Package ‘panelr’

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Title Regression Models and Utilities for Repeated Measures and Panel Data

Version 0.7.4

Description Provides an object type and associated tools for storing and wrangling panel data. Implements several methods for creating regression models that take advantage of the unique aspects of panel data. Among other capabilities, automates the "within-between" (also known as "between-within" and "hybrid") panel regression specification that combines the desirable aspects of both fixed effects and random effects econometric models and fits them as multilevel models (Allison, 2009 <doi:10.4135/9781412993869.d33>; Bell & Jones, 2015 <doi:10.1017/psrm.2014.7>). These models can also be estimated via generalized estimating equations (GEE; McNeish, 2019 <doi:10.1080/00273171.2019.1602504>) and Bayesian estimation is (optionally) supported via 'Stan'. Supports estimation of asymmetric effects models via first differences (Allison, 2019 <doi:10.1177/2378023119826441>) as well as a generalized linear model extension thereof using GEE.

URL https://panelr.jacob-long.com

BugReports https://github.com/jacob-long/panelr

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Author Jacob A. Long [aut, cre] (<https://orcid.org/0000-0002-1582-6214>)

Maintainer Jacob A. Long <jacob.long@sc.edu>

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are_varying  Check if variables are constant or variable over time.

Description
This function is designed for use with panel_data() objects.

Usage
are_varying(data, ..., type = "time")

Arguments
- data: A data frame, typically of panel_data() class.
- ...: Variable names. If none are given, all variables are checked.
- type: Check for variance over time or across individuals? Default is "time". "individual" considers variables like age to be non-varying because everyone ages at the same speed.

Value
A named logical vector. If TRUE, the variable is varying.

Examples
wages <- panel_data(WageData, id = id, wave = t)
wages %>% are_varying(occ, ind, fem, blk)

asym  Estimate asymmetric effects models using first differences

Description
The function fits the asymmetric effects first difference model described in Allison (2019) using GLS estimation.
Usage

```r
asym(
  formula,
  data,
  id = NULL,
  wave = NULL,
  use.wave = FALSE,
  min.waves = 1,
  variance = c("toeplitz-1", "constrained", "unconstrained"),
  error.type = c("CR2", "CR1S"),
  ...
)
```

Arguments

- **formula**: Model formula. See details for crucial info on panelr’s formula syntax.
- **data**: The data, either a panel_data object or data.frame.
- **id**: If data is not a panel_data object, then the name of the individual id column as a string. Otherwise, leave as NULL, the default.
- **wave**: If data is not a panel_data object, then the name of the panel wave column as a string. Otherwise, leave as NULL, the default.
- **use.wave**: Should the wave be included as a predictor? Default is FALSE.
- **min.waves**: What is the minimum number of waves an individual must have participated in to be included in the analysis? Default is 2 and any valid number is accepted. "all" is also acceptable if you want to include only complete panelists.
- **variance**: One of "toeplitz-1", "constrained", or "unconstrained". The toeplitz variance specification estimates a single error variance and a single lag-1 error correlation with other lags having zero correlation. The constrained model assumes no autocorrelated errors or heteroskedastic errors. The unconstrained option allows separate variances for every period as well as every lag of autocorrelation. This can be very computationally taxing as periods increase and will be inefficient when not necessary. See Allison (2019) for more.
- **error.type**: Either "CR2" or "CR1S". See the clubSandwich package for more details.
- **...**: Ignored.

References


Examples

```r
## Not run:
data("teen_poverty")
# Convert to long format
teen <- long_panel(teen_poverty, begin = 1, end = 5)
model <- asym(hours ~ lag(pov) + spouse, data = teen)
```
asyg.gee

summary(model)

## End(Not run)

asyg.gee

Asymmetric effects models fit with GEE

Description

Fit "within-between" and several other regression variants for panel data via generalized estimating equations.

Usage

asyg.gee(
  formula,
  data,
  id = NULL,
  wave = NULL,
  cor.str = c("ar1", "exchangeable", "unstructured"),
  use.wave = FALSE,
  wave.factor = FALSE,
  min.waves = 1,
  family = gaussian,
  weights = NULL,
  offset = NULL,
  ...
)

Arguments

formula Model formula. See details for crucial info on panelr's formula syntax.
data The data, either a panel_data object or data.frame.
id If data is not a panel_data object, then the name of the individual id column as a string. Otherwise, leave as NULL, the default.
wave If data is not a panel_data object, then the name of the panel wave column as a string. Otherwise, leave as NULL, the default.
cor.str Any correlation structure accepted by geepack::geeglm(). Default is "ar1", most useful alternative is "exchangeable". "unstructured" may cause problems due to its computational complexity.
use.wave Should the wave be included as a predictor? Default is FALSE.
wave.factor Should the wave variable be treated as an unordered factor instead of continuous? Default is FALSE.
min.waves  What is the minimum number of waves an individual must have participated in to be included in the analysis? Default is 2 and any valid number is accepted. “all” is also acceptable if you want to include only complete panelists.

family  Use this to specify GLM link families. Default is gaussian, the linear model.

weights  If using weights, either the name of the column in the data that contains the weights or a vector of the weights.

offset  this can be used to specify an a priori known component to be included in the linear predictor during fitting. This should be NULL or a numeric vector of length equal to the number of cases. One or more offset terms can be included in the formula instead or as well, and if more than one is specified their sum is used. See model.offset.

...  Additional arguments provided to geepack::geeglm().

Details

See the documentation for wbm() for many details on formula syntax and other arguments.

Value

An asym_gge object, which inherits from wbgee and geeglm.

Author(s)

Jacob A. Long

References


Examples

data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- asym_gge(lwage ~ lag(union) + wks, data = wages)
summary(model)
**complete_data**

*Filter out entities with too few observations*

**Description**

This function allows you to define a minimum number of waves/periods and exclude all individuals with fewer observations than that.

**Usage**

```
complete_data(data, ..., formula = NULL, vars = NULL, min.waves = "all")
```

**Arguments**

- `data`: A `panel_data()` frame.
- `...`: Optionally, unquoted variable names/expressions separated by commas to be passed to `dplyr::select()`. Otherwise, all columns are included if `formula` and `vars` are also `NULL`.
- `formula`: A formula, like the one you’ll be using to specify your model.
- `vars`: As an alternative to `formula`, a vector of variable names.
- `min.waves`: What is the minimum number of observations to be kept? Default is "all", but it can be any number.

**Details**

If `...` (that is, unquoted variable name(s)) are included, then `formula` and `vars` are ignored. Likewise, `formula` takes precedence over `vars`. These are just different methods for selecting variables and you can choose whichever you prefer/are comfortable with. `...` corresponds with the "tidyverse" way, `formula` is useful for programming or working with model formulas, and `vars` is a "standard" evaluation method for when you are working with strings.

**Value**

A `panel_data` frame.

**Examples**

```r
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
complete_data(wages, wks, lwage, min.waves = 3)
```
Estimate first differences models using GLS

Description

The function fits first difference models using GLS estimation.

Usage

fdm(
  formula,
  data,
  id = NULL,
  wave = NULL,
  use.wave = FALSE,
  min.waves = 1,
  variance = c("toeplitz-1", "constrained", "unconstrained"),
  error.type = c("CR2", "CR1S"),
  ...
)

Arguments

formula Model formula. See details for crucial info on panelr’s formula syntax.
data The data, either a panel_data object or data.frame.
id If data is not a panel_data object, then the name of the individual id column as a string. Otherwise, leave as NULL, the default.
wave If data is not a panel_data object, then the name of the panel wave column as a string. Otherwise, leave as NULL, the default.
use.wave Should the wave be included as a predictor? Default is FALSE.
min.waves What is the minimum number of waves an individual must have participated in to be included in the analysis? Default is 2 and any valid number is accepted. "all" is also acceptable if you want to include only complete panelists.
variance One of "toeplitz-1", "constrained", or "unconstrained". The toeplitz variance specification estimates a single error variance and a single lag-1 error correlation with other lags having zero correlation. The constrained model assumes no autocorrelated errors or heteroskedastic errors. The unconstrained option allows separate variances for every period as well as every lag of autocorrelation. This can be very computationally taxing as periods increase and will be inefficient when not necessary. See Allison (2019) for more.
error.type Either "CR2" or "CR1S". See the clubSandwich package for more details.
...

References

Examples

data("teen_poverty")
# Convert to long format
teen <- long_panel(teen_poverty, begin = 1, end = 5)
model <- fdm(hours ~ lag(pov) + spouse, data = teen)
summary(model)

get_wave

Retrieve panel_data metadata

Description

get_id(), get_wave(), and get_periods() are extractor functions that can be used to retrieve the names of the id and wave variables or time periods of a panel_data frame.
Usage

get_wave(data)
get_id(data)
get_periods(data)

Arguments

data  A panel_data frame

Value

A panel_data frame

Examples

data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
get_wave(wages)
get_id(wages)
get_periods(wages)

heise  Estimate Heise stability and reliability coefficients

Description

This function uses three waves of data to estimate stability and reliability coefficients as described in Heise (1969).

Usage

heise(data, ..., waves = NULL)

Arguments

data  A panel_data frame.
...  unquoted variable names that are passed to dplyr::select()
waves  Which 3 waves should be used? If NULL (the default), the first, middle, and last waves are used.

Value

A tibble with reliability (rel), waves 1-3 stability (stab13), waves 1-2 stability (stab12), and waves 2-3 stability (stab23) and the variable these values refer to (var).
References

Examples
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
heise(wages, wks, lwage) # will use waves 1, 4, and 7 by default

---

**is_panel**

Check if object is panel_data

**Description**
This is a convenience function that checks whether an object is a panel_data object.

**Usage**

`is_panel(x)`

**Arguments**

- `x`: Any object.

**Examples**

data("WageData")
isi_panel(WageData) # FALSE
wages <- panel_data(WageData, id = id, wave = t)
isi_panel(wages) # TRUE

---

**line_plot**

Plot trends in longitudinal variables

**Description**

line_plot allows for flexible visualization of repeated measures variables from panel_data frames.
line_plot

Usage

line_plot(
  data,
  var,
  id = NULL,
  wave = NULL,
  overlay = TRUE,
  show.points = TRUE,
  subset.ids = FALSE,
  n.random.subset = 9,
  add.mean = FALSE,
  mean.function = "lm",
  line.size = 1,
  alpha = if (overlay) 0.5 else 1
)

Arguments

data Either a panel_data frame or another data frame.
var The unquoted name of the variable of interest.
id If data is not a panel_data object, then the id variable.
wave If data is not a panel_data object, then the wave variable.
overlay Should the lines be plotted in the same panel or each in their own facet/panel? Default is TRUE, meaning they are plotted in the same panel.
show.points Plot a point at each wave? Default is TRUE.
subset.ids Plot only a subset of the entities’ lines? Default is NULL, meaning plot all ids. If TRUE, a random subset (the number defined by n.random.subset) are plotted. You may also supply a vector of ids to choose them yourself.
n.random.subset How many entities to randomly sample when subset.ids is TRUE.
add.mean Add a line representing the mean trend? Default is FALSE. Cannot be combined with overlay.
mean.function The mean function to supply to geom_smooth when add.mean is TRUE. Default is "lm", but another option of interest is "loess".
line.size The thickness of the plotted lines. Default: 0.5
alpha The transparency for the lines and points. When overlay = TRUE, it is set to 0.5, otherwise 1, which means non-transparent.

Value

The ggplot object.
long_panel

Examples

```r
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
line_plot(wages, lwage, add.mean = TRUE, subset.ids = TRUE, overlay = FALSE)
```

long_panel

Convert wide panels to long format

Description

This function takes wide format panels as input and converts them to long format.

Usage

```r
long_panel(
  data,
  prefix = NULL,
  suffix = NULL,
  begin = NULL,
  end = NULL,
  id = "id",
  wave = "wave",
  periods = NULL,
  label_location = c("end", "beginning"),
  as_panel_data = TRUE,
  match = ".*",
  use.regex = FALSE,
  check.varying = TRUE
)
```

Arguments

data The wide data frame.

prefix What character(s) go before the period indicator? If none, set this argument to NULL.

suffix What character(s) go after the period indicator? If none, set this argument to NULL.

begin What is the label for the first period? Could be 1, "A", or anything that can be sequenced.

end What is the label for the final period? Could be 2, "B", or anything that can be sequenced and lies further along the sequence than the begin argument.

id The name of the ID variable as a string. If there is no ID variable, then this will be the name of the newly-created ID variable.
wave This will be the name of the newly-created wave variable.

periods If you period indicator does not lie in a sequence or is not understood by the function, then you can supply them as a vector instead. For instance, you could give c("one","three","five") if your variables are labeled var_one, var_three, and var_five.

label_location Where does the period label go on the variable? If the variables are labeled like var_1, var_2, etc., then it is "end". If the labels are more like A_var, B_var, and so on, then it is "beginning".

as_panel_data Should the return object be a panel_data() object? Default is TRUE.

match The regex that will match the part of the variable names other than the wave indicator. By default it will match any character any amount of times. Sometimes you might know that the variable names should start with a digit, for instance, and you might use "\d.*" instead.

use.regex Should the begin and end arguments be treated as regular expressions? Default is FALSE.

check.varying Should the function check to make sure that every variable in the wide data with a wave indicator is actually time-varying? Default is TRUE, meaning that a constant like "race_W1" only measured in wave 1 will be defined in each wave in the long data. With very large datasets, however, sometimes setting this to FALSE can save memory.

Details

There is no easy way to convert panel data from wide to long format because the both formats are basically non-standard for other applications. This function can handle the common case in which the wide data frame has a regular labeling system for each period. The key thing is providing enough information for the function to understand the pattern.

In the end, this function calls stats::reshape() but should be easier to use and able to handle more situations, such as when the label occurs at the beginning of the variable name. Also, just as important, this function has built-in utilities to handle unbalanced data — when variables occur more than once but every single period, which breaks stats::reshape().

Value

Either a data.frame or panel_data frame.

See Also

widen_panel()

Examples

## We need a wide data frame, so we will make one from the long-format data included in the package.

# Convert WageData to panel_data object
wages <- panel_data(WageData, id = id, wave = t)
# Convert wages to wide format
wide_wages <- widen_panel(wages)

# Note: wide_wages has variables in the following format:
# var1_1, var1_2, var1_3, var2_1, var2_2, var2_3, etc.
## Not run:
long_wages <- long_panel(wide_wages, prefix = "_", begin = 1, end = 7,
                        id = "id", label_location = "end")

## End(Not run)
# Note that in this case, the prefix and label_location arguments are
# the defaults but are included just for clarity.

make_diff_data

Generate differenced and asymmetric effects data

Description

This is an interface to the internal functions that process data for fdm(), asym(), and asym.gee().

Usage

make_diff_data(
  formula,
  data,
  id = NULL,
  wave = NULL,
  use.wave = FALSE,
  min.waves = 1,
  weights = NULL,
  offset = NULL,
  asym = FALSE,
  cumulative = FALSE,
  escape.names = FALSE,
  ...
)

Arguments

formula  Model formula. See details for crucial info on panelr’s formula syntax.
data     The data, either a panel_data object or data.frame.
id       If data is not a panel_data object, then the name of the individual id column
          as a string. Otherwise, leave as NULL, the default.
wave     If data is not a panel_data object, then the name of the panel wave column as
          a string. Otherwise, leave as NULL, the default.
use.wave Should the wave be included as a predictor? Default is FALSE.
min.waves What is the minimum number of waves an individual must have participated in to be included in the analysis? Default is 2 and any valid number is accepted. "all" is also acceptable if you want to include only complete panelists.
weights If using weights, either the name of the column in the data that contains the weights or a vector of the weights.
offset this can be used to specify an \textit{a priori} known component to be included in the linear predictor during fitting. This should be NULL or a numeric vector of length equal to the number of cases. One or more offset terms can be included in the formula instead or as well, and if more than one is specified their sum is used. See \code{model.offset}.
asy Return asymmetric effects transformed data? Default is FALSE.
cumulative Return cumulative positive/negative differences, most useful for fixed effects estimation and/or generalized linear models? Default is FALSE.
escape.names Return only syntactically valid variable names? Default is FALSE.
... Ignored.

\section*{Examples}
\begin{verbatim}
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
make_diff_data(wks ~ lwage + union, data = wages)
\end{verbatim}

\section*{Description}
This function allows users to make the changes to their data that occur in \code{wbm()} without having to fit the model.

\section*{Usage}
\begin{verbatim}
make_wb_data(
  formula,
  data,
  id = NULL,
  wave = NULL,
  model = "w-b",
  detrend = FALSE,
  use.wave = FALSE,
  wave.factor = FALSE,
  min.waves = 2,
\end{verbatim}
make_wb_data

balance.correction = FALSE,
dt.random = TRUE,
dt.order = 1,
weights = NULL,
offset = NULL,
interaction.style = c("double-demean", "demean", "raw"),
...
)

Arguments

formula Model formula. See details for crucial info on panelr’s formula syntax.
data The data, either a panel_data object or data.frame.
id If data is not a panel_data object, then the name of the individual id column as a string. Otherwise, leave as NULL, the default.
wave If data is not a panel_data object, then the name of the panel wave column as a string. Otherwise, leave as NULL, the default.
model One of "w-b", "within", "between", "contextual". See details for more on these options.
detrend Adjust within-subject effects for trends in the predictors? Default is FALSE, but some research suggests this is a better idea (see Curran and Bauer (2011) reference).
use.wave Should the wave be included as a predictor? Default is FALSE.
wave.factor Should the wave variable be treated as an unordered factor instead of continuous? Default is FALSE.
min.waves What is the minimum number of waves an individual must have participated in to be included in the analysis? Default is 2 and any valid number is accepted. "all" is also acceptable if you want to include only complete panelists.
balance.correction Correct between-subject effects for unbalanced panels following the procedure in Curran and Bauer (2011)? Default is FALSE.
dt.random Should the detrending procedure be performed with a random slope for each entity? Default is TRUE but for short panels FALSE may be better, fitting a trend for all entities.
dt.order If detrending using detrend, what order polynomial would you like to specify for the relationship between time and the predictors? Default is 1, a linear model.
weights If using weights, either the name of the column in the data that contains the weights or a vector of the weights.
offset this can be used to specify an a priori known component to be included in the linear predictor during fitting. This should be NULL or a numeric vector of length equal to the number of cases. One or more offset terms can be included in the formula instead or as well, and if more than one is specified their sum is used. See model.offset.
The best way to calculate interactions in within models is in some dispute. The conventional way ("demean") is to first calculate the product of the variables involved in the interaction before those variables have their means subtracted and then subtract the mean of the product from the product term (see Schunk and Perales (2017)). Giesselmann and Schmidt-Catran (2020) show this method carries between-entity differences that within models are designed to model out. They suggest an alternate method ("double-demean") in which the product term is first calculated using the de-meaned lower-order variables and then the subject means are subtracted from this product term. Another option is to simply use the product term of the de-meaned variables ("raw"), but Giesselmann and Schmidt-Catran (2020) show this method biases the results towards zero effect. The default is "double-demean" but if emulating other software is the goal, "demean" might be preferred.

Additional arguments provided to \texttt{lme4::lmer()}, \texttt{lme4::glmer()}, or \texttt{lme4::glmer.nb()}. 

\section*{Value}

A \texttt{panel_data} object with the requested specification.

\section*{Examples}

```r
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
make_wb_data(lwage ~ wks + union | fem, data = wages)
```

\section*{Description}

This is similar to \texttt{model.frame}, but is designed specifically for \texttt{panel_data()} data frames. It’s a workhorse in \texttt{wbm()} but may be useful in scripting use as well.

\section*{Usage}

\texttt{model_frame(formula, data)}

\section*{Arguments}

\begin{itemize}
  \item \texttt{formula} A formula. Note that to get an individual-level mean with incomplete data (e.g., panel attrition), you should use \texttt{imean()} rather than \texttt{mean()}.
  \item \texttt{data} A \texttt{panel_data()} frame.
\end{itemize}

\section*{Value}

A \texttt{panel_data()} frame with only the columns needed to fit a model as described by the formula.
Examples

data("WageData")
  wages <- panel_data(WageData, id = id, wave = t)
  model_frame(lwage ~ wks + exp, data = wages)

Description

These data come from the years 1990-1994 in the National Longitudinal Survey of Youth, with information about 581 individuals. These data are in the "wide" format for demonstration purposes.

Usage

nlsy

Format

A data frame with 581 rows and 16 variables:

  momage  Mother's age at birth
  gender  0 if boy, 1 if girl
  momwork 1 if mother works, 0 if not
  married 1 if parents are married, 0 if not
  hispanic 1 if child is Hispanic, 0 if not
  black 1 if child is black, 0 if not
  childage Child's age at first interview
  anti90 A measure of anti-social behavior antisocial behavior measured on a scale from 0 to 6, taken in 1990
  anti92 A measure of anti-social behavior antisocial behavior measured on a scale from 0 to 6, taken in 1992
  anti94 A measure of anti-social behavior antisocial behavior measured on a scale from 0 to 6, taken in 1994
  self90 A measure of self-esteem measured on a scale from 6 to 24, taken in 1990
  self92 A measure of self-esteem measured on a scale from 6 to 24, taken in 1992
  self94 A measure of self-esteem measured on a scale from 6 to 24, taken in 1994
  pov90 1 if family is in poverty, 0 if not, in 1990
  pov92 1 if family is in poverty, 0 if not, in 1992
  pov94 1 if family is in poverty, 0 if not, in 1994
Source
These data originate with the U.S. Department of Labor. The particular subset used here come from Paul Allison via Statistical Horizons: https://statisticalhorizons.com/wp-content/uploads/nlsy.dta

---

### nobs.wbm

**Number of observations used in wbm models**

---

**Description**

This S3 method allows you to retrieve either the number of observations or number of entities in the data used to fit wbm objects.

**Usage**

```r
## S3 method for class 'wbm'
nobs(object, entities = TRUE, ...)
```

**Arguments**

- `object`: A fitted model object.
- `entities`: Should `nobs` return the number of entities in the panel or the number of rows in the panel_data frame? Default is TRUE, returning the number of entities.
- `...`: Further arguments to be passed to methods.

**Examples**

```r
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- wbm(lwage ~ lag(union) + wks, data = wages)
nobs(model)
```

---

### panel_data

**Create panel data frames**

---

**Description**

Format your data for use with **panelr**.
Usage

panel_data(data, id = id, wave = wave, ...)

as_pdata.frame(data)

as_panel_data(data, ...)

## Default S3 method:
as_panel_data(data, id = id, wave = wave, ...)

## S3 method for class 'pdata.frame'
as_panel_data(data, ...)

as_panel(data, ...)

Arguments

data A data frame.
id The name of the column (unquoted) that identifies participants/entities. A new column will be created called id, overwriting any column that already has that name.
wave The name of the column (unquoted) that identifies waves or periods. A new column will be created called wave, overwriting any column that already has that name.
...
Attributes for adding onto this method. See tibble::new_tibble() for a run-through of the logic.

Value

A panel_data object.

Examples

data("WageData")
wages <- panel_data(WageData, id = id, wave = t)

---

predict.wbgee Predictions and simulations from within-between GEE models

Description

These methods facilitate fairly straightforward predictions from wbgee models.
Usage

## S3 method for class 'wbgee'
predict(
  object,
  newdata = NULL,
  se.fit = FALSE,
  raw = FALSE,
  type = c("link", "response"),
  ...
)

Arguments

object Object of class inheriting from "lm"
newdata An optional data frame in which to look for variables with which to predict. If
  omitted, the fitted values are used.
se.fit A switch indicating if standard errors are required.
raw Is newdata a geeglm model frame or panel_data? TRUE indicates a geeglm-
  style newdata, with all of the extra columns created by wbgee.
type Type of prediction (response or model term). Can be abbreviated.
...
  further arguments passed to or from other methods.

Examples

data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- wbgee(lwage ~ lag(union) + wks, data = wages)
# By default, assumes you're using the processed data for newdata
predict(model)

Description

These methods facilitate fairly straightforward predictions and simulations from wbm models.

Usage

## S3 method for class 'wbm'
predict(
  object,
  newdata = NULL,
  se.fit = FALSE,
  raw = FALSE,
  use.re.var = FALSE,
**predict.wbm**

```r
re.form = NULL,
type = c("link", "response"),
allow.new.levels = TRUE,
na.action = na.pass,
...)

## S3 method for class 'wbm'
simulate(
  object,
  nsim = 1,
  seed = NULL,
  use.u = FALSE,
  newdata = NULL,
  raw = FALSE,
  newparams = NULL,
  re.form = NA,
  type = c("link", "response"),
  allow.new.levels = FALSE,
  na.action = na.pass,
  ...
)
```

### Arguments

- **object**: a fitted model object
- **newdata**: data frame for which to evaluate predictions.
- **se.fit**: Include standard errors with the predictions? Note that these standard errors by default include only fixed effects variance. See details for more info. Default is FALSE.
- **raw**: Is newdata a merMod model frame or panel_data? TRUE indicates a merMod-style newdata, with all of the extra columns created by wbm.
- **use.re.var**: If se.fit is TRUE, include random effects variance in standard errors? Default is FALSE.
- **re.form**: (formula, NULL, or NA) specify which random effects to condition on when predicting. If NULL, include all random effects; if NA or ~0, include no random effects.
- **type**: character string - either "link", the default, or "response" indicating the type of prediction object returned.
- **allow.new.levels**: logical if new levels (or NA values) in newdata are allowed. If FALSE (default), such new values in newdata will trigger an error; if TRUE, then the prediction will use the unconditional (population-level) values for data with previously unobserved levels (or NAs).
- **na.action**: function determining what should be done with missing values for fixed effects in newdata. The default is to predict NA: see na.pass.
... When `boot` and `se.fit` are TRUE, any additional arguments are passed to `lme4::bootMer()`.

- **nsim** positive integer scalar - the number of responses to simulate.
- **seed** an optional seed to be used in `set.seed` immediately before the simulation so as to generate a reproducible sample.
- **use.u** (logical) if TRUE, generate a simulation conditional on the current random-effects estimates; if FALSE generate new Normally distributed random-effects values. (Redundant with `re.form`, which is preferred: TRUE corresponds to `re.form = NULL` (condition on all random effects), while FALSE corresponds to `re.form = ~0` (condition on none of the random effects).)
- **newparams** new parameters to use in evaluating predictions, specified as in the `start` parameter for `lmer` or `glmer` – a list with components `theta` and `beta` and (for LMMs or GLMMs that estimate a scale parameter) `sigma`

**Examples**

```r
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- wbm(lwage ~ lag(union) + wks, data = wages)
# By default, assumes you're using the processed data for newdata
predict(model)
```

**Description**

`summary` method for `panel_data` objects.

**Usage**

```r
## S3 method for class 'panel_data'
summary(object, ..., by.wave = TRUE, by.id = FALSE)
```

**Arguments**

- **object** A `panel_data` frame.
- **...** Optionally, unquoted variable names/expressions separated by commas to be passed to `dplyr::select()`. Otherwise, all columns are included.
- **by.wave** (if `skimr` is installed) Separate descriptives by wave? Default is TRUE.
- **by.id** (if `skimr` is installed) Separate descriptives by entity? Default is FALSE. Be careful if you have a large number of entities as the output will be massive.
Examples

data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
summary(wages, lwage, exp, wks)

Description

These data come from the years 1979-1983 in the National Longitudinal Survey of Youth, with information about 1141 teenage women. These data are in the "wide" format for demonstration purposes.

Usage

teen_poverty

Format

A data frame with 1141 rows and 28 variables:

- id  Unique identifier for the respondent
- age  Age at first interview
- black  1 if subject is black, 0 if not
- pov1  1 if subject is in poverty, 0 if not, at time 1
- pov2  1 if subject is in poverty, 0 if not, at time 2
- pov3  1 if subject is in poverty, 0 if not, at time 3
- pov4  1 if subject is in poverty, 0 if not, at time 4
- pov5  1 if subject is in poverty, 0 if not, at time 5
- mother1  1 if subject has had a child, 0 if not, at time 1
- mother2  1 if subject has had a child, 0 if not, at time 2
- mother3  1 if subject has had a child, 0 if not, at time 3
- mother4  1 if subject has had a child, 0 if not, at time 4
- mother5  1 if subject has had a child, 0 if not, at time 5
- spouse1  1 if subject lives with a spouse, 0 if not, at time 1
- spouse2  1 if subject lives with a spouse, 0 if not, at time 2
- spouse3  1 if subject lives with a spouse, 0 if not, at time 3
- spouse4  1 if subject lives with a spouse, 0 if not, at time 4
- spouse5  1 if subject lives with a spouse, 0 if not, at time 5
inschool1 1 if subject is in school, 0 if not, at time 1
inschool2 1 if subject is in school, 0 if not, at time 2
inschool3 1 if subject is in school, 0 if not, at time 3
inschool4 1 if subject is in school, 0 if not, at time 4
inschool5 1 if subject is in school, 0 if not, at time 5
hours1 Hours worked during the week of the survey, at time 1
hours2 Hours worked during the week of the survey, at time 2
hours3 Hours worked during the week of the survey, at time 3
hours4 Hours worked during the week of the survey, at time 4
hours5 Hours worked during the week of the survey, at time 5

Source

These data originate with the U.S. Department of Labor. The particular subset used here come from Paul Allison via Statistical Horizons: [https://statisticalhorizons.com/wp-content/uploads/teenpov.dta](https://statisticalhorizons.com/wp-content/uploads/teenpov.dta)

---

tidy.asym

Tidy methods for fdm and asym models

Description

panelr provides methods to access fdm and asym data in a tidy format

Usage

```r
## S3 method for class 'asym'
tidy(x, conf.int = FALSE, conf.level = 0.95, ...)

## S3 method for class 'fdm'
tidy(x, conf.int = FALSE, conf.level = 0.95, ...)

## S3 method for class 'fdm'
glance(x, ...)
```

Arguments

- **x**
  - An fdm or asym object.
- **conf.int**
  - Logical indicating whether or not to include a confidence interval in the tidied output. Defaults to FALSE.
- **conf.level**
  - The confidence level to use for the confidence interval if conf.int = TRUE. Must be strictly greater than 0 and less than 1. Defaults to 0.95, which corresponds to a 95 percent confidence interval.
- **...**
  - Ignored
Examples

```r
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- fdm(lwage ~ wks + union, data = wages)
if (requireNamespace("generics")) {
  generics::tidy(model)
}
```

Description

`panelr` provides methods to access `wbgee` data in a tidy format.

Usage

```r
## S3 method for class 'asym_gee'
tidy(x, conf.int = FALSE, conf.level = 0.95, ...)
## S3 method for class 'wbgee'
tidy(x, conf.int = FALSE, conf.level = 0.95, ...)
## S3 method for class 'wbgee'
glance(x, ...)
```

Arguments

- `x` A `wbgee` object.
- `conf.int` Logical indicating whether or not to include a confidence interval in the tidied output. Defaults to `FALSE`.
- `conf.level` The confidence level to use for the confidence interval if `conf.int = TRUE`. Must be strictly greater than 0 and less than 1. Defaults to 0.95, which corresponds to a 95 percent confidence interval.
- `...` Ignored

Examples

```r
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- wbgee(lwage ~ lag(union) + wks, data = wages)
if (requireNamespace("generics")) {
  generics::tidy(model)
}
```
Tidy methods for wbm models

Description

`panelr` provides methods to access `wbm` data in a tidy format.

Usage

```r
## S3 method for class 'wbm'
tidy(
  x,
  conf.int = FALSE,
  conf.level = 0.95,
  effects = c("fixed", "ran_pars"),
  conf.method = "Wald",
  ran_prefix = NULL,
  ...
)
## S3 method for class 'wbm'
glance(x, ...)
## S3 method for class 'summ.wbm'
glance(x, ...)
## S3 method for class 'summ.wbm'
tidy(x, ...)
```

Arguments

- `x` An object of class `merMod`, such as those from `lmer`, `glmer`, or `nlmer`
- `conf.int` whether to include a confidence interval
- `conf.level` confidence level for CI
- `effects` A character vector including one or more of "fixed" (fixed-effect parameters); "ran_pars" (variances and covariances or standard deviations and correlations of random effect terms); "ran_vals" (conditional modes/BLUPs/latent variable estimates); or "ran_coefs" (predicted parameter values for each group, as returned by `coef.merMod`.
- `conf.method` method for computing confidence intervals (see `lme4::confint.merMod`)
- `ran_prefix` a length-2 character vector specifying the strings to use as prefixes for self-(variance/standard deviation) and cross- (covariance/correlation) random effects terms
- `...` Additional arguments (passed to `confint.merMod` for `tidy`; `augment_columns` for `augment`; ignored for `glance`
unpanel

Examples

```
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- wbm(lwage ~ lag(union) + wks, data = wages)
if (requireNamespace("broom.mixed")) {
  broom.mixed::tidy(model)
}
```

---

**unpanel**

Convert panel_data to regular data frame

---

**Description**

This convenience function removes the special features of `panel_data`.

**Usage**

```
unpanel(panel)
```

**Arguments**

- `panel` A `panel_data` object.

**Value**

An ungrouped tibble.

**Examples**

```
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
wages_non_panel <- unpanel(wages)
```

---

**WageData**

Earnings data from the Panel Study of Income Dynamics

---

**Description**

These data come from the years 1976-1982 in the Panel Study of Income Dynamics (PSID), with information about the demographics and earnings of 595 individuals.

**Usage**

```
WageData
```
Format

A data frame with 4165 rows and 14 variables:

- **id**: Unique identifier for each survey respondent
- **t**: A number corresponding to each wave of the survey, 1 through 7
- **wks**: Weeks worked in the past year
- **lwage**: Natural logarithm of earnings in the past year
- **union**: Binary indicator whether respondent is a member of union (1 = union member)
- **ms**: Binary indicator for whether respondent is married (1 = married)
- **occ**: Binary indicator for whether respondent is a blue collar (= 0) or white collar (= 1) worker.
- **ind**: Binary indicator for whether respondent works in manufacturing (= 1)
- **south**: Binary indicator for whether respondent lives in the South (= 1)
- **smsa**: Binary indicator for whether respondent lives in a standard metropolitan area (SMSA; = 1)
- **fem**: Binary indicator for whether respondent is female (= 1)
- **blk**: Binary indicator for whether respondent is African-American (= 1)
- **ed**: Years of education
- **exp**: Years in the workforce.

Source

These data are all over the place. This particular file was downloaded from Richard Williams at http://www3.nd.edu/~rwilliam/statafiles/wages.dta, though he doesn’t claim ownership of these data.

The data were shared as a supplement to Baltagi (2005) at https://www.wiley.com/legacy/wileychi/baltagi3e/data_sets.html.

They were also shared as a supplement to Greene (2008) at http://pages.stern.nyu.edu/~wgreene/Text/Edition6/tablelist6.htm.

The data are also available in numerous other locations, including in slightly different formats as Wages in the plm package and PSID7682 in the AER package.

---

**wbgee**

*Panel regression models fit with GEE*

Description

Fit "within-between" and several other regression variants for panel data via generalized estimating equations.
wbgee

Usage

wbgee(
  formula,
  data,
  id = NULL,
  wave = NULL,
  model = "w-b",
  cor.str = c("ar1", "exchangeable", "unstructured"),
  detrend = FALSE,
  use.wave = FALSE,
  wave.factor = FALSE,
  min.waves = 2,
  family = gaussian,
  balance.correction = FALSE,
  dt.random = TRUE,
  dt.order = 1,
  weights = NULL,
  offset = NULL,
  interaction.style = c("double-demean", "demean", "raw"),
  scale = FALSE,
  scale.response = FALSE,
  n.sd = 1,
  calc.fit.stats = TRUE,
  ...
)

Arguments

formula       Model formula. See details for crucial info on panelr’s formula syntax.
data          The data, either a panel_data object or data.frame.
id            If data is not a panel_data object, then the name of the individual id column as a string. Otherwise, leave as NULL, the default.
wave          If data is not a panel_data object, then the name of the panel wave column as a string. Otherwise, leave as NULL, the default.
model         One of "w-b", "within", "between", "contextual". See details for more on these options.
cor.str       Any correlation structure accepted by geepack::geeglm(). Default is "ar1", most useful alternative is "exchangeable". "unstructured" may cause problems due to its computational complexity.
detrend       Adjust within-subject effects for trends in the predictors? Default is FALSE, but some research suggests this is a better idea (see Curran and Bauer (2011) reference).
use.wave      Should the wave be included as a predictor? Default is FALSE.
wave.factor   Should the wave variable be treated as an unordered factor instead of continuous? Default is FALSE.
min.waves What is the minimum number of waves an individual must have participated in to be included in the analysis? Default is 2 and any valid number is accepted. “all” is also acceptable if you want to include only complete panelists.

family Use this to specify GLM link families. Default is gaussian, the linear model.

balance.correction Correct between-subject effects for unbalanced panels following the procedure in Curran and Bauer (2011)? Default is FALSE.

dt.random Should the detrending procedure be performed with a random slope for each entity? Default is TRUE but for short panels FALSE may be better, fitting a trend for all entities.

dt.order If detrending using detrend, what order polynomial would you like to specify for the relationship between time and the predictors? Default is 1, a linear model.

weights If using weights, either the name of the column in the data that contains the weights or a vector of the weights.

offset this can be used to specify an a priori known component to be included in the linear predictor during fitting. This should be NULL or a numeric vector of length equal to the number of cases. One or more offset terms can be included in the formula instead or as well, and if more than one is specified their sum is used. See model.offset.

interaction.style The best way to calculate interactions in within models is in some dispute. The conventional way ("demean") is to first calculate the product of the variables involved in the interaction before those variables have their means subtracted and then subtract the mean of the product from the product term (see Schunk and Perales (2017)). Giesselmann and Schmidt-Catran (2020) show this method carries between-entity differences that within models are designed to model out. They suggest an alternate method ("double-demean") in which the product term is first calculated using the de-meaned lower-order variables and then the subject means are subtracted from this product term. Another option is to simply use the product term of the de-meaned variables ("raw"), but Giesselmann and Schmidt-Catran (2020) show this method biases the results towards zero effect. The default is "double-demean" but if emulating other software is the goal, "demean" might be preferred.

scale If TRUE, reports standardized regression coefficients. Default is FALSE.

scale.response Should the response variable also be rescaled? Default is FALSE.

n.sd How many standard deviations should you divide by for standardization? Default is 1, though some prefer 2.

calc.fit.stats Calculate fit statistics? Default is TRUE, but occasionally poor-fitting models might trip up here.

... Additional arguments provided to geepack::geglm().

Details

See the documentation for wbm() for many details on formula syntax and other arguments.
Value

A `wbgee` object, which inherits from `geeglm`.

Author(s)

Jacob A. Long

References


Examples

data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- wbgee(lwage ~ lag(union) + wks | blk + fem | blk * lag(union),
              data = wages)
summary(model)

Description

Fit "within-between" and several other regression variants for panel data in a multilevel modeling framework.
Usage

```r
wbm(
  formula,
  data,
  id = NULL,
  wave = NULL,
  model = "w-b",
  detrend = FALSE,
  use.wave = FALSE,
  wave.factor = FALSE,
  min.waves = 2,
  family = gaussian,
  balance.correction = FALSE,
  dt.random = TRUE,
  dt.order = 1,
  pR2 = TRUE,
  pvals = TRUE,
  t.df = "Satterthwaite",
  weights = NULL,
  offset = NULL,
  interaction.style = c("double-demean", "demean", "raw"),
  scale = FALSE,
  scale.response = FALSE,
  n.sd = 1,
  dt_random = dt.random,
  dt_order = dt.order,
  balance.correction = balance.correction,
  ...
)
```

Arguments

- `formula` Model formula. See details for crucial info on panelr’s formula syntax.
- `data` The data, either a `panel_data` object or `data.frame`.
- `id` If `data` is not a `panel_data` object, then the name of the individual id column as a string. Otherwise, leave as NULL, the default.
- `wave` If `data` is not a `panel_data` object, then the name of the panel wave column as a string. Otherwise, leave as NULL, the default.
- `model` One of "w-b", "within", "between", "contextual". See details for more on these options.
- `detrend` Adjust within-subject effects for trends in the predictors? Default is FALSE, but some research suggests this is a better idea (see Curran and Bauer (2011) reference).
- `use.wave` Should the wave be included as a predictor? Default is FALSE.
- `wave.factor` Should the wave variable be treated as an unordered factor instead of continuous? Default is FALSE.
min.waves: What is the minimum number of waves an individual must have participated in to be included in the analysis? Default is 2 and any valid number is accepted. "all" is also acceptable if you want to include only complete panelists.

family: Use this to specify GLM link families. Default is gaussian, the linear model.

balance.correction: Correct between-subject effects for unbalanced panels following the procedure in Curran and Bauer (2011)? Default is FALSE.

dt.random: Should the detrending procedure be performed with a random slope for each entity? Default is TRUE but for short panels FALSE may be better, fitting a trend for all entities.

dt.order: If detrending using detrend, what order polynomial would you like to specify for the relationship between time and the predictors? Default is 1, a linear model.

pR2: Calculate a pseudo R-squared? Default is TRUE, but in some cases may cause errors or add computation time.

pvals: Calculate p values? Default is TRUE but for some complex linear models, this may take a long time to compute using the pbkrtest package.

t.df: For linear models only. User may choose the method for calculating the degrees of freedom in t-tests. Default is "Satterthwaite", but you may also choose "Kenward-Roger". Kenward-Roger standard errors/degrees of freedom requires the pbkrtest package.

weights: If using weights, either the name of the column in the data that contains the weights or a vector of the weights.

offset: this can be used to specify an a priori known component to be included in the linear predictor during fitting. This should be NULL or a numeric vector of length equal to the number of cases. One or more offset terms can be included in the formula instead or as well, and if more than one is specified their sum is used. See model.offset.

interaction.style: The best way to calculate interactions in within models is in some dispute. The conventional way ("demean") is to first calculate the product of the variables involved in the interaction before those variables have their means subtracted and then subtract the mean of the product from the product term (see Schunk and Perales (2017)). Giesselmann and Schmidt-Catran (2020) show this method carries between-entity differences that within models are designed to model out. They suggest an alternate method ("double-demean") in which the product term is first calculated using the de-meaned lower-order variables and then the subject means are subtracted from this product term. Another option is to simply use the product term of the de-meaned variables ("raw"), but Giesselmann and Schmidt-Catran (2020) show this method biases the results towards zero effect. The default is "double-demean" but if emulating other software is the goal, "demean" might be preferred.

scale: If TRUE, reports standardized regression coefficients. Default is FALSE.

scale.response: Should the response variable also be rescaled? Default is FALSE.

n.sd: How many standard deviations should you divide by for standardization? Default is 1, though some prefer 2.
dt_random  Deprecated. Equivalent to dt.random.
dt_order  Deprecated. Equivalent to dt.order.
balance_correction  Deprecated. Equivalent to balance.correction.
...  Additional arguments provided to lme4::lmer(), lme4::glmer(), or lme4::glmer.nb().

Details

Formula syntax

The within-between models, and multilevel panel models more generally, distinguish between time-varying and time-invariant predictors. These are, as they sound, variables that are either measured repeatedly (in every wave) in the case of time-varying predictors or only once in the case of time-invariant predictors. You need to specify these separately in the formula to tell the model which variables you expect to change over time and which will not. The primary way of doing so is via the | operator.

As an example, we can look at the WageData included in this package. We will create a model that predicts the logarithm of the individual’s wages (lwage) with their union status (union), which can change over time, and their race (blk; dichotomized as black or non-black), which does not change throughout the period of study. Our formula will look like this:

lwage ~ union | blk

Put time-varying variables before the first | and time-invariant variables afterwards. You can specify lags like lag(union) for time-varying variables; for more than 1 lag, include the number: lag(union,2).

After the first | go the time-invariant variables. Note that if you put a time-varying variable here, what you get is the observed value rather than one adjusted to isolate within-entity effects. You may also take a time-varying variable — let’s say weeks worked (wks) — and use imean(wks) to include the individual’s mean across all waves as a predictor while omitting the per-wave measures.

There is also a place for a second |. Here you can specify cross-level interactions (within-level interactions can be specified here as well). If I wanted the interaction term for union and blk — to see whether the effect of union status depended on one’s race — I would specify the formula this way:

lwage ~ union | blk | union * blk

Another use for the post-second | section of the formula is for changing the random effects specification. By default, only a random intercept is specified in the call to lme4::lmer() / lme4::glmer(). If you would like to specify other random slopes, include them here using the typical lme4 syntax:

lwage ~ union | blk | (union | id)

You can also include the wave variable in a random effects term to specify a latent growth curve model:

lwage ~ union | blk + t | (t | id)

One last thing to know: If you want to use the second | but not the first, put a 1 or 0 after the first, like this:

lwage ~ union | 1 | (union | id)

Of course, with no time-invariant variables, you need no | operators at all.
As a convenience, wbm does the heavy lifting for specifying the within-between model correctly. As a side effect it only takes a few easy tweaks to specify the model slightly differently. You can change this behavior with the model argument.

By default, the argument is "w-b" (equivalently, "within-between"). This means, for each time-varying predictor, you have two types of variables in the model. The "between" effect is represented by the individual-level mean for each entity (e.g., each respondent to a panel survey). The "within" effect is represented by each wave’s measure with the individual-level mean subtracted. Some refer to this as "de-meaning." Thinking in a Hausman test framework — with the within-between model as described here — you should expect the within and between coefficients to be the same if a random effects model were appropriate.

The contextual model is very similar (use argument "contextual"). In some situations, this will be more intuitive to interpret. Empirically, the only difference compared to the within-between specification is that the contextual model does not subtract the individual-level means from the wave-level measures. This also changes the interpretation of the between-subject coefficients: In the contextual model, they are the difference between the within and between effects. If there’s no difference between within and between effects, then, the coefficients will be 0.

To fit a random effects model, use either "between" or "random". This involves no de-meaning and no individual-level means whatsoever.

To fit a fixed effects model, use either "within" or "fixed". Any between-subjects terms in the formula will be ignored. The time-varying variables will be de-meaned, but the individual-level mean is not included in the model.

A wbm object, which inherits from merMod.

Jacob A. Long


See Also

`wbm_stan()` for a Bayesian estimation option.

Examples

```r
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- wbm(lwage ~ lag(union) + wks | blk + fem | blk * lag(union),
             data = wages)
summary(model)
```

wbm-class

Within-Between Model (wbm) class

Description

Models fit using `wbm()` return values of this class, which inherits from `merMod-class`.

Slots

call_info  A list of metadata about the arguments used.
call      The actual function call.
summ     The `jtools::summ()` object returned from calling it on the `merMod` object.
summ_atts The attributes of the `summ` object.
orig_data The data provided to the `data` argument in the function call.

wbm_stan

Bayesian estimation of within-between models

Description

A near-equivalent of `wbm()` that instead uses Stan, via `rstan` and `brms`.

Usage

```r
wbm_stan(
  formula,
  data,
  id = NULL,
  wave = NULL,
  model = "w-b",
  detrend = FALSE,
  use.wave = FALSE,
```
```r
wave.factor = FALSE,
min.waves = 2,
model.cor = FALSE,
family = gaussian,
fit_model = TRUE,
balance.correction = FALSE,
dt.random = TRUE,
dt.order = 1,
chains = 3,
iter = 2000,
scale = FALSE,
save_ranef = FALSE,
interaction.style = c("double-demean", "demean", "raw"),
weights = NULL,
offset = NULL,
...
)
```

Arguments

- **formula**: Model formula. See details for crucial info on `panelr`'s formula syntax.
- **data**: The data, either a `panel_data` object or `data.frame`.
- **id**: If `data` is not a `panel_data` object, then the name of the individual id column as a string. Otherwise, leave as NULL, the default.
- **wave**: If `data` is not a `panel_data` object, then the name of the panel wave column as a string. Otherwise, leave as NULL, the default.
- **model**: One of "w-b", "within", "between", "contextual". See details for more on these options.
- **detrend**: Adjust within-subject effects for trends in the predictors? Default is FALSE, but some research suggests this is a better idea (see Curran and Bauer (2011) reference).
- **use.wave**: Should the wave be included as a predictor? Default is FALSE.
- **wave.factor**: Should the wave variable be treated as an unordered factor instead of continuous? Default is FALSE.
- **min.waves**: What is the minimum number of waves an individual must have participated in to be included in the analysis? Default is 2 and any valid number is accepted. "all" is also acceptable if you want to include only complete panelists.
- **model.cor**: Do you want to model residual autocorrelation? This is often appropriate for linear models (family = gaussian). Default is FALSE to be consistent with `wbm()` reduce runtime, and avoid warnings for non-linear models.
- **family**: Use this to specify GLM link families. Default is gaussian, the linear model.
- **fit_model**: Fit the model? Default is TRUE. If FALSE, only the model code is returned.
- **balance.correction**: Correct between-subject effects for unbalanced panels following the procedure in Curran and Bauer (2011)? Default is FALSE.
Should the detrending procedure be performed with a random slope for each entity? Default is TRUE but for short panels FALSE may be better, fitting a trend for all entities.

If detrending using detrend, what order polynomial would you like to specify for the relationship between time and the predictors? Default is 1, a linear model.

How many Markov chains should be used? Default is 3, to leave you with one unused thread if you’re on a typical dual-core machine.

How many iterations, including warmup? Default is 2000, leaving 1000 per chain after warmup. For some models and data, you may need quite a few more.

Standardize predictors? This can speed up model fit. Default is FALSE.

Save random effect estimates? This can be crucial for predicting from the model and for certain post-estimation procedures. On the other hand, it drastically increases the size of the resulting model. Default is FALSE.

The best way to calculate interactions in within models is in some dispute. The conventional way ("demean") is to first calculate the product of the variables involved in the interaction before those variables have their means subtracted and then subtract the mean of the product from the product term (see Schunk and Perales (2017)). Giesselmann and Schmidt-Catran (2020) show this method carries between-entity differences that within models are designed to model out. They suggest an alternate method ("double-demean") in which the product term is first calculated using the de-meaned lower-order variables and then the subject means are subtracted from this product term. Another option is to simply use the product term of the de-meaned variables ("raw"), but Giesselmann and Schmidt-Catran (2020) show this method biases the results towards zero effect. The default is "double-demean" but if emulating other software is the goal, "demean" might be preferred.

If using weights, either the name of the column in the data that contains the weights or a vector of the weights.

this can be used to specify an a priori known component to be included in the linear predictor during fitting. This should be NULL or a numeric vector of length equal to the number of cases. One or more offset terms can be included in the formula instead or as well, and if more than one is specified their sum is used. See model.offset.

Additional arguments passed on to brms::brm(). This can include specification of priors.

See wbm() for details on the formula syntax, model types, and some other stuff.

A wbm_stan object, which is a list containing a model object with the brm model and a stan_code object with the model code.

If fit_model = FALSE, instead a list is returned containing a stan_code object and a stan_data object, leaving you with the tools you need to run the model yourself using rstan.
**widen_panel**

**Author(s)**
Jacob A. Long

**See Also**
wbm()

**Examples**

```r
## Not run:
data("WageData")
wages <- panel_data(WageData, id = id, wave = t)
model <- wbm_stan(lwage ~ lag(union) + wks | blk + fem | blk * lag(union),
        data = wages, chains = 1, iter = 2000)
summary(model)
## End(Not run)
```

**widen_panel**

*Convert long panel data to wide format*

**Description**

This function takes `panel_data()` objects as input as converts them to wide format for use in SEM and other situations when such a format is needed.

**Usage**

```r
widen_panel(data, separator = ",", ignore.attributes = FALSE, varying = NULL)
```

**Arguments**

- **data**
  The `panel_data` frame.

- **separator**
  When the variables are labeled with the wave number, what should separate the variable name and wave number? By default, it is ",". In other words, a variable named `var` will be `var_1`, `var_2`, and so on in the wide data frame.

- **ignore.attributes**
  If the data was created by `long_panel()`, it stores information about which variables vary over time and which are constants. Sometimes, though, this information is not accurate (it is only based on the wide data’s variable names) and you may want to force this function to check again based on the actual values of the variables.

- **varying**
  If you want to skip the checks for whether variables are varying and specify yourself, as is done with `stats::reshape()`, you can supply them as a vector here.
Details

This is a wrapper for `stats::reshape()`, which is renowned for being pretty confusing to use. This function automatically detects which of the variables vary over time and which don’t, not appending wave information to constants.

Value

A data.frame with 1 row per respondent.

See Also

`reshape`

Examples

```r
wages <- panel_data(WageData, id = id, wave = t)
wide_wages <- widen_panel(wages)
```
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