

# Package ‘codebook’

November 23, 2018

**Title** Automatic Codebooks from Survey Metadata Encoded in Attributes

**Description** Easily automate the following tasks to describe data frames:  
summarise the distributions of scales and items graphically and using descriptive statistics,  
compute reliabilities (internal consistencies, retest, multilevel) for psychological scales,  
combine this information with metadata (such as item labels and labelled values) that is derived from R attributes.  
To do so, the package relies on 'rmarkdown' partials, so you can generate HTML, PDF, and Word documents. Codebooks are also available as tables (CSV, Excel, etc.).  
The metadata and codebooks are also available at your fingertips via RStudio Addins.

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**Depends** R (>= 3.0.1)

**URL** <https://github.com/rubenarслан/codebook>

**BugReports** <https://github.com/rubenarслан/codebook/issues>

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---

aggregate\_and\_document\_scale

*Aggregate variables and remember which variables this were*

---

## Description

The resulting variables will have the attribute `scale_item_names` containing the basis for aggregation. Its `label` attribute will refer to the common stem of the aggregated variable names (if any), the number of variables, and the aggregation function.

## Usage

```
aggregate_and_document_scale(items, fun = rowMeans, stem = NULL)
```

## Arguments

<code>items</code>	data.frame of the items that should be aggregated
<code>fun</code>	aggregation function, defaults to <code>rowMeans</code> with <code>na.rm = FALSE</code>
<code>stem</code>	common stem for the variables, specify if it should not be auto-detected as the longest common stem of the variable names

## Examples

```
testdf <- data.frame(bfi_neuro_1 = rnorm(20), bfi_neuro_2 = rnorm(20),
                     bfi_neuro_3R = rnorm(20), age = rpois(20, 30))
item_names <- c('bfi_neuro_1', 'bfi_neuro_2', 'bfi_neuro_3R')
testdf$bfi_neuro <- aggregate_and_document_scale(testdf[, item_names])
testdf$bfi_neuro
```

---

<code>asis_knit_child</code>	<i>Knit a child document and output as is (render markup)</i>
------------------------------	---

---

## Description

This slightly modifies the `knitr::knit_child()` function to have different defaults.

- the environment defaults to the calling environment.
- the output receives the class `knit_asis`, so that the output will be rendered "as is" by knitr when calling inside a chunk (no need to set `results='asis'` as a chunk option).
- defaults to `quiet = TRUE`

## Usage

```
asis_knit_child(input = NULL, text = NULL, ..., quiet = TRUE,
               options = NULL, envir = parent.frame(), use_strings = TRUE)
```

## Arguments

<code>input</code>	if you specify a file path here, it will be read in before being passed to knitr (to avoid a working directory mess)
<code>text</code>	passed to <code>knitr::knit_child()</code>
<code>...</code>	passed to <code>knitr::knit_child()</code>
<code>quiet</code>	passed to <code>knitr::knit_child()</code>
<code>options</code>	defaults to <code>NULL</code> .
<code>envir</code>	passed to <code>knitr::knit_child()</code>
<code>use_strings</code>	whether to read in the child file as a character string (solves working directory problems but harder to debug)

## Details

Why default to the calling environment? Typically this function defaults to the global environment. This makes sense if you want to use knit children in the same context as the rest of the document. However, you may also want to use knit children inside functions to e.g. summarise a regression using a set of commands (e.g. plot some diagnostic graphs and a summary for a regression nicely formatted).

Some caveats:

- the function has to return to the top-level. There's no way to `cat()` this from loops or an if-condition without setting `results='asis'`. You can however concatenate these objects with `paste.knit_asis()`

## Examples

```
## Not run:
# an example of a wrapper function that calls asis_knit_child with an argument
# ensures distinct paths for cache and figures, so that these calls can be looped in para
regression_summary <- function(model) {
  hash <- digest::digest(model)
  options <- list(
    fig.path = paste0(knitr::opts_chunk$get("fig.path"), hash, "-"),
    cache.path = paste0(knitr::opts_chunk$get("cache.path"), hash, "-")
  )
  asis_knit_child("_regression_summary.Rmd", options = options)
}

## End(Not run)
```

---

bfi

---

*Mock BFI data*


---

## Description

a small mock BFI dataset with realistic values, exported from formr. The dataset is self-documenting via its attributes.

## Usage

```
bfi
```

## Format

A data frame with 28 rows and 29 variables:

## Description

Pass a data frame to this function to make a codebook for that dataset. If the dataset has metadata (attributes) set on its variables, these will be used to make the codebook more informative. Examples are item, value, and missing labels. Data frames imported via `haven::read_dta()`, `haven::read_sav()`, or from [formr.org](https://formr.org) will have these attributes in the right format. By calling this function inside a knitr code chunk, the codebook will become part of the document you are generating.

## Usage

```
codebook(results, reliabilities = NULL, survey_repetition = c("auto",
  "single", "repeated_once", "repeated_many"), survey_overview = TRUE,
  missingness_report = TRUE, metadata_table = TRUE,
  metadata_json = TRUE, indent = "#")
```

## Arguments

<code>results</code>	a data frame, ideally with attributes set on variables
<code>reliabilities</code>	a named list with one entry per scale and one or several printable reliability computations for this scale. if NULL, computed on-the-fly using <code>compute_reliabilities</code>
<code>survey_repetition</code>	defaults to "auto" which is to try to determine the level of repetition from the "session" and "created" variables. Other values are: single, repeated_once, repeated_many
<code>survey_overview</code>	whether to print an overview of survey entries, durations (depends on presence of columns session, created, modified, ended, expired)
<code>missingness_report</code>	whether to print a missingness report. Turn off if this gets too complicated and you need a custom solution (e.g. in case of random missings).
<code>metadata_table</code>	whether to print a metadata table/tabular codebook.
<code>metadata_json</code>	whether to include machine-readable metadata as JSON-LD (not visible)
<code>indent</code>	add # to this to make the headings in the components lower-level. defaults to beginning at h2

## Examples

```
# will generate figures in a temporary directory
old_base_dir <- knitr::opts_knit$get("base.dir")
knitr::opts_knit$set(base.dir = tempdir())
on.exit(knitr::opts_knit$set(base.dir = old_base_dir))
data("bfi")
bfi <- bfi[, c("BFIK_open_1", "BFIK_open_1")]
md <- codebook(bfi, survey_repetition = "single", metadata_table = FALSE)
```

---

codebook\_browser      *Browse and search codebook*


---

### Description

Usable as an Addin in RStudio. You can select it from a menu at the top, when this package is installed. If you're currently selecting the name of a data frame in your source code, this will be the dataset shown by default. If you don't select text, you can pick a dataset from a dropdown. You can add a keyboard shortcut for this command by following the [instructions](#) by RStudio. How about Cmd+Ctrl+C?

### Usage

```
codebook_browser(data = NULL, labels_only = FALSE,
  title = "Codebook metadata", viewer = rstudioapi::viewer)
```

### Arguments

data	the dataset to display. If left empty will try to use selected text in RStudio or offer a dropdown
labels_only	defaults to false called with TRUE from <code>label_browser()</code>
title	title of the gadget
viewer	defaults to displaying in the RStudio viewer

---

codebook\_component\_scale  
*Codebook component for scales*


---

### Description

Codebook component for scales

### Usage

```
codebook_component_scale(scale, scale_name, items, reliabilities,
  indent = "##")
```

### Arguments

scale	a scale with attributes set
scale_name	the variable name of this scale
items	a data.frame with the items constituting the scale
reliabilities	a list with one or several results from calls to psych package functions for computing reliability
indent	add # to this to make the headings in the components lower-level. defaults to beginning at h2

**Examples**

```
# will generate figures in a temporary directory
old_base_dir <- knitr::opts_knit$get("base.dir")
knitr::opts_knit$set(base.dir = tempdir())
on.exit(knitr::opts_knit$set(base.dir = old_base_dir))
data("bfi")
bfi <- bfi[,c("BFIK_open", paste0("BFIK_open_", 1:4))]
codebook_component_scale(bfi[,1], "BFIK_open", bfi[, -1],
  reliabilities = list(BFIK_open = psych::alpha(bfi[, -1])))
```

---

codebook\_component\_single\_item

*Codebook component for single items*


---

**Description**

Codebook component for single items

**Usage**

```
codebook_component_single_item(item, item_name, indent = "##")
```

**Arguments**

item	an item with attributes set
item_name	the item name
indent	add # to this to make the headings in the components lower-level. defaults to beginning at h2

**Examples**

```
# will generate figure in a temporary directory
old_base_dir <- knitr::opts_knit$get("base.dir")
knitr::opts_knit$set(base.dir = tempdir())
on.exit(knitr::opts_knit$set(base.dir = old_base_dir))
data("bfi")
codebook_component_single_item(bfi$BFIK_open_1, "BFIK_open_1")
```

---

codebook\_items

*Tabular codebook*


---

**Description**

Renders a tabular codebook including attributes and data summaries. The table is generated using `DT::datatable()` and can be exported to CSV, Excel, etc.

**Usage**

```
codebook_items(results, indent = "##")
```

**Arguments**

<code>results</code>	a data frame, ideally with attributes set on variables
<code>indent</code>	add # to this to make the headings in the components lower-level. defaults to beginning at h2

**Examples**

```
data("bfi")
## Not run:
# doesn't show interactively, because a html widget needs to be registered
codebook_items(bfi)

## End(Not run)
```

---

```
codebook_missingness
      Codebook missingness
```

---

**Description**

An overview table of missingness patterns generated using `md_pattern()`.

**Usage**

```
codebook_missingness(results, indent = "##")
```

**Arguments**

<code>results</code>	a data frame
<code>indent</code>	add # to this to make the headings in the components lower-level. defaults to beginning at h2

**Examples**

```
data("bfi")
codebook_missingness(bfi)
```

---

```
codebook_survey_overview
      Codebook survey overview
```

---

**Description**

An overview of the number of rows and groups, and of the durations participants needed to respond (if those data are available).

**Usage**

```
codebook_survey_overview(results, survey_repetition = "single",
  indent = "##")
```



**Arguments**

`results` a data frame which has the following columns: session, created, modified, expired, ended

`survey_repetition` defaults to single (other values: repeated\_once, repeated\_many). controls whether internal consistency, retest reliability or multilevel reliability is computed

`indent` add # to this to make the headings in the components lower-level. defaults to beginning at h2

**Examples**

```
# will generate figures in a figure/ subdirectory
data("bfi")
codebook_survey_overview(bfi)
```

---

codebook_table	<i>Codebook metadata table</i>
----------------	--------------------------------

---

**Description**

will generate a table combining metadata from variable attributes with data summaries generated using `skimr::skim_to_wide()`

**Usage**

```
codebook_table(results)
```

**Arguments**

`results` a data frame, ideally with attributes set on variables

**Examples**

```
data("bfi")
codebook_table(bfi)
```

---

compute_reliabilities	<i>Compute reliabilities</i>
-----------------------	------------------------------

---

**Description**

If you pass the object resulting from a call to `formr_results` to this function, it will compute reliabilities for each scale. Internally, each reliability computation is passed to a `future::future()`. If you are calculating multilevel reliabilities, it may be worthwhile to parallelise this operation using `future::plan()`. If you don't plan on any complicated parallelisation, you probably do not need to call this function directly, but can rely on it being automatically called during codebook generation. If you do plan to do that, you can pass the results of this operation to the codebook function.

**Usage**

```
compute_reliabilities(results, survey_repetition = "single")
```

**Arguments**

`results` a formr results table with attributes set on items and scales  
`survey_repetition` defaults to "single". Can also be "repeated\_once" or "repeated\_many"

**Examples**

```
data("bfi", package = "codebook")
reliabilities <- compute_reliabilities(bfi)
```

---

<code>detect_missings</code>	<i>Detect missing values</i>
------------------------------	------------------------------

---

**Description**

SPSS users frequently label their missing values, but don't set them as missing. This function will rectify that for negative values and for the values 99 and 999 (only if they're 5\*MAD away from the median). Using different settings, you can also easily tag other missings.

**Usage**

```
detect_missings(data, only_labelled_missings = TRUE,
  negative_values_are_missing = TRUE, ninety_nine_problems = TRUE,
  learn_from_labels = TRUE, missing = c(), non_missing = c(),
  vars = names(data), use_labelled_spss = FALSE)
```

**Arguments**

`data` the data frame with labelled missings  
`only_labelled_missings` don't set values to missing if there's no label for them  
`negative_values_are_missing` by default we label negative values as missing  
`ninety_nine_problems` SPSS users often store values as 99/999, should we do this for values with 5\*MAD of the median  
`learn_from_labels` if there are labels for missings of the form [-1] no answer, set -1 in the data to the corresponding tagged missing  
`missing` also set these values to missing (or enforce for 99/999 within 5\*MAD)  
`non_missing` don't set these values to missing  
`vars` only edit these variables  
`use_labelled_spss` the labelled\_spss class has a few drawbacks. Since R can't store missings like -1 and 99, we're replacing them with letters unless this option is enabled. If you prefer to keep your -1 etc, turn this on.

---

detect_scales	<i>Detect item scales</i>
---------------	---------------------------

---

### Description

Did you create aggregates of items like this `scale <- scale_1 + scale_2R + scale_3R`? If you run this function on a dataset, it will detect these relationships and set the appropriate attributes. Once they are set, the codebook package can perform reliability computations for you.

### Usage

```
detect_scales(data, quiet = FALSE)
```

### Arguments

<code>data</code>	the data frame
<code>quiet</code>	defaults to false. Suppresses messages about found items.

### Examples

```
bfi <- data.frame(matrix(data = rnorm(500), ncol = 5))
names(bfi) <- c("bfi_e1", "bfi_e2R", "bfi_e3", "bfi_n1", "bfi_n2")
bfi$bfi_e <- rowMeans(bfi[, c("bfi_e1", "bfi_e2R", "bfi_e3")])
bfi <- detect_scales(bfi)
bfi$bfi_e
```

---

ended	<i>How many surveys were ended?</i>
-------	-------------------------------------

---

### Description

Just a simple to check how many times a survey (e.g. diary) was finished. It defaults to checking the "ended" variable for this.

### Usage

```
ended(survey, variable = "ended")
```

### Arguments

<code>survey</code>	which survey are you asking about?
<code>variable</code>	which variable should be filled out, defaults to "ended"

### Examples

```
survey <- data.frame(ended = c("2016-05-28 10:11:00", NA, "2016-05-30 11:18:28"))
ended(survey = survey)
```

---

 expired

*How many surveys were expired?*


---

### Description

Just a simple to check how many times a survey (e.g. diary) has expired (i.e. user missed it). It defaults to checking the "expired" variable for this.

### Usage

```
expired(survey, variable = "expired")
```

### Arguments

survey	which survey are you asking about?
variable	which variable should be filled out, defaults to "expired"

### Examples

```
survey <- data.frame(expired = c(NA, "2016-05-29 10:11:00", NA))
expired(survey = survey)
```

---

 has\_label

*Has label*


---

### Description

Has label

### Usage

```
has_label(x)
```

### Arguments

x	a vector
---	----------

### Examples

```
example("labelled", "haven")
has_label(x)
```

---

has_labels	<i>Has labels</i>
------------	-------------------

---

**Description**

Has labels

**Usage**

```
has_labels(x)
```

**Arguments**

x	a vector
---	----------

**Examples**

```
example("labelled", "haven")
has_labels(x)
```

---

knitr::knit_print.alpha	<i>Pretty-print a Cronbach's alpha object</i>
-------------------------	---

---

**Description**

Turn a `psych::alpha()` object into HTML tables.

**Usage**

```
knit_print.alpha(x, indent = "#####", ...)
```

**Arguments**

x	a psych alpha object
indent	add # to this to make the headings in the components lower-level. defaults to beginning at h5
...	ignored

**Examples**

```
example("alpha", "psych")
knitr::knit_print(a4)
```

---

```
knit_print.htest
```

*Print a `stats::cor.test()` object for knitr*

---

### Description

Just prints the normal output of `stats::cor.test()`.

### Usage

```
knit_print.htest(x, indent = "#####", ...)
```

### Arguments

<code>x</code>	a psych alpha object
<code>indent</code>	add # to this to make the headings in the components lower-level. defaults to beginning at h5
<code>...</code>	ignored

### Examples

```
knitr::knit_print(cor.test(rnorm(100), rnorm(100)))
```

---

```
knit_print.multilevel
```

*Print a `psych::multilevel.reliability()` object for knitr*

---

### Description

Just prints the normal output of `psych::multilevel.reliability()`.

### Usage

```
knit_print.multilevel(x, indent = "#####", ...)
```

### Arguments

<code>x</code>	a psych alpha object
<code>indent</code>	add # to this to make the headings in the components lower-level. defaults to beginning at h5
<code>...</code>	ignored

### Examples

```
example("mlr", "psych")
knitr::knit_print(mg)
```

---

label_browser	<i>Browse and search variable and value labels</i>
---------------	--

---

### Description

Same as the `codebook_browser()`, but doesn't show data summaries and additional attributes.

### Usage

```
label_browser(data = NULL, viewer = rstudioapi::viewer)
```

### Arguments

data	the dataset to display. If left empty will try to use selected text in RStudio or offer a dropdown
viewer	defaults to displaying in the RStudio viewer

---

label_browser_static	<i>Browse and search variable and value labels</i>
----------------------	--

---

### Description

Same as the `codebook_browser()`, but doesn't show data summaries and additional attributes. This yields a static table, so you can continue to edit code while viewing the labels, but you cannot switch the dataset via a dropdown menu.

### Usage

```
label_browser_static(data = NULL, viewer = rstudioapi::viewer)
```

### Arguments

data	data frame. if left empty, will use the text you currently select in RStudio as the label or the first data frame in your environment
viewer	where to show. defaults to viewer tab

### Examples

```
label_browser_static(bfi)
```

---

```
likert_from_items
```

*Derive a likert object from items*


---

### Description

Pass a data.frame containing several items composing one scale, get a `likert::likert()` object, which you can plot. Intelligently makes use of labels and value labels if present.

### Usage

```
likert_from_items(items)
```

### Arguments

`items`                      a data frame of items composing one scale

### Examples

```
data("bfi", package = "codebook")
open_items <- paste0("BFIK_open_", 1:4)
graphics::plot(likert_from_items(bfi[, open_items]))
```

---

```
load_data_and_render_codebook
```

*Render codebook based on file*


---

### Description

Submit a data file and an rmarkdown template as a file to generate a codebook. Used chiefly in the webapp.

### Usage

```
load_data_and_render_codebook(file, text, remove_file = FALSE, ...)
```

### Arguments

`file`                      path to a file to make codebook from (sav, rds, dta, por, xpt, csv, csv2, tsv, etc.)

`text`                      codebook template

`remove_file`              whether to remove file after rendering

`...`                      all other arguments passed to `rmarkdown::render()`



---

md_pattern	<i>Missing data patterns</i>
------------	------------------------------

---

### Description

Generate missingness patterns using `mice::md.pattern()`, with options to reduce the complexity of the output.

### Usage

```
md_pattern(data, only_vars_with_missings = TRUE, min_freq = 0.01)
```

### Arguments

<code>data</code>	the dataset
<code>only_vars_with_missings</code>	defaults to TRUE, omitting variables that have no missings
<code>min_freq</code>	minimum number of rows to have this missingness pattern

### Examples

```
data("nhanes", package = "mice")
md_pattern(nhanes)
md_pattern(nhanes, only_vars_with_missings = FALSE, min_freq = 0.2)
```

---

metadata_jsonld	<i>Metadata as JSON-LD</i>
-----------------	----------------------------

---

### Description

Echo a list of a metadata, generated using `metadata_list()` as JSON-LD in a script tag.

### Usage

```
metadata_jsonld(results)
```

### Arguments

<code>results</code>	a data frame, ideally with attributes set on variables
----------------------	--

### Examples

```
data("bfi")
metadata_jsonld(bfi)
```

---

metadata_list	<i>Metadata from dataframe</i>
---------------	--------------------------------

---

### Description

Returns a list containing variable metadata (attributes) and data summaries.

### Usage

```
metadata_list(results)
```

### Arguments

`results` a data frame, ideally with attributes set on variables

### Examples

```
data("bfi")
md_list <- metadata_list(bfi)
md_list$variableMeasured[[20]]
```

---

modified	<i>How many surveys were modified?</i>
----------	--

---

### Description

Just a simple to check how many times a survey (e.g. diary) has expired (i.e. user missed it). It defaults to checking the "expired" variable for this.

### Usage

```
modified(survey, variable = "modified")
```

### Arguments

`survey` which survey are you asking about?  
`variable` which variable should be filled out, defaults to "modified"

### Examples

```
survey <- data.frame(modified = c(NA, "2016-05-29 10:11:00", NA))
modified(survey = survey)
```

---

paste.knit_asis	<i>Paste and output as is (render markup)</i>
-----------------	---

---

### Description

Helper function for knit\_asis objects, useful when e.g. `asis_knit_child()` was used in a loop.

### Usage

```
paste.knit_asis(..., sep = "\n\n\n", collapse = "\n\n\n")
```

### Arguments

<code>...</code>	passed to <code>base::paste()</code>
<code>sep</code>	defaults to two empty lines, passed to <code>base::paste()</code>
<code>collapse</code>	defaults to two empty lines, passed to <code>base::paste()</code>

### Details

Works like `base::paste()` with both the `sep` and the `collapse` argument set to two empty lines

### Examples

```
paste.knit_asis("# Headline 1", "## Headline 2")
```

---

plot_labelled	<i>Plot labelled vector</i>
---------------	-----------------------------

---

### Description

Plot a labelled vector, making use of the variable name, label and value labels to make the plot more readable. This function also works for other vectors, but provides little benefit.

### Usage

```
plot_labelled(item, item_name = NULL, wrap_at = 50,
  go_vertical = FALSE)
```

### Arguments

<code>item</code>	a vector
<code>item_name</code>	item name, defaults to name of first argument
<code>wrap_at</code>	the subtitle (the label) will be wrapped at this number of characters
<code>go_vertical</code>	defaults to FALSE. Whether to show choices on the Y axis instead.

### Examples

```
data("bfi", package = "codebook")
plot_labelled(bfi$BFIK_open_1)
```

---

<code>print.knit_asis</code>	<i>Print new lines in knit_asis outputs</i>
------------------------------	---

---

### Description

Print new lines in `knit_asis` outputs

### Usage

```
## S3 method for class 'knit_asis'  
print(x, ...)
```

### Arguments

<code>x</code>	the <code>knit_asis</code> object
<code>...</code>	ignored

---

<code>rescue_attributes</code>	<i>Rescue lost attributes</i>
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---

### Description

You can use this function if some of your items have lost their attributes during wrangling. Variables have to have the same name (Duh) and no attributes should be overwritten. But use with care. Similar to `labelled::copy_labels()`.

### Usage

```
rescue_attributes(df_no_attributes, df_with_attributes)
```

### Arguments

<code>df_no_attributes</code>	the data frame with missing attributes
<code>df_with_attributes</code>	the data frame from which you want to restore attributes

---

```
reverse_labelled_values
```

*Reverse labelled values reverse the underlying value for a numeric `haven::labelled()` vector while keeping the labels correct*

---

### Description

Reverse labelled values reverse the underlying value for a numeric `haven::labelled()` vector while keeping the labels correct

### Usage

```
reverse_labelled_values(x)
```

### Arguments

`x`                      a labelled vector

### Value

return the labelled vector with the underlying values having been reversed

### Examples

```
x <- haven::labelled(rep(1:3, each = 3), c(Bad = 1, Good = 5))
x
reverse_labelled_values(x)
```

---

```
summary.labelled      Summary function for labelled vector
```

---

### Description

Summary function for labelled vector

### Usage

```
## S3 method for class 'labelled'
summary(object, ...)
```

### Arguments

`object`                a labelled vector  
`...`                    passed to `summary.factor`

### Examples

```
example("labelled", "haven")
summary(x)
```

---

```
summary.labelled_spss
```

*Summary function for labelled\_spss vector*

---

### Description

Summary function for labelled\_spss vector

### Usage

```
## S3 method for class 'labelled_spss'
summary(object, ...)
```

### Arguments

<code>object</code>	a labelled_spss vector
<code>...</code>	passed to summary.factor

### Examples

```
example("labelled", "haven")
summary(x)
```

---

<code>zap_attributes</code>	<i>Zap attributes</i>
-----------------------------	-----------------------

---

### Description

Modelled on `haven::zap_labels()`, but more encompassing. By default removes the following attributes: `format.spss`, `format.sas`, `format.stata`, `label`, `labels`, `na_values`, `na_range`, `display_width`

### Usage

```
zap_attributes(x, attributes = c("format.spss", "format.sas",
  "format.stata", "label", "labels", "na_values", "na_range",
  "display_width"))
```

### Arguments

<code>x</code>	the data frame or variable
<code>attributes</code>	character vector of attributes to zap. NULL if everything (including factor levels etc) should be zapped

**Examples**

```
bfi <- data.frame(matrix(data = rnorm(300), ncol = 3))
names(bfi) <- c("bfi_e1", "bfi_e2R", "bfi_e3")
attributes(bfi$bfi_e1)$label <- "I am outgoing."
attributes(bfi$bfi_e2R)$label <- "I prefer books to people."
attributes(bfi$bfi_e3)$label <- "I love to party."
bfi$bfi_e <- rowMeans(bfi[, c("bfi_e1", "bfi_e2R", "bfi_e3")])
bfi <- detect_scales(bfi, quiet = TRUE) # create attributes
str(zap_attributes(bfi, "label"))
zap_attributes(bfi$bfi_e)
```

---

zap_label	<i>Zap variable label</i>
-----------	---------------------------

---

**Description**

Modelled on `haven::zap_labels()`, zaps variable labels (not value labels).

**Usage**

```
zap_label(x)
```

**Arguments**

`x` the data frame or variable

**Examples**

```
x <- haven::labelled(rep(1:5, each = 1), c(Bad = 1, Good = 5))
zap_label(x)
```

---

zap_labelled	<i>Zap labelled class</i>
--------------	---------------------------

---

**Description**

Modelled on `haven::zap_labels()`, zaps labelled class (not other attributes).

**Usage**

```
zap_labelled(x)
```

**Arguments**

`x` the data frame or variable

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