sepal.Length ~ poly(Sepal.Width, 2) + Species,
  data = iris,
  contrasts = list(Species = contr.sum)
) %>%
  tidy_and_attach(conf.int = TRUE) %>%
  tidy_identify_variables()
```

Description

This function will apply sequentially:

- `tidy_and_attach()`
- `tidy_identify_variables()`
- `tidy_add_contrasts()`
- `tidy_add_reference_rows()`
- `tidy_add_estimate_to_reference_rows()`
- `tidy_add_variable_labels()`
- `tidy_add_term_labels()`
- `tidy_add_header_rows()`
- `tidy_remove_intercept()`
- `tidy_select_variables()`
- `tidy_add_coefficients_type()`
- `tidy_detach_model()`

Usage

```
tidy_plus_plus(  
  model,  
  tidy_fun = broom::tidy,  
  conf.int = TRUE,  
  exponentiate = FALSE,  
  variable_labels = NULL,  
  term_labels = NULL,  
  interaction_sep = " * ",  
  categorical_terms_pattern = "{level}",  
  add_reference_rows = TRUE,  
  no_reference_row = NULL,  
  add_estimate_to_reference_rows = TRUE,  
  add_header_rows = FALSE,  
  show_single_row = NULL,  
  intercept = FALSE,  
  include = everything(),  
  keep_model = FALSE,  
  quiet = FALSE,  
  strict = FALSE,  
  ...  
)
```

Arguments

<code>model</code>	a model to be attached/tidied
<code>tidy_fun</code>	option to specify a custom tidier function
<code>conf.int</code>	should confidence intervals be computed? (see <code>broom::tidy()</code>)

exponentiate	logical indicating whether or not to exponentiate the coefficient estimates. This is typical for logistic, Poisson and Cox models, but a bad idea if there is no log or logit link; defaults to FALSE.
variable_labels	a named list or a named vector of custom variable labels
term_labels	a named list or a named vector of custom term labels
interaction_sep	separator for interaction terms
categorical_terms_pattern	a glue pattern for labels of categorical terms with treatment or sum contrasts (see model_list_terms_levels())
add_reference_rows	should reference rows be added?
no_reference_row	variables (accepts tidyselect notation) for those no reference row should be added, when add_reference_rows = TRUE
add_estimate_to_reference_rows	should an estimate value be added to reference rows?
add_header_rows	should header rows be added?
show_single_row	variables that should be displayed on a single row (accepts tidyselect notation), when add_header_rows is TRUE
intercept	should the intercept(s) be included?
include	variables to include. Accepts tidyselect syntax. Use - to remove a variable. Default is everything(). See also all_continuous() , all_categorical() , all_dichotomous() and all_interaction()
keep_model	should the model be kept as an attribute of the final result?
quiet	logical argument whether broom.helpers should not return a message when requested output cannot be generated. Default is FALSE
strict	logical argument whether broom.helpers should return an error when requested output cannot be generated. Default is FALSE
...	other arguments passed to tidy_fun()

See Also

Other tidy_helpers: [tidy_add_coefficients_type\(\)](#), [tidy_add_contrasts\(\)](#), [tidy_add_estimate_to_reference_rows\(\)](#), [tidy_add_header_rows\(\)](#), [tidy_add_reference_rows\(\)](#), [tidy_add_term_labels\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_attach_model\(\)](#), [tidy_identify_variables\(\)](#), [tidy_remove_intercept\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
ex1 <- lm(Sepal.Length ~ Sepal.Width + Species, data = iris) %>%
  tidy_plus_plus()
ex1
```



```

df <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(
    Survived = factor(Survived, c("No", "Yes"))
  ) %>%
  labelled::set_variable_labels(
    Class = "Passenger's class",
    Sex = "Gender"
  )

ex2 <- glm(
  Survived ~ Class + Age * Sex,
  data = df, weights = df$n,
  family = binomial
) %>%
  tidy_plus_plus(
    exponentiate = TRUE,
    add_reference_rows = FALSE,
    categorical_terms_pattern = "{level} / {reference_level}"
  )
ex2

if (requireNamespace("gtsummary")) {
  ex3 <- glm(
    response ~ poly(age, 3) + stage + grade * trt,
    na.omit(gtsummary::trial),
    family = binomial,
    contrasts = list(
      stage = contr.treatment(4, base = 3),
      grade = contr.sum
    )
  ) %>%
  tidy_plus_plus(
    exponentiate = TRUE,
    variable_labels = c(age = "Age (in years)"),
    add_header_rows = TRUE,
    show_single_row = all_dichotomous(),
    term_labels = c("poly(age, 3)3" = "Cubic age"),
    keep_model = TRUE
  )
  ex3
}

```

tidy_remove_intercept *Remove intercept(s)*

Description

Will remove terms where `var_type == "intercept"`.

Usage

```
tidy_remove_intercept(x, model = tidy_get_model(x))
```

Arguments

x a tidy tibble
 model the corresponding model, if not attached to x

Details

If the variable column is not yet available in x, [tidy_identify_variables\(\)](#) will be automatically applied.

See Also

Other tidy_helpers: [tidy_add_coefficients_type\(\)](#), [tidy_add_contrasts\(\)](#), [tidy_add_estimate_to_reference_rows\(\)](#), [tidy_add_header_rows\(\)](#), [tidy_add_reference_rows\(\)](#), [tidy_add_term_labels\(\)](#), [tidy_add_variable_labels\(\)](#), [tidy_attach_model\(\)](#), [tidy_identify_variables\(\)](#), [tidy_plus_plus\(\)](#), [tidy_select_variables\(\)](#)

Examples

```
Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived)) %>%
  glm(Survived ~ Class + Age + Sex, data = ., weights = .$n, family = binomial) %>%
  tidy_and_attach() %>%
  tidy_remove_intercept()
```

tidy_select_variables *Select variables to keep/drop*

Description

Will remove unselected variables from the results. To remove the intercept, use [tidy_remove_intercept\(\)](#).

Usage

```
tidy_select_variables(x, include = everything(), model = tidy_get_model(x))
```

Arguments

x a tidy tibble
 include variables to include. Accepts [tidyselect](#) syntax. Use - to remove a variable. Default is [everything\(\)](#). See also [all_continuous\(\)](#), [all_categorical\(\)](#), [all_dichotomous\(\)](#) and [all_interaction\(\)](#)
 model the corresponding model, if not attached to x

Details

If the variable column is not yet available in `x`, `tidy_identify_variables()` will be automatically applied.

See Also

Other tidy_helpers: `tidy_add_coefficients_type()`, `tidy_add_contrasts()`, `tidy_add_estimate_to_reference_rows()`, `tidy_add_header_rows()`, `tidy_add_reference_rows()`, `tidy_add_term_labels()`, `tidy_add_variable_labels()`, `tidy_attach_model()`, `tidy_identify_variables()`, `tidy_plus_plus()`, `tidy_remove_intercept()`

Examples

```
res <- Titanic %>%
  dplyr::as_tibble() %>%
  dplyr::mutate(Survived = factor(Survived)) %>%
  glm(Survived ~ Class + Age * Sex, data = ., weights = .$n, family = binomial) %>%
  tidy_and_attach() %>%
  tidy_identify_variables()

res
res %>% tidy_select_variables()
res %>% tidy_select_variables(include = "Class")
res %>% tidy_select_variables(include = -c("Age", "Sex"))
res %>% tidy_select_variables(include = starts_with("A"))
res %>% tidy_select_variables(include = all_categorical())
res %>% tidy_select_variables(include = all_dichotomous())
res %>% tidy_select_variables(include = all_interaction())
res %>% tidy_select_variables(
  include = c("Age", all_categorical(dichotomous = FALSE), all_interaction())
)
```

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