Package ‘janitor’

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Title Simple Tools for Examining and Cleaning Dirty Data
Version 1.2.1

Description The main janitor functions can: perfectly format data.frame column names; provide quick counts of variable combinations (i.e., frequency tables and crosstabs); and isolate duplicate records. Other janitor functions nicely format the tabulation results. These tabulate-and-report functions approximate popular features of SPSS and Microsoft Excel. This package follows the principles of the "tidyverse" and works well with the pipe function %>% . janitor was built with beginning-to-intermediate R users in mind and is optimized for user-friendliness. Advanced R users can already do everything covered here, but with janitor they can do it faster and save their thinking for the fun stuff.

URL https://github.com/sfirke/janitor

BugReports https://github.com/sfirke/janitor/issues

Depends R (>= 3.1.2)
Imports dplyr (>= 0.7.0), tidyr (>= 0.7.0), snakecase (>= 0.9.2), magrittr, purrr, rlang

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add_totals_col

### Description
This function is deprecated, use adorn_totals instead.

### Usage
```r
add_totals_col(dat, na.rm = TRUE)
```
**add_totals_row**

**Arguments**

- **dat**: an input data.frame with at least one numeric column.
- **na.rm**: should missing values (including NaN) be omitted from the calculations?

**Value**

Returns a data.frame with a totals column containing row-wise sums.

---

**adorn_crosstab**

**Add presentation formatting to a crosstabulation table.**

**Description**

This function is deprecated, use the adorn_ family of functions instead.

**Usage**

```r
adorn_crosstab(
  dat, 
  denom = "row", 
  show_n = TRUE, 
  digits = 1, 
  show_totals = FALSE, 
  rounding = "half to even"
)
```
Adorn_ns

Arguments

dat a data.frame with row names in the first column and numeric values in all other columns. Usually the piped-in result of a call to crosstab that included the argument percent = "none".
denom the denominator to use for calculating percentages. One of "row", "col", or "all".
show_n should counts be displayed alongside the percentages?
digits how many digits should be displayed after the decimal point?
show_totals display a totals summary? Will be a row, column, or both depending on the value of denom.
rounding method to use for truncating percentages - either "half to even", the base R default method, or "half up", where 14.5 rounds up to 15.

Value

Returns a data.frame.

Description

This function adds back the underlying Ns to a tabyl whose percentages were calculated using adorn_percentages(), to display the Ns and percentages together. You can also call it on a non-tabyl data.frame with tabyl-like format to which you wish to append Ns.

Usage

adorn_ns(dat, position = "rear", ns = attr(dat, "core"))

Arguments

dat a data.frame of class tabyl that has had adorn_percentages and/or adorn_pct_formatting called on it. If given a list of data.frames, this function will apply itself to each data.frame in the list (designed for 3-way tabyl lists).
position should the N go in the front, or in the rear, of the percentage?
ns the Ns to append. The default is the "core" attribute of the input tabyl dat, where the original Ns of a two-way tabyl are stored. However, if you need to modify the numbers, e.g., to format 4000 as 4,000 or 4k, you can do that separately and supply the formatted result here.

Value

a data.frame with Ns appended
adorn_pct_formatting

Examples

```r
mtcars %>%
  tabyl(am, cyl) %>%
  adorn_percentages("col") %>%
  adorn_pct_formatting() %>%
  adorn_ns(position = "front")
```

---

adorn_pct_formatting  Format a data.frame of decimals as percentages.

Description

Numeric columns get multiplied by 100 and formatted as percentages according to user specifications. This function excludes the first column of the input data.frame, assuming that it contains a descriptive variable. Other non-numeric columns are also excluded.

Usage

```r
adorn_pct_formatting(
  dat,
  digits = 1,
  rounding = "half to even",
  affix_sign = TRUE
)
```

Arguments

dat  a data.frame with decimal values, typically the result of a call to `adorn_percentages` on a `tabyl`. If given a list of data.frames, this function will apply itself to each data.frame in the list (designed for 3-way `tabyl` lists).

digits  how many digits should be displayed after the decimal point?

rounding  method to use for rounding - either "half to even", the base R default method, or "half up", where 14.5 rounds up to 15.

affix_sign  should the % sign be affixed to the end?

Value

a data.frame with formatted percentages

Examples

```r
mtcars %>%
  tabyl(am, cyl) %>%
  adorn_percentages("col") %>%
  adorn_pct_formatting()
```
adorn_percentages  

Convert a data.frame of counts to percentages.

Description

This function excludes the first column of the input data.frame, assuming that it contains a descriptive variable. If the input data.frame is not a tabyl, it will convert to one in order to preserve the underlying values in the core attribute.

Usage

adorn_percentages(dat, denominator = "row", na.rm = TRUE)

Arguments

dat  
a tabyl or other data.frame with a tabyl-like layout. If given a list of data.frames, this function will apply itself to each data.frame in the list (designed for 3-way tabyl lists).

denominator  
the direction to use for calculating percentages. One of "row", "col", or "all".

na.rm  
should missing values (including NaN) be omitted from the calculations?

Value

Returns a data.frame of percentages, expressed as numeric values between 0 and 1.

Examples

```r
mtcars %>%
  tabyl(am, cyl) %>%
  adorn_percentages("col")

# calculates correctly even with totals column and/or row:
mtcars %>%
  tabyl(am, cyl) %>%
  adorn_totals("row") %>%
  adorn_percentages()
```
adorn_rounding

Round the numeric columns in a data.frame.

Description

Can run on any data.frame with at least one numeric column. This function defaults to excluding the first column of the input data.frame, assuming that it contains a descriptive variable, but this can be overridden with the argument `skip_first_col = FALSE`.

If you’re formatting percentages, e.g., the result of `adorn_percentages()`, use `adorn_pct_formatting()` instead. This is a more flexible variant for ad-hoc usage. Compared to `adorn_pct_formatting()`, it can run on the first column and does not multiply by 100 or pad the numbers with spaces for alignment in the results data.frame. This function retains the class of numeric input columns.

Usage

```r
adorn_rounding(
  dat,
  digits = 1,
  rounding = "half to even",
  skip_first_col = TRUE
)
```

Arguments

- `dat`: a data.frame with at least one numeric column
- `digits`: how many digits should be displayed after the decimal point?
- `rounding`: method to use for rounding - either "half to even", the base R default method, or "half up", where 14.5 rounds up to 15.
- `skip_first_col`: should the first column be left unrounded, assuming it contains values of a descriptive variable as in a `tabyl`? Defaults to `TRUE`.

Value

Returns the data.frame with rounded numeric columns.

Examples

```r
mtcars %>%
  tabyl(am, cyl) %>%
  adorn_percentages() %>
  adorn_rounding(digits = 2, rounding = "half up")

# tolerates non-numeric columns:
library(dplyr)
mtcars %>%
tabyl(am, cyl) %>
```
adorn_title

Add column name to the top of a two-way tabyl.

Description

This function adds the column variable name to the top of a tabyl for a complete display of information. This makes the tabyl prettier, but renders the data.frame less useful for further manipulation.

Usage

adorn_title(dat, placement = "top", row_name, col_name)

Arguments

dat a data.frame of class tabyl or other data.frame with a tabyl-like layout. If given a list of data.frames, this function will apply itself to each data.frame in the list (designed for 3-way tabyl lists).

placement whether the column name should be added to the top of the tabyl in an otherwise-empty row "top" or appended to the already-present row name variable ("combined"). The formatting in the "top" option has the look of base R's table(); it also wipes out the other column names, making it hard to further use the data.frame besides formatting it for reporting. The "combined" option is more conservative in this regard.

row_name (optional) default behavior is to pull the row name from the attributes of the input tabyl object. If you wish to override that text, or if your input is not a tabyl, supply a string here.

col_name (optional) default behavior is to pull the column_name from the attributes of the input tabyl object. If you wish to override that text, or if your input is not a tabyl, supply a string here.

Value

the input tabyl, augmented with the column title. Non-tabyl inputs that are of class tbl_df are downgraded to basic data.frames so that the title row prints correctly.

Examples

mtcars %>%
tabyl(am, cyl) %>%
adorn_title(placement = "top")

# Adding a title to a non-tabyl
adorn_totals

```r
library(tidyr); library(dplyr)
mtcars %>%
  group_by(gear, am) %>%
  summarise(avg_mpg = mean(mpg)) %>%
  spread(gear, avg_mpg) %>%
  adorn_title("top", row_name = "Gears", col_name = "Cylinders")
```

adorn_totals

**Append a totals row and/or column to a data.frame.**

**Description**

This function excludes the first column of the input data.frame, assuming it’s a descriptive variable not to be summed. Non-numeric columns are converted to character class and have a user-specified fill character inserted in the totals row.

**Usage**

```r
adorn_totals(dat, where = "row", fill = "-", na.rm = TRUE, name = "Total")
```

**Arguments**

- `dat`: an input data.frame with at least one numeric column. If given a list of data.frames, this function will apply itself to each data.frame in the list (designed for 3-way tabyl lists).
- `where`: one of "row", "col", or c("row", "col")
- `fill`: if there are multiple non-numeric columns, what string should fill the bottom row of those columns?
- `na.rm`: should missing values (including NaN) be omitted from the calculations?
- `name`: name of the totals column or row

**Value**

Returns a data.frame augmented with a totals row, column, or both. The data.frame is now also of class tabyl and stores information about the attached totals and underlying data in the tabyl attributes.

**Examples**

```r
mtcars %>%
  tabyl(am, cyl) %>%
  adorn_totals()
```
**as_tabyl**

Add tabyl attributes to a data.frame.

**Description**

A tabyl is a data.frame containing counts of a variable or co-occurrences of two variables (a.k.a., a contingency table or crosstab). This specialized kind of data.frame has attributes that enable adorn_ functions to be called for precise formatting and presentation of results. E.g., display results as a mix of percentages, Ns, add totals rows or columns, rounding options, in the style of Microsoft Excel PivotTable.

A tabyl can be the result of a call to janitor::tabyl(), in which case these attributes are added automatically. This function adds tabyl class attributes to a data.frame that isn’t the result of a call to tabyl but meets the requirements of a two-way tabyl:

1) First column contains values of variable 1
2) Column names 2:n are the values of variable 2
3) Numeric values in columns 2:n are counts of the co-occurrences of the two variables.*

* = this is the ideal form of a tabyl, but janitor’s adorn_ functions tolerate and ignore non-numeric columns in positions 2:n.

For instance, the result of dplyr::count() followed by tidyr::spread() can be treated as a tabyl.

The result of calling tabyl() on a single variable is a special class of one-way tabyl; this function only pertains to the two-way tabyl.

**Usage**

`as_tabyl(dat, axes = 2, row_var_name = NULL, col_var_name = NULL)`

**Arguments**

- `dat`: a data.frame with variable values in the first column and numeric values in all other columns.
- `axes`: is this a two_way tabyl or a one_way tabyl? If this function is being called by a user, this should probably be "2". One-way tabyls are created by tabyl but are a special case.
- `row_var_name`: (optional) the name of the variable in the row dimension; used by adorn_title().
- `col_var_name`: (optional) the name of the variable in the column dimension; used by adorn_title().

**Value**

Returns the same data.frame, but with the additional class of "tabyl" and the attribute "core".

**Examples**

```r
as_tabyl(mtcars)
```
chisq.test

Apply stats::chisq.test to a two-way tabyl

Description
This generic function overrides stats::chisq.test. If the passed table is a two-way tabyl, it runs it through janitor::chisq.test.tabyl, otherwise it just calls stats::chisq.test.

Usage
chisq.test(x, ...)

## Default S3 method:
chisq.test(x, y = NULL, ...)

## S3 method for class 'tabyl'
chisq.test(x, tabyl_results = TRUE, ...)

Arguments
x a two-way tabyl, a numeric vector or a factor
...
other parameters passed to stats::chisq.test
y if x is a vector, must be another vector or factor of the same length
tabyl_results if TRUE and x is a tabyl object, also return 'observed', 'expected', 'residuals' and 'stdres' as tabyl

Value
The result is the same as the one of stats::chisq.test. If 'tabyl_results' is TRUE, the returned tables 'observed', 'expected', 'residuals' and 'stdres' are converted to tabyls.

Examples
tab <- tabyl(mtcars, gear, cyl)
chisq.test(tab)
chisq.test(tab)$residuals
clean_names

Cleans names of a data.frame.

Description

Resulting names are unique and consist only of the _ character, numbers, and letters. Capitalization preferences can be specified using the case parameter.

Accented characters are transliterated to ASCII. For example, an "ö" with a German umlaut over it becomes "o", and the Spanish character "enye" becomes "n".

This function takes and returns a data.frame, for ease of piping with `%>%`. For the underlying function that works on a character vector of names, see `make_clean_names`.

Usage

```r
clean_names(dat, case)
```

## Default S3 method:
clean_names(
  dat,
  case = c("snake", "lower_camel", "upper_camel", "screaming_snake", "lower_upper",
             "upper_lower", "all_caps", "small_camel", "big_camel", "old_janitor", "parsed",
             "mixed", "none")
)
```

Arguments

dat the input data.frame.

case The desired target case (default is "snake"), indicated by these possible values:
  • "snake" produces snake_case
  • "lower_camel" or "small_camel" produces lowerCamel
  • "upper_camel" or "big_camel" produces UpperCamel
  • "screaming_snake" or "all_caps" produces ALL_CAPS
  • "lower_upper" produces lowerUPPER
  • "upper_lower" produces UPPERlower
  • old_janitor: legacy compatibility option to preserve behavior of clean_names prior to addition of the "case" argument(janitor versions <= 0.3.1 ). Provided as a quick fix for old scripts broken by the changes to clean_names in janitor v1.0.
  • "parsed", "mixed", "none": less-common cases offered by snakecase::to_any_case. See `to_any_case` for details.

Value

Returns the data.frame with clean names.
**compare_df_cols**

Generate a comparison of data.frames (or similar objects) that indicates if they will successfully bind together by rows.

**Usage**

```r
compare_df_cols(
  ..., 
  return = c("all", "match", "mismatch"), 
  bind_method = c("bind_rows", "rbind"), 
  strict_description = FALSE
)
```

**Arguments**

- `...` A combination of data.frames, tibbles, and lists of data.frames/tibbles. The values may optionally be named arguments; if named, the output column will be the name; if not named, the output column will be the data.frame name (see examples section).
- `return` Should a summary of "all" columns be returned, only return "match"ing columns, or only "mismatch"ing columns?
- `bind_method` What method of binding should be used to determine matches? With "bind_rows", columns missing from a data.frame would be considered a match (as in `dplyr::bind_rows()`); with "rbind", columns missing from a data.frame would be considered a mismatch (as in `base::rbind()`).
- `strict_description` Passed to `describe_class`. Also, see the Details section.
Details

Due to the returned "column_name" column, no input data.frame may be named "column_name".

The strict_description argument is most typically used to understand if factor levels match or are bindable. Factors are typically bindable, but the behavior of what happens when they bind differs based on the binding method ("bind_rows" or "rbind"). Even when strict_description is FALSE, data.frames may still bind because some classes (like factors and characters) can bind even if they appear to differ.

Value

A data.frame with a column named "column_name" with a value named after the input data.frames' column names, and then one column per data.frame (named after the input data.frame). If more than one input has the same column name, the column naming will have suffixes defined by sequential use of base::merge() and may differ from expected naming. The rows within the data.frame-named columns are descriptions of the classes of the data within the columns (generated by describe_class).

See Also

Other Data frame type comparison: compare_df_cols_same(), describe_class()

Examples

```r
compare_df_cols(data.frame(A=1), data.frame(B=2))
# user-defined names
compare_df_cols(dfA=data.frame(A=1), dfB=data.frame(B=2))
# a combination of list and data.frame input
compare_df_cols(listA=list(dfA=data.frame(A=1), dfB=data.frame(B=2)), data.frame(A=3))
```

Description

Check whether a set of data.frames are row-bindable. Calls compare_df_cols() and returns TRUE if there are no mis-matching rows.

Usage

```r
compare_df_cols_same(
  ..., 
  bind_method = c("bind_rows", "rbind"),
  verbose = TRUE
)
```
**Arguments**

...  
bind_method  
verbose  

**Value**  
TRUE if row binding will succeed or FALSE if it will fail.

**See Also**  
Other Data frame type comparison: compare_df_cols(), describe_class()

**Examples**

```r
compare_df_cols_same(data.frame(A=1), data.frame(A=2))
compare_df_cols_same(data.frame(A=1), data.frame(B=2))
compare_df_cols_same(data.frame(A=1), data.frame(B=2), verbose=FALSE)
compare_df_cols_same(data.frame(A=1), data.frame(B=2), bind_method="rbind")
```

---

**convert_to_NA**  
*Convert string values to true NA values.*

**Description**  
Converts instances of user-specified strings into NA. Can operate on either a single vector or an entire data frame.

**Usage**

```r
convert_to_NA(dat, strings)
```

**Arguments**

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>dat</td>
<td>vector or data.frame to operate on.</td>
</tr>
<tr>
<td>strings</td>
<td>character vector of strings to convert.</td>
</tr>
</tbody>
</table>

**Value**  
Returns a cleaned object. Can be a vector, data.frame, or tibble::tbl_df depending on the provided input.
Warning

Deprecated, do not use in new code. Use dplyr::na_if() instead.

See Also

janitor Deprecated

crosstab

Generate a crosstabulation of two vectors.

description

This function is deprecated, use tabyl(dat, var1, var2) instead.

Usage

crosstab(...)

## Default S3 method:
crosstab(vec1, vec2, percent = "none", show_na = TRUE, ...)

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

Arguments

... additional arguments, if calling crosstab on a data.frame.
vec1 the vector to place on the crosstab column. If supplying a data.frame, this should be an unquoted column name.
vec2 the vector to place on the crosstab row. If supplying a data.frame, this should be an unquoted column name.
percent which grouping to use for percentages, if desired (defaults to "none", which returns simple counts). Must be one of "none", "row", "col", or "all".
show_na a logical value indicating whether counts should be displayed where either variable is NA.
.data (optional) a data.frame, in which case vec1 and vec2 should be unquoted column names.

Value

Returns a data.frame with the frequencies of the crosstabulated variables.
describe_class

**Description**

Describe the class(es) of an object

**Usage**

```r
describe_class(x, strict_description = TRUE)
```

## S3 method for class 'factor'
```r
describe_class(x, strict_description = TRUE)
```

## Default S3 method:
```r
describe_class(x, strict_description = TRUE)
```

**Arguments**

- **x**
  The object to describe
- **strict_description**
  Should differing factor levels be treated as differences for the purposes of identifying mismatches? `strict_description = TRUE` is stricter and factors with different levels will be treated as different classes. `FALSE` is more lenient: for class comparison purposes, the variable is just a "factor".

**Details**

For package developers, an S3 generic method can be written for `describe_class()` for custom classes that may need more definition than the default method. This function is called by `compare_df_cols`.

**Value**

A character scalar describing the class(es) of an object where if the scalar will match, columns in a data.frame (or similar object) should bind together without issue.

**Methods (by class)**

- **factor**: Describe factors with their levels and if they are ordered.
- **default**: List all classes of an object.

**See Also**

Other Data frame type comparison: `compare_df_cols_same()`, `compare_df_cols()`
**Examples**

```r
describe_class(1)
describe_class(factor("A"))
describe_class(ordered(c("A", "B")))
describe_class(ordered(c("A", "B")), strict_description=FALSE)
```

---

**Description**

Converts numbers like 42370 into date values like 2016-01-01.

Defaults to the modern Excel date encoding system. However, Excel for Mac 2008 and earlier Mac versions of Excel used a different date system. To determine what platform to specify: if the date 2016-01-01 is represented by the number 42370 in your spreadsheet, it’s the modern system. If it’s 40908, it’s the old Mac system. More on date encoding systems at http://support.office.com/en-us/article/Date-calculations-in-Excel-e7fe7167-48a9-4b96-bb53-5612a800b487.

A list of all timezones is available from `base::OlsonNames()`, and the current timezone is available from `base::Sys.timezone()`.

---

**Usage**

```r
excel_numeric_to_date(
  date_num,
  date_system = "modern",
  include_time = FALSE,
  round_seconds = TRUE,
  tz = ""
)
```

---

**Arguments**

- `date_num`: numeric vector of serial numbers to convert.
- `date_system`: the date system, either "modern" or "mac pre-2011".
- `include_time`: Include the time (hours, minutes, seconds) in the output? (See details)
- `round_seconds`: Round the seconds to an integer (only has an effect when include_time is TRUE)?
- `tz`: Time zone, used when include_time = TRUE (see details for more information on timezones).

---

**Details**

When using include_time=TRUE, days with leap seconds will not be accurately handled as they do not appear to be accurately handled by Windows (as described in https://support.microsoft.com/en-us/help/2722715/support-for-the-leap-second).
Value

Returns a vector of class Date if include_time is FALSE. Returns a vector of class POSIXlt if include_time is TRUE.

Examples

eexcel_numeric_to_date(40000)
eexcel_numeric_to_date(40000.5) # No time is included
eexcel_numeric_to_date(40000.5, include_time = TRUE) # Time is included
eexcel_numeric_to_date(40000.521, include_time = TRUE) # Time is included
eexcel_numeric_to_date(40000.521, include_time = TRUE,
   round_seconds = FALSE) # Time with fractional seconds is included

f

f

Description

This generic function overrides stats::fisher.test. If the passed table is a two-way tabyl, it runs it through janitor::fisher.test.tabyl, otherwise it just calls stats::fisher.test.

Usage

fisher.test(x, ...)

## Default S3 method:
fisher.test(x, y = NULL, ...)

## S3 method for class 'tabyl'
fisher.test(x, ...)

Arguments

x a two-way tabyl, a numeric vector or a factor

... other parameters passed to stats::fisher.test

y if x is a vector, must be another vector or factor of the same length

Value

The result is the same as the one of stats::fisher.test.

Examples

tab <- tabyl(mtcars, gear, cyl)
fisher.test(tab)
get_dupes

Get rows of a data.frame with identical values for the specified variables.

Description

For hunting duplicate records during data cleaning. Specify the data.frame and the variable combination to search for duplicates and get back the duplicated rows.

Usage

get_dupes(dat, ...)

Arguments

dat      the input data.frame.
...
unquoted variable names to search for duplicates.

Value

Returns a data.frame (actually a tbl_df) with the full records where the specified variables have duplicated values, as well as a variable dupe_count showing the number of rows sharing that combination of duplicated values.

Examples

get_dupes(mtcars, mpg, hp)
# or called with the magrittr pipe %>% :
mtcars %>% get_dupes(wt)

janitor_deprecated

Deprecated Functions in Package janitor

Description

These functions are provided for compatibility with older versions of janitor only, and may be defunct as soon as the next release.

Details

- use_first_valid_of
- convert_to_NA
**make_clean_names**  
>Cleans a vector of text, typically containing the names of an object.

**Description**

Resulting strings are unique and consist only of the `_` character, numbers, and letters. Capitalization preferences can be specified using the `case` parameter.

For use on the names of a data.frame, e.g., in a `%%` pipeline, call the convenience function `clean_names`.

Accented characters are transliterated to ASCII. For example, an "ö" with a German umlaut over it becomes "o", and the Spanish character "enye" becomes "n".

**Usage**

```r
make_clean_names(
  string,
  case = c("snake", "lower_camel", "upper_camel", "screaming_snake", "lower_upper",
           "upper_lower", "all_caps", "small_camel", "big_camel", "old_janitor", "parsed",
           "mixed", "none")
)
```

**Arguments**

- `string` A character vector of names to clean.
- `case` The desired target case (default is "snake"), indicated by these possible values:
  - "snake" produces snake_case
  - "lower_camel" or "small_camel" produces lowerCamel
  - "upper_camel" or "big_camel" produces UpperCamel
  - "screaming_snake" or "all_caps" produces ALL_CAPS
  - "lower_upper" produces lowerUPPER
  - "upper_lower" produces UPPERlower
  - `old_janitor`: legacy compatibility option to preserve behavior of `clean_names` prior to addition of the "case" argument (janitor versions < 0.3.1). Provided as a quick fix for old scripts broken by the changes to `clean_names` in janitor v1.0.
  - "parsed", "mixed", "none": less-common cases offered by `snakecase::to_any_case`. See `to_any_case` for details.

**Value**

Returns the “cleaned” character vector.
Examples

# cleaning the names of a vector:
x <- structure(1:3, names = c("name with space", "TwoWords", "total $ (2009)"))
x
names(x) <- make_clean_names(names(x))
x # now has cleaned names
# if you prefer camelCase variable names:
make_clean_names(names(x), "small_camel")

# similar to janitor::clean_names(poorly_named_df):
# not run:
# make_clean_names(names(poorly_named_df))

remove_constant Remove constant columns from a data.frame or matrix.

Description

Remove constant columns from a data.frame or matrix.

Usage

remove_constant(dat, na.rm = FALSE)

Arguments

dat the input data.frame or matrix.

na.rm should NA values be removed when considering whether a column is constant? The default value of FALSE will result in a column not being removed if it’s a mix of a single value and NA.

See Also

remove_empty() for removing empty columns or rows.

Other remove functions: remove_empty()

Examples

remove_constant(data.frame(A=1, B=1:3))

# To find the columns that are constant
data.frame(A=1, B=1:3) %>%
dplyr::select_at(setdiff(names(.), names(remove_constant(.)))) %>%
unique()
**remove_empty**

Remove empty rows and/or columns from a data.frame or matrix.

**Description**

Removes all rows and/or columns from a data.frame or matrix that are composed entirely of NA values.

**Usage**

```r
remove_empty(dat, which = c("rows", "cols"))
```

**Arguments**

- `dat`: the input data.frame or matrix.
- `which`: one of "rows", "cols", or c("rows", "cols"). Where no value of which is provided, defaults to removing both empty rows and empty columns, declaring the behavior with a printed message.

**Value**

Returns the object without its missing rows or columns.

**See Also**

`remove_constant()` for removing constant columns.

Other remove functions: `remove_constant()`

**Examples**

```r
# not run:
# dat %>% remove_empty("rows")
```

---

**remove_empty_cols**

Removes empty columns from a data.frame.

**Description**

This function is deprecated, use `remove_empty("cols")` instead.

**Usage**

```r
remove_empty_cols(dat)
```
remove_empty_rows

Arguments

  dat          the input data.frame.

Value

  Returns the data.frame with no empty columns.

Examples

  # not run:
  # dat %>% remove_empty_cols

remove_empty_rows  Removes empty rows from a data.frame.

Description

  This function is deprecated, use remove_empty("rows") instead.

Usage

  remove_empty_rows(dat)

Arguments

  dat          the input data.frame.

Value

  Returns the data.frame with no empty rows.

Examples

  # not run:
  # dat %>% remove_empty_rows
**round_half_up**

*Round a numeric vector; halves will be rounded up, ala Microsoft Excel.*

**Description**

In base R `round()`, halves are rounded to even, e.g., 12.5 and 11.5 are both rounded to 12. This function rounds 12.5 to 13 (assuming `digits = 0`). Negative halves are rounded away from zero, e.g., -0.5 is rounded to -1.

This may skew subsequent statistical analysis of the data, but may be desirable in certain contexts. This function is implemented exactly from [http://stackoverflow.com/a/12688836](http://stackoverflow.com/a/12688836); see that question and comments for discussion of this issue.

**Usage**

```r
round_half_up(x, digits = 0)
```

**Arguments**

- `x` a numeric vector to round.
- `digits` how many digits should be displayed after the decimal point?

**Examples**

```r
round_half_up(12.5)
round_half_up(1.125, 2)
round_half_up(1.125, 1)
round_half_up(-0.5, 0) # negatives get rounded away from zero
```

---

**round_to_fraction**

*Round to the nearest fraction of a specified denominator.*

**Description**

Round a decimal to the precise decimal value of a specified fractional denominator. Common use cases include addressing floating point imprecision and enforcing that data values fall into a certain set.

E.g., if a decimal represents hours and values should be logged to the nearest minute, `round_to_fraction(x, 60)` would enforce that distribution and 0.57 would be rounded to 0.566667, the equivalent of 34/60. 0.56 would also be rounded to 34/60.

Set `denominator = 1` to round to whole numbers.

The `digits` argument allows for rounding of the subsequent result.
Usage

round_to_fraction(x, denominator, digits = Inf)

Arguments

x A numeric vector
denominator The denominator of the fraction for rounding (a scalar or vector positive integer).
digits Integer indicating the number of decimal places to be used after rounding to the fraction. This is passed to base::round(). Negative values are allowed (see Details). (Inf indicates no subsequent rounding)

Details

If digits is Inf, x is rounded to the fraction and then kept at full precision. If digits is "auto", the number of digits is automatically selected as ceiling(log10(denominator)) + 1.

Value

the input x rounded to a decimal value that has an integer numerator relative to denominator (possibly subsequently rounded to a number of decimal digits).

Examples

round_to_fraction(1.6, denominator = 2)
round_to_fraction(pi, denominator = 7) # 22/7
round_to_fraction(c(8.1, 9.2), denominator = c(7, 8))
round_to_fraction(c(8.1, 9.2), denominator = c(7, 8), digits = 3)
round_to_fraction(c(8.1, 9.2, 10.3), denominator = c(7, 8, 1001), digits = "auto")

row_to_names

Elevate a row to be the column names of a data.frame.

Description

Elevate a row to be the column names of a data.frame.

Usage

row_to_names(dat, row_number, remove_row = TRUE, remove_rows_above = TRUE)

Arguments

dat The input data.frame
row_number The row of dat containing the variable names
remove_row Should the row row_number be removed from the resulting data.frame?
remove_rows_above If row_number != 1, should the rows above row_number - that is, between 1:(row_number-1) - be removed from the resulting data.frame?
tabyl

Value
A data.frame with new names (and some rows removed, if specified)

Examples

```r
x <- data.frame(X_1 = c(NA, "Title", 1:3),
                 X_2 = c(NA, "Title2", 4:6))
x %>%
  row_to_names(row_number = 2)
```

Description

A fully-featured alternative to `table()`. Results are data.frames and can be formatted and enhanced with janitor's family of `adorn_` functions.

Specify a data.frame and the one, two, or three unquoted column names you want to tabulate. Three variables generates a list of 2-way tabyls, split by the third variable.

Alternatively, you can tabulate a single variable that isn’t in a data.frame by calling `tabyl` on a vector, e.g., `tabyl(mtcars$gear)`.

Usage

```r
tabyl(dat, ...)
## Default S3 method:
tabyl(dat, show_na = TRUE, show_missing_levels = TRUE, ...)
## S3 method for class `data.frame`
tabyl(dat, var1, var2, var3, show_na = TRUE, show_missing_levels = TRUE, ...)
```

Arguments

dat a data.frame containing the variables you wish to count. Or, a vector you want to tabulate.

... the arguments to `tabyl` (here just for the sake of documentation compliance, as all arguments are listed with the vector- and data.frame-specific methods)

show_na should counts of NA values be displayed? In a one-way tabyl, the presence of NA values triggers an additional column showing valid percentages (calculated excluding NA values).

show_missing_levels should counts of missing levels of factors be displayed? These will be rows and/or columns of zeroes. Useful for keeping consistent output dimensions even when certain factor levels may not be present in the data.
```
var1  the column name of the first variable.
var2  (optional) the column name of the second variable (the rows in a 2-way tabulation).
var3  (optional) the column name of the third variable (the list in a 3-way tabulation).
```

**Value**

Returns a data.frame with frequencies and percentages of the tabulated variable(s). A 3-way tabulation returns a list of data.frames.

**Examples**

```r
tabyl(mtcars, cyl)
tabyl(mtcars, cyl, gear)
tabyl(mtcars, cyl, gear, am)

# or using the %>% pipe
mtcars %>%
tabyl(cyl, gear)

# illustrating show_na functionality:
my_cars <- rbind(mtcars, rep(NA, 11))
my_cars %>%
tabyl(cyl, show_na = FALSE)

# Calling on a single vector not in a data.frame:
val <- c("hi", "med", "med", "lo")
tabyl(val)
```

---

**top_levels**

*Generate a frequency table of a factor grouped into top-n, bottom-n, and all other levels.*

**Description**

Get a frequency table of a factor variable, grouped into categories by level.

**Usage**

```r
top_levels(input_vec, n = 2, show_na = FALSE)
```

**Arguments**

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>input_vec</td>
<td>the factor variable to tabulate.</td>
</tr>
<tr>
<td>n</td>
<td>number of levels to include in top and bottom groups</td>
</tr>
<tr>
<td>show_na</td>
<td>should cases where the variable is NA be shown?</td>
</tr>
</tbody>
</table>
Value

Returns a data.frame (actually a tbl_df) with the frequencies of the grouped, tabulated variable. Includes counts and percentages, and valid percentages (calculated omitting NA values, if present in the vector and show_na = TRUE.)

Examples

```r
top_levels(as.factor(mtcars$hp), 2)
```

---

**untabyl**

Remove tabyl attributes from a data.frame.

Description

Strips away all tabyl-related attributes from a data.frame.

Usage

```r
untabyl(dat)
```

Arguments

dat a data.frame of class tabyl.

Value

Returns the same data.frame, but without the tabyl class and attributes.

Examples

```r
mtcars %>%
tabyl(am) %>%
untabyl() %>%
attributes() # tabyl-specific attributes are gone
```
use_first_valid_of

Returns first non-NA value from a set of vectors.

Description

At each position of the input vectors, iterates through in order and returns the first non-NA value. This is a robust replacement of the common `ifelse(!is.na(x), x, ifelse(!is.na(y), y, z))`. It's more readable and handles problems like `ifelse`'s inability to work with dates in this way.

Usage

use_first_valid_of(..., if_all_NA = NA)

Arguments

... the input vectors. Order matters: these are searched and prioritized in the order they are supplied.

if_all_NA what value should be used when all of the vectors return NA for a certain index? Default is NA.

Value

Returns a single vector with the selected values.

Warning

Deprecated, do not use in new code. Use `dplyr::coalesce()` instead.

See Also

`janitor_deprecated`
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