

# Package ‘basecamb’

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**Type** Package

**Title** Utilities for Streamlined Data Import, Imputation and Modelling

**Version** 1.1.2

**Description** Provides functions streamlining the data analysis workflow:  
Outsourcing data import, renaming and type casting to a \*.csv.  
Manipulating imputed datasets and fitting models on them. Summarizing models.

**Depends** R (>= 4.0.0)

**Imports** assertive.types, assertthat, dplyr, mice, Hmisc, survival,  
stats, purrr

**License** GPL (>= 3)

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**Config/testthat/edition** 3

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<https://github.com/codeblue-team/basecamb>

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**NeedsCompilation** no

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<code>.scale_variable</code>	<i>Scaling a variable</i>
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---

## Description

A helper function to scale a variable in a dataframe. Divides 'variable' by 'scaling\_denominator'.

## Usage

```
.scale_variable(data, variable, scaling_denominator)
```

## Arguments

<code>data</code>	data.frame
<code>variable</code>	a char indicating the variable to be scaled
<code>scaling_denominator</code>	a numeric indicating the scaling. The variable is divided by the <code>scaling_denominator</code> .

## Value

the input dataframe with the newly scaled 'variable'

---

apply\_data\_dictionary *Clean column names, types and levels*

---

## Description

Use a data dictionary data.frame to apply the following tidying steps to your data.frame:

- Remove superfluous columns
- Rename columns
- Ensure/coerce correct data type for each column
- Assign factorial levels, including renaming and grouping

## Usage

```
apply_data_dictionary(  
  data,  
  data_dictionary,  
  na_action_default = "keep_NA",  
  print_coerced_NA = TRUE  
)
```

## Arguments

data data.frame to be cleaned

data\_dictionary

data.frame with the following columns:

- old\_column\_name : character with the old column name
- new\_data\_type : character denoting the tidy data type. Supported types are:
  - character
  - integer
  - float
  - factor
  - date
- new\_column\_name : tidy column name. Can be left blank to keep the old column name
- coding (factor and date columns only):
  - factor columns: character denoting old value (key) and new value (value) in a standardised fashion:
    - \* key-value pairs are separated from other key-value-pairs by a comma (",")
    - \* key and value of the same pair are separated by an equal sign ("=")
    - \* quotations around individual keys and values are recommended for clarity, but do not affect functionality.

- \* all values will be coerced to type character, with the exception of "NA" being parsed as type NA
- \* using "default" as a key will assign the specified value to all current values that do not match any of the specified keys, excluding NA
- \* using "NA" as a key will assign the specified value to all current NA values
- \* example coding: "'key1' = 'val1', 'key2' = 'val2', 'default' = 'Other', 'NA' = NA"
- \* if no coding is specified for a column, the coding remains unchanged
  - date columns: character denoting coding (see format argument in as.Date)
- Optional other columns (do not affect behaviour)

na\_action\_default

character: Specify what to do with NA values. Defaults to 'keep\_NA'. Options are:

- 'keep\_NA' NA values remain NA values
- 'assign\_default' NA values are assigned the value specified as 'default'. Requires a 'default' value to be specified Can be overwritten for individual columns by specifying a value for key 'NA'

print\_coerced\_NA

logical indicating whether a message specifying the location of NAs that are introduced by apply\_data\_dictionary() to data should be printed.

## Value

clean data.frame

## Author(s)

J. Peter Marquardt

---

apply\_function\_to\_imputed\_data

*Apply function to dataframes in a mice object*

---

## Description

Wrapper function to apply a function on each dataframe in an imputed dataset created with `mice::mice()`.

## Usage

`apply_function_to_imputed_data(mice_data, fun, ...)`

**Arguments**

mice_data	a mice object generated by <code>mice::mice()</code> .
fun	the function to apply to each dataframe. May only take one positional argument of type <code>data.frame</code> .
...	other arguments passed to <code>fun()</code>

**Value**

a mice object with transformed data.

**Author(s)**

J. Peter Marquardt

---

assign\_factorial\_levels

*Assign custom values for key levels in factorial columns*

---

**Description**

Use a named vector of keys (current value) and values for factorial columns to assign meaningful levels and/or group levels

**Usage**

```
assign_factorial_levels(
  data,
  factor_keys_values,
  na_action_default = "keep_NA"
)
```

**Arguments**

data            data.frame to modify

factor\_keys\_values

named list with:

- Keys: Names of factor columns
- values: Named vectors with
  - keys: current value (string representation)
  - values: new value to be assigned
  - if a 'default' key is passed, all existing values not conforming to the new scheme will be converted to the 'default' value
  - if a 'NA' key is passed, all NA values will be converted to the value specified here. Overwrites `na_action_default` for the specified column.

na\_action\_default

character: Specify what to do with NA values. Defaults to 'keep\_NA'. Options are:

- 'keep\_NA' NA values remain NA values
- 'assign\_default' NA values are assigned the value specified as 'default'. Requires a 'default' value to be specified Can be overwritten for individual columns by specifying a value for key 'NA'

### Value

data frame with new levels

### Author(s)

J. Peter Marquardt

### Examples

```
data <- data.frame(col1 = as.factor(rep(c('1', '2', '4'), 5)))
keys_1 <- list('col1' = c('1' = 'One', '2' = 'Two', '4' = 'Four'))
data_1 <- assign_factorial_levels(data, keys_1)
keys_2 <- list('col1' = c('1' = 'One', 'default' = 'Not_One'))
data_2 <- assign_factorial_levels(data, keys_2)
```

---

assign\_types\_names      *Assign tidy types and names to a data.frame*

---

### Description

Verbosely assign tidy name and data type for each column of a data.frame and get rid of superfluous columns. Uses a .csv file for assignments to encourage a data dictionary based workflow. CAVE! Requires 'Date' type columns to already be read in as Date.

### Usage

```
assign_types_names(data, meta_data)
```

### Arguments

data	data.frame to be tidied. Dates must already be of type date.
meta_data	data.frame specifying old column names, new column names and datatypes of data. Has the following columns: <ul style="list-style-type: none"> <li>• old_column_name : character with the old column name.</li> <li>• new_data_type : character denoting the tidy data type. Supported types are: <ul style="list-style-type: none"> <li>– character (will be coerced using <code>as.character()</code>).</li> <li>– integer (will be coerced using <code>as.integer()</code>).</li> </ul> </li> </ul>

- float (will be coerced using `as.double()`).
- factor (will be coerced using `as.factor()`). Will result in a warning if the new factor variable will have more than 10 levels.
- date (can only confirm correct datatype assignment or coerce characters with format `'%Y-%m-%d'`).
- `new_column_name` : tidy column name. Can be left blank to keep the old column name.
- Optional other columns (do not affect behavior).

**Value**

clean data.frame

**Author(s)**

J. Peter Marquardt

---

build\_model\_formula    *Build formula for statistical models*

---

**Description**

Build formula used in statistical models from vectors of strings with the option to specify an environment.

**Usage**

```
build_model_formula(  
  outcome,  
  predictors,  
  censor_event = NULL,  
  env = parent.frame()  
)
```

**Arguments**

<code>outcome</code>	character denoting the column with the outcome.
<code>predictors</code>	vector of characters denoting the columns with the predictors.
<code>censor_event</code>	character denoting the column with the censoring event, for use in Survival-type models.
<code>env</code>	environment to be used in formula creation

**Value**

formula for use in statistical models

**Author(s)**

J. Peter Marquardt

**Examples**

```
build_model_formula("outcome", c("pred_1", "pred_2"))
build_model_formula("outcome", c("pred_1", "pred_2"), censor_event = "cens_event")
```

---

cox.zph.mids	<i>Test cox proportional odds assumption on models using multiple imputation.</i>
--------------	---

---

**Description**

Constructs a model and conducts a cox.zph test for each imputation of the data set.

**Usage**

```
cox.zph.mids(
  model,
  imputations,
  p_level = 0.05,
  global_only = TRUE,
  return_raw = FALSE,
  p_only = TRUE,
  verbose = TRUE
)
```

**Arguments**

model	cox proportional model to be evaluated
imputations	mids object containing imputations
p_level	value below which violation of proportional odds assumption is assumed. Defaults to .05
global_only	return global p-value only. Implies p_only to be TRUE
return_raw	return cox.zph objects in a list. If TRUE, function will not return anything else
p_only	returns p-values of test only. If FALSE returns Chi <sup>2</sup> and degrees of freedom as well
verbose	Set to FALSE to deactivate messages

**Value**

depending on specified options, this function can return

- default: A vector of global p-values
- global\_only = FALSE: a data.frame with p-values for all variables plus the global
- return\_raw = TRUE: list of cox.zph objects



**Author(s)**

J. Peter Marquardt

**Examples**

```
data <- data.frame(time = 101:200, status = rep(c(0,1), 50), pred = rep(c(1:9, NA), 10))
imputed_data <- mice::mice(data)
cox_mod <- Hmisc::fit.mult.impute(survival::Surv(time, status) ~ pred,
fitter = rms::cph, xtrans = imputed_data)
cox.zph.mids(cox_mod, imputed_data)
```

---

deconstruct\_formula     *Deconstruct formula*

---

**Description**

Deconstruct a formula object into strings of its components. Predictors are split by '+', so interaction terms will be returned as a single string.

**Usage**

```
deconstruct_formula(formula)
```

**Arguments**

formula            formula object for use in statistical models.

**Value**

a named list with fields:

- outcome (character)
- predictors (vector of characters)
- censor\_event (character) (optional) censor event, only for formulas including a Surv() object

**Author(s)**

J. Peter Marquardt

**Examples**

```
deconstruct_formula(stats::as.formula("outcome ~ predictor1 + predictor2 + predictor3"))
deconstruct_formula(stats::as.formula("Surv(outcome, censor_event) ~ predictor"))
```

---

filter_nth_entry	<i>Filter dataframe for nth entry</i>
------------------	---------------------------------------

---

### Description

Filter a dataframe for the nth entry of each subject in it. A typical use cases would be to filter a dataset for the first or last measurement of a subject.#'

### Usage

```
filter_nth_entry(data, ID_column, entry_column, n = 1, reverse_order = FALSE)
```

### Arguments

data	the data.frame to filter
ID_column	character column identifying subjects
entry_column	character column identifying order of entries. That column can be of types Date, numeric, or any other type suitable for order()
n	integer number of entry to keep after ordering
reverse_order	logical when TRUE sorts entries last to first before filtering

### Value

data.frame with  $\leq 1$  entry per subject

### Author(s)

J. Peter Marquardt

### Examples

```
data <- data.frame(list(ID = rep(1:5, 3), encounter = rep(1:3, each=5), value = rep(4:6, each=5)))
filter_nth_entry(data, 'ID', 'encounter')
filter_nth_entry(data, 'ID', 'encounter', n = 2)
filter_nth_entry(data, 'ID', 'encounter', reverse_order = TRUE)
```

---

`fit_mult_impute_obs_outcome`

*Fit a model on multiply imputed data using only observations with non-missing outcome(s)*

---

### Description

This function is a wrapper for fitting models with `Hmisc::fit.mult.impute()` on a multiply imputed dataset generated with `mice::mice()`. Cases with a missing outcome in the original dataset are removed from the `mids` object by using the "subset" argument in `Hmisc::fit.mult.impute()`.

### Usage

```
fit_mult_impute_obs_outcome(mids, formula, fitter, ...)
```

### Arguments

<code>mids</code>	a <code>mids</code> object, i.e. the imputed dataset.
<code>formula</code>	a formula that describes the model to be fit. The outcome (y variable) in the formula will be used to remove missing cases.
<code>fitter</code>	a modeling function (not in quotes) that is compatible with <code>Hmisc::fit.mult.impute()</code> .
<code>...</code>	additional arguments to <code>Hmisc::fit.mult.impute()</code> .

### Value

mod a `fit.mult.impute` object.

### Author(s)

Till D. Best

### Examples

```
# create an imputed dataset
imputed_data <- mice::mice(airquality)

fit_mult_impute_obs_outcome(mids = imputed_data, formula = Ozone ~ Solar.R + Wind, fitter = glm)
```

---

`or_model_summary`*Summarise a logistic regression model on the odds ratio scale*

---

### Description

This function summarises regression models that return data on the log-odds scale and returns a dataframe with estimates, and confidence intervals as odds ratios. P value are also provided. Additionally, intercepts can be removed from the summary. This comes in handy when ordinal logistic regression models are fit. Ordinal regression models (such as proportional odds models) usually result in many intercepts that are not really of interest. This function is also compatible with models obtained from multiply imputed datasets, for example models fitted with `Hmisc::fit.mult.impute()`.

### Usage

```
or_model_summary(  
  model,  
  conf_int = 1.96,  
  print_intercept = FALSE,  
  round_est = 3,  
  round_p = 4  
)
```

### Arguments

<code>model</code>	a model object with estimates on the log-odds scale.
<code>conf_int</code>	a numeric used to calculate the confidence intervals. The default of 1.96 gives the 95% confidence interval.
<code>print_intercept</code>	a logical flag indicating whether intercepts shall be removed. All variables that start with "y>=" will be removed. If there is a variable matching this pattern, it will also be removed!
<code>round_est</code>	the number of decimals returned for estimates (odds ratios) and confidence intervals.
<code>round_p</code>	the number of decimals provided for p-values.

### Details

CAVE! The function does not check whether your estimates are on the log-odds scale. It will do the transformation no matter what!

### Value

a dataframe with the adjusted odds ratio, confidence intervals and p-values.

### Author(s)

Till D. Best

**Examples**

```
# fit a logistic model
mod <- glm(formula = am ~ mpg + cyl, data = mtcars, family = binomial())

or_model_summary(model = mod)
```

---

parse\_date\_columns      *Parse values in date columns as Dates*

---

**Description**

Parse date columns in a data.frame as Date. Use a named list to specify each date column (key) and the format (value) it is coded in.

**Usage**

```
parse_date_columns(data, date_formats)
```

**Arguments**

data	data.frame to modify
date_formats	named list with: <ul style="list-style-type: none"><li>• Keys: Names of date columns</li><li>• values: character specifying the format</li></ul>

**Value**

data.frame with date columns in Date type

**Author(s)**

J. Peter Marquardt

**Examples**

```
data <- data.frame(date = rep('01/23/4567', 5))
data <- parse_date_columns(data, list(date = '%m/%d/%Y'))
```

---

quantile\_group      *Stratify a numeric vector into quantile groups*

---

### Description

Transforms a numeric vector into quantile groups. For each input value, the output value corresponds to the quantile that value is in. When grouping into  $n$  quantiles, the lowest  $1/n$  of values are assigned 1, the highest  $1/n$  are assigned  $n$ .

### Usage

```
quantile_group(data, n, na.rm = TRUE)
```

### Arguments

data	a vector of type numeric with values to be grouped into quantiles
n	integer indicating number of quantiles, minimum of 2. Must be smaller than <code>length(data)</code>
na.rm	logical; if TRUE all NA values will be removed before calculating groups, if FALSE no NA values are permitted.

### Details

Tied values will be assigned to the lower quantile group rather than estimating a distribution. In extreme cases this can mean one or more quantile groups are not represented.

If uneven group sizes cannot be avoided, values will be assigned the higher quantile group.

### Value

vector of length `length(data)` with the quantile groups

### Author(s)

J. Peter Marquardt

### Examples

```
quantile_group(10:1, 3)
quantile_group(c(rep(1,3), 10:1, NA), 5)
```

---

`remove_missing_from_mids`*Remove missing cases from a mids object*

---

## Description

Deprecated, use [apply\\_function\\_to\\_imputed\\_data](#) instead.

## Usage

```
remove_missing_from_mids(mids, var)
```

## Arguments

<code>mids</code>	mids objects that is filtered.
<code>var</code>	a string or vector of strings specifying the variable(s). All cases (i.e. rows) for which there are missing values are removed.

## Details

`Remove_missing_from_mids` is used to filter a mids object for missing cases in the original dataset in the variable `var`. This is useful for situations where you want to use as many observations as possible for imputation but only fit your model on a subset of these. Or, if you want to create one large imputed dataset from which multiple analyses with multiple outcomes are derived.

## Value

a mids object filtered for observed cases of `var`.

## Author(s)

Till D. Best

## See Also

[apply\\_function\\_to\\_imputed\\_data](#)

---

scale\_continuous\_predictors  
*Scale continuous predictors*

---

**Description**

This function linearly scales variables in data objects according to a data dictionary. The data dictionary has at least two columns, "variable" and "scaling\_denominator". "Variable" is divided by "scaling\_denominator".

**Usage**

```
scale_continuous_predictors(data, scaling_dictionary)
```

**Arguments**

`data` a data object with variables.  
`scaling_dictionary` a data.frame with two columns that are called "variable" and "scaling\_denominator".

**Value**

The data with the newly scaled 'variables'.

**Author(s)**

Till D. Best



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