

Package ‘partialling.out’

October 17, 2025

Title Residuals from Partial Regressions

Version 0.2.0

Description Creates a data frame with the residuals of partial regressions of the main explanatory variable and the variable of interest. This method follows the Frisch-Waugh-Lovell theorem, as explained in Lovell (2008) <[doi:10.3200/JECE.39.1.88-91](https://doi.org/10.3200/JECE.39.1.88-91)>.

License GPL (>= 3)

BugReports <https://github.com/ropensci/partialling.out/issues/>

Suggests tinytest, tinsnapshot, knitr, rmarkdown, palmerpenguins, tinytable, fwlplot, tsibble, units, purrr, fontquiver, rsvg, svglite

Config/testthat.edition 3

Encoding UTF-8

RoxygenNote 7.3.2

VignetteBuilder knitr

Imports glue, lifecycle, rlang, fixest, lfe, stats, tinyplot

URL <https://docs.ropensci.org/partialling.out/>,
<https://github.com/ropensci/partialling.out/>

NeedsCompilation no

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Repository CRAN

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<i>partialalling_out</i>	<i>partialalling_out</i> : partialling out variable of interest and main
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Description

Creates a data.frame of the residualised main explanatory variable and, if wanted, variable of interest of a linear or fixed effects model

Usage

```
partialalling_out(model, data, weights, both, na.rm, ...)
```

Arguments

<code>model</code>	object for which we want to residualise variables
<code>data</code>	data.frame used in the original model. Using different data will return unexpected results or an error.
<code>weights</code>	a numeric vector for weighting the partial models. Length must be equal to number of rows of data
<code>both</code>	if TRUE will residualise both the variable of interest and the first explanatory variable in the model. If FALSE, only the latter. Set to TRUE by default
<code>na.rm</code>	if TRUE will remove observations with NA before any models are run. If FALSE, the underlying <code>lm</code> , <code>feols</code> , or <code>felm</code> will remove NA values but errors may arise if weights are used.
<code>...</code>	Any other <code>lm</code> , <code>feols</code> , or <code>felm</code> parameters that will be passed to the partial regressions

Details

The function regresses the main (i.e. first in the model) explanatory variable and the variable of interest (if parameter `both` is set to TRUE) against all other control variables and fixed effects and returns the residuals in a data.frame

Will accept `lm`, `felm` (`lfe` package), and `feols` (`fixest` package) objects

Value

a data.frame with the (residualised) variable of interest and residualised main explanatory variable

Examples

```
library(palmerpenguins)
library(fixest)
model <- feols(bill_length_mm ~ bill_depth_mm | species + island,
               data = penguins)
partial_df <- partialling_out(model, penguins, both = TRUE)
```

plot_partial_residuals

plot_partial_residuals: scatterplot of partial residuals

Description

Function for plotting partial residuals Uses `tinyplot` as backend

Usage

```
plot_partial_residuals(x, add_lm = TRUE, quantile = FALSE, probs = 0.02, ...)
```

Arguments

x	a <code>partial_residuals</code> objects from <code>partialling_out()</code>
add_lm	if TRUE, a <code>lm</code> will be plotted
quantile	if TRUE, will plot only the mean values of the quantiles of the mean explanatory variable specified by <code>probs</code>
probs	numeric vector of length one that specifies the number of quantiles to be computed if <code>quantile</code> is TRUE. by default, 0.02, which will give 50 quantiles.
...	Any other <code>tinyplot::plt()</code> params

Value

invisibly, x

Examples

```
library(palmerpenguins)
library(fixest)
model <- feols(bill_length_mm ~ bill_depth_mm | species + island,
               data = penguins)
partial_df <- partialling_out(model, penguins, both = TRUE)
plot_partial_residuals(partial_df)
```

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