

# Package ‘autoslider.core’

June 13, 2025

**Type** Package

**Title** Slide Automation for Tables, Listings and Figures

**Version** 0.2.5

**Description** The normal process of creating clinical study slides is that a statistician manually type in the numbers from outputs and a separate statistician to double check the typed in numbers. This process is time consuming, resource intensive, and error prone. Automatic slide generation is a solution to address these issues. It reduces the amount of work and the required time when creating slides, and reduces the risk of errors from manually typing or copying numbers from the output to slides. It also helps users to avoid unnecessary stress when creating large amounts of slide decks in a short time window.

**License** Apache License 2.0

**URL** <https://github.com/insightengineering/autoslider.core>

**BugReports** <https://github.com/insightengineering/autoslider.core/issues>

**Depends** R (>= 4.1.0)

**Imports** assertthat (>= 0.2.1), checkmate (>= 2.2.0), cli (>= 3.6.3), dplyr (>= 1.1.3), flextable (>= 0.9.4), forcats (>= 1.0.0), ggplot2 (>= 3.5.0), ggpubr (>= 0.6.0), grid, gridExtra (>= 2.3), gtsummary (>= 2.0.3), methods, officer (>= 0.3.18), rlang (>= 1.1.1), rlistings (>= 0.2.11), rtables (>= 0.6.12), rvg (>= 0.2.5), stringr (>= 1.5.0), survival, tern (>= 0.9.8), tidyr (>= 1.3.0), yaml (>= 2.3.7)

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gagolews/stringi, tidymodels/broom, cran/car, tidyverse/dplyr,  
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autoslider.core-package

*autoslider.core Package*

---

## Description

The normal process of creating clinical study slides is that a statistician manually type in the numbers from outputs and a separate statistician to double check the typed in numbers. This process is time consuming, resource intensive, and error prone. Automatic slide generation is a solution to address these issues. It reduces the amount of work and the required time when creating slides, and reduces the risk of errors from manually typing or copying numbers from the output to slides. It also helps users to avoid unnecessary stress when creating large amounts of slide decks in a short time window.

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**See Also**

Useful links:

- <https://github.com/insightengineering/autoslider.core>
- Report bugs at <https://github.com/insightengineering/autoslider.core/issues>

---

adding\_ai\_footnotes     *Update footnote with AI response*

---

**Description**

Update footnote with AI response

**Usage**

```
adding_ai_footnotes(outputs, prompt_list, platform, base_url, api_key, model)
```

**Arguments**

outputs	Output objects
prompt_list	List of prompt
platform	Platform provider
base_url	Base url
api_key	API key
model	Model of choice

---

autoslider_error	<i>autoslider_error class</i>
------------------	-------------------------------

---

**Description**

autoslider\_error class

**Usage**

```
autoslider_error(x, spec, step)
```

**Arguments**

x	character scaler
spec	spec should be a list containing "program" and "suffix"
step	step is a character indicating in which step the pipeline encounter error

**Details**

this function is used to create autoslider\_error object. this function is for internal use only to create the autoslider\_error object. It enable us for further functionalities, like providing help on easy debugging, e.g. if the error is inside the user function, provide the call and let the user run the code outside the pipeline.

**Value**

autoslider\_error object

---

autoslider\_format      *Table color and font*

---

## Description

Zebra themed color

## Usage

```
autoslider_format(
  ft,
  odd_header = "#0EAED5",
  odd_body = "#EBF5FA",
  even_header = "#0EAED5",
  even_body = "#D0E4F2",
  font_name = "arial",
  body_font_size = 12,
  header_font_size = 14
)

blue_format(ft, ...)

orange_format(ft, ...)

red_format(ft, ...)

purple_format(ft, ...)

autoslider_dose_format(ft, header_vals = names(ft))

black_format_ae(ft, body_font_size = 8, header_font_size = 8, ...)

black_format_tb(ft, body_font_size = 8, header_font_size = 8, ...)
```

## Arguments

ft	flextable object
odd_header	Hex color code, default to deep sky blue
odd_body	Hex color code, default to alice blue
even_header	Hex color code, default to slate gray
even_body	Hex color code, default to slate gray
font_name	Font name, default to arial
body_font_size	Font size of the table content, default to 12
header_font_size	Font size of the table header, default to 14

... arguments passed to program  
 header\_vals Header

### Value

A flextable with applied theme.

### Functions

- autoslider\_format(): User defined color code and font size
- blue\_format(): Blue color theme
- orange\_format(): Orange color theme
- red\_format(): Red color theme
- purple\_format(): Purple color theme
- autoslider\_dose\_format(): 'AutoslideR' dose formats
- black\_format\_ae(): Black color theme for AE listing
- black\_format\_tb(): Black color theme

### Author(s)

Nina Qi and Jasmina Uzunovic

---

build\_table\_header *Build table header, a utility function to help with construct structured header for table layout*

---

### Description

Build table header, a utility function to help with construct structured header for table layout

### Usage

```
build_table_header(anl, arm, split_by_study, side_by_side)
```

### Arguments

anl analysis data object  
 arm Arm variable for column split  
 split\_by\_study if true, construct structured header with the study ID  
 side\_by\_side A logical value indicating whether to display the data side by side.

### Value

A 'rtables' layout with desired header.



---

center_figure_loc	<i>Create location container to center the figure, based on ppt size and user specified figure size</i>
-------------------	---

---

**Description**

Create location container to center the figure, based on ppt size and user specified figure size

**Usage**

```
center_figure_loc(fig_width, fig_height, ppt_width, ppt_height)
```

**Arguments**

fig_width	Figure width
fig_height	Figure height
ppt_width	Slide width
ppt_height	Slide height

**Value**

Location for a placeholder from scratch

---

center_table_loc	<i>create location container to center the table</i>
------------------	--

---

**Description**

create location container to center the table

**Usage**

```
center_table_loc(ft, ppt_width, ppt_height)
```

**Arguments**

ft	Flextable object
ppt_width	Powerpoint width
ppt_height	Powerpoint height

**Value**

Location for a placeholder

---

check\_and\_set\_cutoff *Assert function to check the cutoff*

---

**Description**

Assert function to check the cutoff

**Usage**

```
check_and_set_cutoff(data, cutoff)
```

**Arguments**

data	dataframe
cutoff	cutoff threshold

**Value**

Set the cutoff value

---

decorate *generic function decorate*

---

**Description**

generic function decorate  
s3 method for decorate

**Usage**

```
decorate(x, ...)
```

```
decorate(x, ...)
```

**Arguments**

x	object to decorate
...	additional arguments passed to methods

**Value**

No return value, called for side effects

---

decorate,listing\_df-method  
*decorate listing*

---

**Description**

decorate listing

**Usage**

```
## S4 method for signature 'listing_df'
decorate(x, titles = "", footnotes = "", paper = "P8", for_test = FALSE, ...)
```

**Arguments**

x	A listing_df object representing the data to be decorated.
titles	Title to be added to the table.
footnotes	Footnote to be added to the table
paper	Orientation and font size as string, e.g. "P8"; "L11"
for_test	'logic' CICD parameter
...	Additional arguments. not used.

**Value**

No return value, called for side effects

---

decorate,VTableTree-method  
*Decorate TableTree*

---

**Description**

Decorate TableTree

**Usage**

```
## S4 method for signature 'VTableTree'
decorate(x, titles = "", footnotes = "", paper = "P8", for_test = FALSE, ...)
```

**Arguments**

x	A VTableTree object representing the data to be decorated.
titles	Title to be added to the table.
footnotes	Footnote to be added to the table
paper	Orientation and font size as string, e.g. "P8"; "L11"
for_test	'logic' CICD parameter
...	Additional arguments passed to the decoration function.

**Value**

No return value, called for side effects

---

`decorate.autoslider_error`

*decorate method for autoslider\_error class*

---

**Description**

decorate method for autoslider\_error class

**Usage**

`decorate.autoslider_error(x, ...)`

**Arguments**

x	object to decorate
...	additional arguments. not used.

**Value**

No return value, called for side effects

---

decorate.default	<i>default method to decorate</i>
------------------	-----------------------------------

---

**Description**

default method to decorate

**Usage**

```
decorate.default(x, ...)
```

**Arguments**

x	object to decorate
...	additional arguments. not used.

**Value**

No return value, called for side effects

---

decorate.ggplot	<i>Decorate ggplot object</i>
-----------------	-------------------------------

---

**Description**

Decorate ggplot object

**Usage**

```
decorate.ggplot(
  x,
  titles = "",
  footnotes = "",
  paper = "L11",
  for_test = FALSE,
  ...
)
```

**Arguments**

x	An object to decorate
titles	Plot titles
footnotes	Plot footnotes
paper	Paper size, by default "L11"
for_test	'logic' CICD parameter
...	additional arguments. not used.

**Details**

The paper default paper size, 'L11', indicate that the fontsize is 11. The fontsize of the footnotes, is the fontsize of the titles minus 2.

**Value**

No return value, called for side effects

---

decorate.grob	<i>decorate grob</i>
---------------	----------------------

---

**Description**

decorate grob

**Usage**

```
decorate.grob(
  x,
  titles = "",
  footnotes = "",
  paper = "L11",
  for_test = FALSE,
  ...
)
```

**Arguments**

x	object to decorate
titles	graph titles
footnotes	graph footnotes
paper	paper size. default is "L8".
for_test	'logic' CICD parameter
...	Additional arguments. not used.

**Details**

The paper default paper size, 'L11', indicate that the fontsize is 11. The fontsize of the footnotes, is the fontsize of the titles minus 2.

**Value**

No return value, called for side effects

---

decorate.gtsummary     *decorate gtsummary*

---

## Description

decorate gtsummary

## Usage

```
decorate.gtsummary(  
  x,  
  titles = "",  
  footnotes = "",  
  paper = "L11",  
  for_test = FALSE,  
  ...  
)
```

## Arguments

x	gtsummary object to decorate
titles	graph titles
footnotes	graph footnotes
paper	paper size. default is "L8".
for_test	'logic' CICD parameter
...	Additional arguments. not used.

## Details

The paper default paper size, 'L11', indicate that the fontsize is 11. The fontsize of the footnotes, is the fontsize of the titles minus 2.#'

## Value

No return value, called for side effects

---

decorate.list	<i>decorate list of grobs</i>
---------------	-------------------------------

---

**Description**

decorate list of grobs

**Usage**

```
decorate.list(x, titles, footnotes, paper = "L11", for_test = FALSE, ...)
```

**Arguments**

x	object to decorate
titles	graph titles
footnotes	graph footnotes
paper	paper size. default is "L11".
for_test	'logic' CICD parameter
...	additional arguments. not used

**Details**

The paper default paper size, 'L11', indicate that the fontsize is 11. The fontsize of the footnotes, is the fontsize of the titles minus 2.

**Value**

No return value, called for side effects

---

decorate_outputs	<i>Decorate outputs</i>
------------------	-------------------------

---

**Description**

Decorate outputs with titles and footnotes

**Usage**

```
decorate_outputs(
  outputs,
  generic_title = NULL,
  generic_footnote = "Confidential and for internal use only",
  version_label = get_version_label_output(),
  for_test = FALSE
)
```



**Arguments**

outputs            'list' of output objects as created by 'generate\_outputs'  
 generic\_title    'character' vector of titles  
 generic\_footnote            'character' vector of footnotes  
 version\_label    'character'. A version label to be added to the title.  
 for\_test            'logic' CICD parameter

**Details**

'generic\_title' and 'generic\_footnote' will be added to *\*all\** outputs. The use case is to add information such as protocol number and snapshot date defined in a central place (e.g. metadata.yml) to *\*every\** output.

'version\_label' must be either "DRAFT", "APPROVED" or 'NULL'. By default, when outputs are created on the master branch it is set to 'NULL', i.e. no version label will be displayed. Otherwise "DRAFT" will be added. To add "APPROVED" to the title you will need to explicitly set 'version\_label = "APPROVED"'.

**Value**

No return value, called for side effects

---

dec_paste	<i>Concatenate arguments into a string</i>
-----------	--

---

**Description**

Concatenate arguments into a string

**Usage**

```
dec_paste(...)
```

**Arguments**

...            arguments passed to program

**Value**

No return value, called for side effects

---

eg_adae	<i>Cached ADAE</i>
---------	--------------------

---

**Description**

Cached ADAE data

**Usage**

```
data(eg_adae)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 1934 rows and 93 columns.

---

eg_adeg	<i>Cached ADEG</i>
---------	--------------------

---

**Description**

Cached ADEG data

**Usage**

```
data(eg_adeg)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 13600 rows and 88 columns.

---

eg_adex	<i>Cached ADEX</i>
---------	--------------------

---

**Description**

Cached ADEX data

**Usage**

```
data(eg_adex)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 6400 rows and 79 columns.

---

eg_adlb	<i>Cached ADLB</i>
---------	--------------------

---

**Description**

Cached ADLB data

**Usage**

```
data(eg_adlb)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 8400 rows and 102 columns.

---

eg_adrs	<i>Cached ADRS</i>
---------	--------------------

---

**Description**

Cached ADRS data

**Usage**

```
data(eg_adrs)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 3200 rows and 65 columns.

---

eg_ads1	<i>Cached ADSL</i>
---------	--------------------

---

**Description**

Cached ADSL data

**Usage**

```
data(eg_ads1)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 400 rows and 56 columns.

---

eg_adtr	<i>Cached ADTR</i>
---------	--------------------

---

**Description**

Cached ADTR data

**Usage**

```
data(eg_adtr)
```

**Format**

An object of class `data.frame` with 2800 rows and 76 columns.

---

eg_adtte	<i>Cached ADTTE</i>
----------	---------------------

---

**Description**

Cached ADTTE data

**Usage**

```
data(eg_adtte)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 2000 rows and 67 columns.

---

eg_adv	<i>Cached ADVS</i>
--------	--------------------

---

**Description**

Cached ADVS data

**Usage**

```
data(eg_adv)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 16800 rows and 87 columns.

---

fastDoCall	<i>Does do.call quicker, and avoids issues with debug mode within do.call</i>
------------	---

---

**Description**

copied from ms showcase app

**Usage**

```
fastDoCall(what, args, quote = FALSE, envir = parent.frame())
```

**Arguments**

what	either a function or a non-empty character string naming the function to be called.
args	a list of arguments to the function call. The names attribute of args gives the argument names.
quote	a logical value indicating whether to quote the arguments.
envir	an environment within which to evaluate the call. This will be most useful if what is a character string and the arguments are symbols or quoted expressions.

**Value**

No return value, called for side effects

---

figure_to_slide	<i>Add figure to slides</i>
-----------------	-----------------------------

---

**Description**

Add figure to slides

**Usage**

```
figure_to_slide(
  ppt,
  content,
  decor = TRUE,
  fig_width,
  fig_height,
  figure_loc = ph_location_type("body"),
  ...
)
```

**Arguments**

ppt	slide page
content	content to be added
decor	should decoration be added
fig_width	user specified figure width
fig_height	user specified figure height
figure_loc	location of the figure. Defaults to 'ph_location_type("body")'
...	arguments passed to program

**Value**

slide with the added content

---

filter_spec	<i>Filter a spec object</i>
-------------	-----------------------------

---

**Description**

Filter a spec object

**Usage**

```
filter_spec(spec, filter_expr, verbose = TRUE)
```

**Arguments**

spec	A 'spec' object as returned by 'read_spec()'
filter_expr	A 'logical' expression indicating outputs to keep
verbose	Should a message about the number of outputs matching 'filter_spec' be printed? Defaults to 'TRUE'.

**Value**

A 'spec' object containing only the outputs matching 'filter\_expr'

**Author(s)**

Thomas Neitmann ('neitmant')

## Examples

```
library(dplyr)
spec_file <- system.file("spec.yml", package = "autoslider.core")
spec <- spec_file %>% read_spec()

## Keep only the t_dm_IT output
filter_spec(spec, output == "t_dm_IT")

## Same as above but more verbose
filter_spec(spec, program == "t_dm" && suffix == "IT")

## Keep all t_ae outputs
filter_spec(spec, program == "t_ae")

## Keep all output run on safety population
filter_spec(spec, "SE" %in% suffix)

## Keep t_dm_CHN_IT and t_dm_CHN_SE
filter_spec(spec, program == "t_dm" && suffix %in% c("CHN_IT", "CHN_SE"))

## Keep all tables
filter_spec(spec, grepl("^t_", program))
```

---

format_3d	<i>Format of xx.xx (xx.xx, xx.xx)</i>
-----------	---------------------------------------

---

## Description

Format of xx.xx (xx.xx, xx.xx)

## Usage

```
format_3d(x, output)
```

## Arguments

x	input array
output	output handle

## Value

formatted values

---

format_date	<i>Convert dates from 'yyyy-mm-dd' format into 20APR2019 format 'Datetime' format removes the time and outputs date in the same way Able to handle truncated dates as well (e.g. just the year or year and month)</i>
-------------	---

---

### Description

'dplyr::case\_when()' will check all RHS expressions on the input, this means if these expressions return warnings, they will happen even then the input doesn't satisfy the LHS. For this reason, I had to 'quiet' all 'lubridate' functions. This 'format\_date()' function was tested with the inputs in the examples, all gave the expected returned value, so there should be no issues.

### Usage

```
format_date(x)
```

### Arguments

x                      vector of dates in character, in 'yyyy-mm-dd' format

### Value

A vector.

### Examples

```
require(lubridate)

# expected to return "2019"
format_date("2019")

# expected to return "20APR2019"
format_date("2019-04-20")

# expected to return ""
format_date("")

# expected to return "18JUN2019"
format_date("2019-06-18T10:32")

# expected to return "APR2019"
format_date("2019-04")
```



---

func_wrapper	<i>function wrapper to pass filtered data</i>
--------------	---

---

**Description**

function wrapper to pass filtered data

**Usage**

```
func_wrapper(func, datasets, spec, verbose = TRUE)
```

**Arguments**

func	function name
datasets	list of raw datasets
spec	spec
verbose	whether to show verbose information

**Value**

a wrapped function using filtered adam

---

generate_output	<i>Generate output and apply filters, titles, and footnotes</i>
-----------------	---

---

**Description**

Generate output and apply filters, titles, and footnotes

**Usage**

```
generate_output(program, datasets, spec, verbose_level = 2, ...)
```

**Arguments**

program	program name
datasets	list of datasets
spec	spec
verbose_level	Verbose level of messages be displayed. See details for further information.
...	arguments passed to program

**Details**

'verbose\_level' is used to control how many messages are printed out. By default, '2' will show all filter messages and show output generation message. '1' will show output generation message only. '0' will display no message.

**Value**

No return value, called for side effects

**Author(s)**

Liming Li ('Lil128')

**Examples**

```
library(dplyr)
filters::load_filters(
  yaml_file = system.file("filters.yml", package = "autoslider.core"),
  overwrite = TRUE
)

spec_file <- system.file("spec.yml", package = "autoslider.core")
spec <- spec_file %>% read_spec()

data <- list(
  adsl = eg_adsl,
  adae = eg_adae
)
generate_output("t_ae_slide", data, spec$t_ae_slide_SE)
```

---

generate\_outputs      *Generate all outputs from a spec*

---

**Description**

Generate all outputs from a spec

**Usage**

```
generate_outputs(spec, datasets, verbose_level = 2)
```

**Arguments**

spec	Specification list generated by 'read_spec'
datasets	A 'list' of datasets
verbose_level	Verbose level of messages be displayed. See details for further information.

## Details

'verbose\_level' is used to control how many messages are printed out. By default, '2' will show all filter messages and show output generation message. '1' will show output generation message only. '0' will display no message.

## Value

No return value, called for side effects

## Author(s)

- Thomas Neitmann ('neitmant') - Liming Li ('Lil128')

## Examples

```
library(dplyr, warn.conflicts = FALSE)
data <- list(
  adsl = eg_adsl,
  adae = eg_adae
)
filters::load_filters(
  yaml_file = system.file("filters.yml", package = "autoslider.core"),
  overwrite = TRUE
)

spec_file <- system.file("spec.yml", package = "autoslider.core")
spec_file %>%
  read_spec() %>%
  filter_spec(output %in% c("t_dm_slide_IT", "t_ae_slide_SE")) %>%
  generate_outputs(datasets = data)
```

---

generate\_slides

*generate slides based on output*

---

## Description

generate slides based on output

## Usage

```
generate_slides(
  outputs,
  outfile = paste0(tempdir(), "/output.pptx"),
  template = file.path(system.file(package = "autoslider.core"), "theme/basic.pptx"),
  fig_width = 9,
  fig_height = 6,
  t_lpp = 20,
```

```

    t_cpp = 200,
    l_lpp = 20,
    l_cpp = 150,
    ...
  )

```

### Arguments

outputs	List of output
outfile	Out file path
template	Template file path
fig_width	figure width in inch
fig_height	figure height in inch
t_lpp	An integer specifying the table lines per page Specify this optional argument to modify the length of all of the table displays
t_cpp	An integer specifying the table columns per page Specify this optional argument to modify the width of all of the table displays
l_lpp	An integer specifying the listing lines per page Specify this optional argument to modify the length of all of the listings display
l_cpp	An integer specifying the listing columns per page Specify this optional argument to modify the width of all of the listings display
...	arguments passed to program

### Value

No return value, called for side effects

### Examples

```

# Example 1. When applying to the whole pipeline
library(dplyr)
data <- list(
  adsl = eg_adsl %>% dplyr::mutate(FASFL = SAFFL),
  adae = eg_adae
)

filters::load_filters(
  yaml_file = system.file("filters.yml", package = "autoslider.core"),
  overwrite = TRUE
)

spec_file <- system.file("spec.yml", package = "autoslider.core")
spec_file %>%
  read_spec() %>%
  filter_spec(program %in% c("t_dm_slide")) %>%

```

```

generate_outputs(datasets = data) %>%
  decorate_outputs() %>%
  generate_slides()

# Example 2. When applying to an rtable object or an rlisting object
adsl <- eg_adsl
t_dm_slide(adsl, "TRT01P", c("SEX", "AGE")) %>%
  generate_slides()

```

---

gen\_notes

*General notes*


---

### Description

General notes

### Usage

```
gen_notes()
```

### Note

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

---

get\_ellmer\_chat

*Get an 'ellmer' chat API with given platform*


---

### Description

Get an 'ellmer' chat API with given platform

### Usage

```

get_ellmer_chat(
  platform = "deepseek",
  base_url = "https://api.deepseek.com",
  api_key = get_deepseek_key(),
  model = "deepseek-chat"
)

```

### Arguments

platform	Platform provider
base_url	Base url
api_key	API key
model	Model of choice

---

get_prompt_list	<i>Read prompt list from yaml file</i>
-----------------	--

---

**Description**

Read prompt list from yaml file

**Usage**

```
get_prompt_list(filename)
```

**Arguments**

filename	File name
----------	-----------

---

get_proper_title	<i>Adjust title line break and font size</i>
------------------	--

---

**Description**

Adjust title line break and font size

**Usage**

```
get_proper_title(title, max_char = 60, title_color = "#1C2B39")
```

**Arguments**

title	Character string
max_char	Integer specifying the maximum number of characters in one line
title_color	Title color

---

gt_t_dm_slide	<i>Demographic table with gtsummary</i>
---------------	---

---

**Description**

Demographic table with gtsummary

**Usage**

```
gt_t_dm_slide(adsl, arm = "TRT01P", vars = c("AGE", "SEX", "RACE"))
```

**Arguments**

adsl	ADSL data set, dataframe
arm	Arm variable, character, "TRT01P" by default.
vars	Characters of variables

**Value**

gtsummary object

**Note**

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

**Examples**

```
library(dplyr)
adsl <- eg_adsl
out1 <- gt_t_dm_slide(adsl, "TRT01P", c("SEX", "AGE", "RACE", "ETHNIC", "COUNTRY"))
print(out1)
generate_slides(out1, paste0(tempdir(), "/dm.pptx"))
```

---

g_eg_slide	<i>Plot mean values of EG</i>
------------	-------------------------------

---

**Description**

Wrapper for 'g\_mean\_general()'. Requires filtering of the datasets (e.g. using SUFFIX in spec.yml)

**Usage**

```
g_eg_slide(
  adsl,
  adeg,
  arm = "TRT01P",
  paramcd = "PARAM",
  subtitle = "Plot of Mean and 95% Confidence Limits by Visit.",
  ...
)
```

**Arguments**

adsl	ADSL data
adeq	ADVS data
arm	"TRT01P" by default
paramcd	Which variable to use for plotting. By default "PARAM"
subtitle	character scalar forwarded to g_lineplot
...	Gets forwarded to 'tern::g_lineplot()'. This lets you specify additional arguments to 'tern::g_lineplot()'

**Author(s)**

Stefan Thoma ('thomas7')

**Examples**

```
library(dplyr)

adeq_filtered <- eg_adeq %>% filter(
  PARAMCD == "HR"
)
plot_eg <- g_eg_slide(
  adsl = eg_adsl,
  adeg = adeq_filtered,
  arm = "TRT01P",
  paramcd = "PARAM",
  subtitle_add_unit = FALSE
) +
  ggplot2::theme(axis.text.x = ggplot2::element_text(angle = 45, hjust = 1))

generate_slides(plot_eg, paste0(tempdir(), "/g_eg.pptx"))
```



g\_lb\_slide

*Plot mean values of LB***Description**

Wrapper for 'g\_mean\_general()'. Requires filtering of the datasets (e.g. using SUFFIX in spec.yml)

**Usage**

```
g_lb_slide(
  adsl,
  adlb,
  arm = "TRT01P",
  paramcd = "PARAM",
  y = "AVAL",
  subtitle = "Plot of Mean and 95% Confidence Limits by Visit.",
  ...
)
```

**Arguments**

adsl	ADSL data
adlb	ADLB data
arm	"TRT01P" by default
paramcd	character scalar. defaults to By default "PARAM" Which variable to use for plotting.
y	character scalar. Variable to plot on the Y axis. By default "AVAL"
subtitle	character scalar forwarded to g_lineplot
...	Gets forwarded to 'tern::g_lineplot()'. This lets you specify additional arguments to 'tern::g_lineplot()'

**Author(s)**

Stefan Thoma ('thomas7')

**Examples**

```
library(dplyr)

adlb_filtered <- eg_adlb %>% filter(
  PARAMCD == "CRP"
)
plot_lb <- g_lb_slide(
  adsl = eg_adsl,
  adlb = adlb_filtered,
  paramcd = "PARAM",
```

```

  subtitle_add_unit = FALSE
) +
  ggplot2::theme(axis.text.x = ggplot2::element_text(angle = 45, hjust = 1))
generate_slides(plot_lb, paste0(tempdir(), "/g_lb.pptx"))

# Let's plot change values:
plot_lb_chg <- g_lb_slide(
  adsl = eg_adsl,
  adlb = adlb_filtered,
  paramcd = "PARAM",
  y = "CHG",
  subtitle = "Plot of change from baseline and 95% Confidence Limit by Visit."
)
generate_slides(plot_lb_chg, paste0(tempdir(), "/g_lb_chg.pptx"))

```

---

g_mean_general	<i>Plot mean values general function used by wrappers 'g_vs_slide', 'g_lb_slide', &amp; 'g_eg_slide'</i>
----------------	--

---

## Description

adapted from <https://insightengineering.github.io/tlg-catalog/stable/graphs/other/mng01.html>

## Usage

```

g_mean_general(
  adsl,
  data,
  variables = control_lineplot_vars(group_var = "TRT01P"),
  by_vars = c("USUBJID", "STUDYID"),
  subtitle = "Plot of Mean and 95% Confidence Limits by Visit.",
  ...
)

```

## Arguments

adsl	ADSL dataset
data	dataset containing the variable of interest in PARAMCD and AVAL
variables	(named character) vector of variable names in df which should include: <ul style="list-style-type: none"> <li>• x (string) name of x-axis variable.</li> <li>• y (string) name of y-axis variable.</li> <li>• group_var (string or NULL) name of grouping variable (or strata), i.e. treatment arm. Can be NA to indicate lack of groups.</li> </ul>

- `subject_var` (string or NULL)  
name of subject variable. Only applies if `group_var` is not NULL.
- `paramcd` (string or NA)  
name of the variable for parameter's code. Used for y-axis label and plot's subtitle. Can be NA if `paramcd` is not to be added to the y-axis label or subtitle.
- `y_unit` (string or NA)  
name of variable with units of y. Used for y-axis label and plot's subtitle. Can be NA if y unit is not to be added to the y-axis label or subtitle.
- `facet_var` (string or NA)  
name of the secondary grouping variable used for plot faceting, i.e. treatment arm. Can be NA to indicate lack of groups.

`by_vars` variables to merge the two datasets by

`subtitle` character scalar forwarded to `g_lineplot`

`...` additional arguments passed to `'tern::g_lineplot'`

**Author(s)**

Stefan Thoma ('thomas7')

**Examples**

```
library(dplyr)
advs_filtered <- eg_advs %>% filter(
  PARAMCD == "SYSBP"
)
out1 <- g_mean_general(eg_adsl, advs_filtered)
generate_slides(out1, paste0(tempdir(), "/g_mean.pptx"))
```

---

g\_vs\_slide

*Plot mean values of VS*

---

**Description**

Wrapper for `'g_mean_general()'`. Requires filtering of the datasets (e.g. using SUFFIX in `spec.yml`)

**Usage**

```
g_vs_slide(
  adsl,
  advs,
  arm = "TRT01P",
  paramcd = "PARAM",
  subtitle = "Plot of Mean and 95% Confidence Limits by Visit.",
  ...
)
```

**Arguments**

adsl	ADSL data
advs	ADVS data
arm	“TRT01P” by default
paramcd	Which variable to use for plotting. By default “PARAM”
subtitle	character scalar forwarded to <code>g_lineplot</code>
...	Gets forwarded to <code>tern::g_lineplot()</code> . This lets you specify additional arguments to <code>tern::g_lineplot()</code>

**Author(s)**

Stefan Thoma (‘thomas7’)

**Examples**

```
library(dplyr)
advs_filtered <- eg_advs %>% filter(
  PARAMCD == "SYSBP"
)

plot_vs <- g_vs_slide(
  adsl = eg_adsl,
  advs = advs_filtered,
  paramcd = "PARAM",
  subtitle_add_unit = FALSE
) +
  ggplot2::theme(axis.text.x = ggplot2::element_text(angle = 45, hjust = 1))

generate_slides(plot_vs, paste0(tempdir(), "/g_vs.pptx"))
```

---

lyt\_to\_side\_by\_side *Build side by side layout by cbind*

---

**Description**

Build side by side layout by cbind

**Usage**

```
lyt_to_side_by_side(lyt, anl, side_by_side = NULL)
```

**Arguments**

lyt	layout object
anl	analysis data object
side_by_side	A logical value indicating whether to display the data side by side.

**Value**

An ‘rtables’ layout

---

```
lyt_to_side_by_side_two_data
  Build side by side layout by cbind
```

---

**Description**

Build side by side layout by cbind

**Usage**

```
lyt_to_side_by_side_two_data(lyt, anl, alt_counts_df, side_by_side = NULL)
```

**Arguments**

lyt	layout object
anl	analysis data object
alt_counts_df	alternative data frame for counts
side_by_side	A logical value indicating whether to display the data side by side.

**Value**

An ‘rtables’ layout

---

```
l_ae_slide      Adverse      Events      listing      adapted      from
                 https://insightsengineering.github.io/tlg-
                 catalog/stable/listings/adverse-events/ael02.html
```

---

**Description**

Adverse Events listing adapted from <https://insightsengineering.github.io/tlg-catalog/stable/listings/adverse-events/ael02.html>

**Usage**

```
l_ae_slide(ads1, adae)
```

**Arguments**

ads1	ADSL data
adae	ADAE data

## Examples

```
library(dplyr)
library(rlistings)
adsl <- eg_adsl
adae <- eg_adae

out <- l_ae_slide(adsl = adsl, adae = adae)
head(out)
```

---

mutate\_actarm

*Refactor active arm*

---

## Description

Refactor active arm

## Usage

```
mutate_actarm(
  df,
  arm_var = "TRT01A",
  levels = c("PLACEBO + PACLITAXEL + CISPLATIN",
             "ATEZOLIZUMAB + TIRAGOLUMAB + PACLITAXEL + CISPLATIN"),
  labels = c("Pbo+Pbo+PC", "Tira+Atezo+PC")
)
```

## Arguments

df	Input dataframe
arm_var	Arm variable
levels	factor levels
labels	factor labels

## Value

Dataframe with re-level and re-labelled arm variable.

---

na_replace	<i>Replace NAs to NA</i>
------------	--------------------------

---

**Description**

Replace NAs to NA

**Usage**

```
na_replace(table_df)
```

**Arguments**

table\_df      Table dataframe

**Value**

Input dataframe with both column replaced to NA

---

new_round	<i>Founding method</i>
-----------	------------------------

---

**Description**

Founding method

**Usage**

```
new_round(x, digits = 1)
```

**Arguments**

x              number need to be rounded  
digits        number of digits

**Value**

rounded value

---

`null_report`*Null report*

---

**Description**

Null report

**Usage**

```
null_report()
```

**Details**

This will create a null report similar as STREAM does. You can use it inside output functions as shown in the example below.

**Value**

An empty 'rtables' object

**Author(s)**

Thomas Neitmann ('neitmant')

**Examples**

```
library(dplyr)
library(filters)
data <- list(
  adsl = eg_adsl,
  adae = eg_adae %>% mutate(AREL = "")
)

null_report()

## An example how to use the `null_report()` inside an output function
t_ae <- function(datasets) {
  trt <- "ACTARM"
  anl <- semi_join(
    datasets$adae,
    datasets$adsl,
    by = c("STUDYID", "USUBJID")
  )

  return(null_report())
}

data %>%
  filters::apply_filter("SER_SE") %>%
```



t\_ae()

---

perc\_perc                      *Format of (xx%, xx%)*

---

### Description

Format of (xx%, xx%)

### Usage

perc\_perc(x, output)

### Arguments

x	input array
output	output handle

### Value

formatted values

---

ph\_with\_img                      *Placeholder for ph\_with\_img*

---

### Description

Placeholder for ph\_with\_img

### Usage

ph\_with\_img(ppt, figure, fig\_width, fig\_height, figure\_loc)

### Arguments

ppt	power point file
figure	image object
fig_width	width of figure
fig_height	height of figure
figure_loc	location of figure

### Value

Location for a placeholder

---

preprocess\_t\_dd      *Preprocess t\_dd function*

---

**Description**

Preprocess t\_dd function

**Usage**

```
preprocess_t_dd(  
  df,  
  levels = c("PROGRESSIVE DISEASE", "ADVERSE EVENT", "OTHER", "<Missing>"),  
  labels = c("Progressive Disease", "Adverse Events", "Other", "<Missing>")  
)
```

**Arguments**

df	Input dataframe
levels	factor levels
labels	factor labels

**Value**

dataframe

---

preprocess\_t\_ds      *Preprocess t\_ds function*

---

**Description**

Preprocess t\_ds function

**Usage**

```
preprocess_t_ds(  
  df,  
  levels = c("Alive: On Treatment", "Alive: In Follow-up", "<Missing>"),  
  labels = c("Alive: On Treatment", "Alive: In Follow-up", "<Missing>")  
)
```

**Arguments**

df	Input dataframe
levels	factor levels
labels	factor labels

**Value**

dataframe

---

*print.decoratedGrob*    *Print decorated grob*

---

**Description**

Print decorated grob

**Usage**

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

**Arguments**

x                    An object of class 'decoratedGrob'  
...                   not used.

**Value**

No return value, called for side effects

---

*print.decoratedGrobSet*  
                          *Print decorated grob set*

---

**Description**

Print decorated grob set

**Usage**

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

**Arguments**

x                    An object of class 'decoratedGrobSet'  
...                   not used.

**Value**

No return value, called for side effects

---

read_spec	<i>Read yaml spec file</i>
-----------	----------------------------

---

**Description**

Read yaml spec file and split according to filter lists

**Usage**

```
read_spec(spec_file = "spec.yml", metadata = NULL)
```

**Arguments**

spec_file	'character'. Path to a yaml spec file
metadata	Metadata of study

**Value**

An object of class 'spec' which is a 'list' where each element corresponds to one output, e.g. 't\_dm\_IT'.

**Author(s)**

- Liming Li ('Lil128') - Thomas Neitmann ('neitmant') - Joe Zhu

**Examples**

```
spec_file <- system.file("spec.yml", package = "autoslider.core")

## Take a look at the 'raw' content of the spec file
cat(readLines(spec_file)[1:24], sep = "\n")

## This is how it looks once read into R
spec <- read_spec(spec_file)
spec[1:3]
```

---

save_output	<i>Save an Output</i>
-------------	-----------------------

---

**Description**

Save an Output

**Usage**

```

save_output(output, file_name, save_rds = TRUE)

save_output(output, file_name, save_rds = TRUE)

save_output.autoslider_error(output, file_name, save_rds = TRUE)

## S4 method for signature 'dVTableTree'
save_output(output, file_name, save_rds = TRUE)

save_output.decoratedGrob(output, file_name, save_rds = TRUE)

save_output.decoratedGrobSet(output, file_name, save_rds = TRUE)

save_output.dgtsummary(output, file_name, save_rds = TRUE)

save_output.dlisting(output, file_name, save_rds = TRUE)

```

**Arguments**

output	Output object, e.g. an 'rtable' or 'grob'
file_name	Full path of the new file <i>*excluding*</i> the extension
save_rds	Saved as an '.rds' files

**Details**

Tables are saved as RDS file

**Value**

The input 'object' invisibly  
 No return value, called for side effects  
 The input 'object' invisibly  
 The input 'object' invisibly  
 The input 'object' invisibly  
 The input 'object' invisibly

**Examples**

```

library(dplyr)
adsl <- eg_adsl %>%
  filter(SAFFL == "Y") %>%
  mutate(TRT01P = factor(TRT01P, levels = c("A: Drug X", "B: Placebo")))
output_dir <- tempdir()
t_dm_slide(adsl, "TRT01P", c("SEX", "AGE", "RACE", "ETHNIC", "COUNTRY")) %>%
  decorate(
    title = "Demographic table",

```

```

    footnote = ""
  ) %>%
  save_output(
    file_name = file.path(output_dir, "t_dm_SE"),
    save_rds = TRUE
  )

```

---

 save\_outputs

*Save a list of outputs*


---

### Description

Save a list of outputs

### Usage

```

save_outputs(
  outputs,
  outfolder = file.path("output"),
  generic_suffix = NULL,
  save_rds = TRUE,
  verbose_level = 1
)

```

### Arguments

**outputs**            'list' of outputs as created by 'generate\_outputs'  
**outfolder**          Folder in which to store the 'outputs'  
**generic\_suffix**    generic suffix. must be length 1 character or NULL.  
**save\_rds**            Should the input 'outputs' be saved as '.rds' files in addition to '.out' or '.pdf' files? Defaults to 'FALSE'.  
**verbose\_level**    Level of verbose information displayed. Default set to '1'.

### Value

The input 'object' invisibly

### Examples

```

## As `save_outputs` is the last step in the pipeline we have to run
## the 'whole machinery' in order to show its functionality.
library(dplyr, warn.conflicts = FALSE)

data <- list(
  adsl = eg_adsl,
  adae = eg_adae,

```

```
    adtte = eg_adtte
  )

  filters::load_filters(
    yaml_file = system.file("filters.yml", package = "autoslider.core"),
    overwrite = TRUE
  )

  ## For this example the outputs will be saved in a temporary directory. In a
  ## production run this should be the reporting event's 'output' folder instead.
  output_dir <- tempdir()

  spec_file <- system.file("spec.yml", package = "autoslider.core")
  read_spec(spec_file) %>%
    filter_spec(program == "t_dm_slide") %>%
    generate_outputs(datasets = data) %>%
    decorate_outputs() %>%
    save_outputs(outfolder = output_dir)
```

---

slides\_from\_rds

*Generate slides from rds files*

---

## Description

Generate slides from rds files

## Usage

```
slides_from_rds(
  filenames,
  outfile = paste0(tempdir(), "/output.pptx"),
  template = file.path(system.file(package = "autoslider.core"), "theme/basic.pptx")
)
```

## Arguments

filenames	List of file names
outfile	Out file path
template	Template file path

## Value

No return value, called for side effects

## Examples

```
library(dplyr, warn.conflicts = FALSE)

data <- list(
  adsl = eg_adsl,
  adae = eg_adae,
  adtte = eg_adtte
)

filters::load_filters(
  yaml_file = system.file("filters.yml", package = "autoslider.core"),
  overwrite = TRUE
)

## For this example the outputs will be saved in a temporary directory. In a
## production run this should be the reporting event's 'output' folder instead.
output_dir <- tempdir()

spec_file <- system.file("spec.yml", package = "autoslider.core")
read_spec(spec_file) %>%
  filter_spec(program == "t_dm_slide") %>%
  generate_outputs(datasets = data) %>%
  decorate_outputs() %>%
  save_outputs(outfolder = output_dir)

slides_from_rds(list.files(file.path(output_dir, "t_dm_slide_FAS.rds")))
```

---

slides\_preview

*Generate flextable for preview first page*

---

## Description

Generate flextable for preview first page

## Usage

```
slides_preview(x)
```

## Arguments

x                    rtables or data.frame

## Value

A flextable or a ggplot object depending to the input.



**Examples**

```
# Example 1. preview table
library(dplyr)
adsl <- eg_adsl
t_dm_slide(adsl, "TRT01P", c("SEX", "AGE")) %>% slides_preview()
```

---

s_surv_time_1	<i>survival time afun</i>
---------------	---------------------------

---

**Description**

survival time afun

**Usage**

```
s_surv_time_1(df, .var, is_event, control = control_surv_time())
```

**Arguments**

df	data
.var	variable of interest
is_event	vector indicating event
control	'control_surv_time()' by default

**Value**

A function suitable for use in `rtables::analyze()` with element selection, reformatting, and relabeling performed automatically.

---

table_to_slide	<i>Add decorated flextable to slides</i>
----------------	--

---

**Description**

Add decorated flextable to slides

**Usage**

```
table_to_slide(
  ppt,
  content,
  decor = TRUE,
  table_loc = ph_location_type("body"),
  ...
)
```

**Arguments**

ppt	Slide
content	Content to be added
decor	Should table be decorated
table_loc	Table location
...	additional arguments

**Value**

Slide with added content

---

to_vector	<i>Convert list of numbers to vectors</i>
-----------	---

---

**Description**

Convert list of numbers to vectors

**Usage**

```
to_vector(num_list)
```

**Arguments**

num_list	list of numbers
----------	-----------------

**Value**

No return value, called for side effects

---

trial	<i>Cached trial</i>
-------	---------------------

---

**Description**

Cached trial data

**Usage**

```
trial
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 200 rows and 8 columns.

---

trim_perc	<i>Format of xx.xx (xx.x)</i>
-----------	-------------------------------

---

**Description**

Format of xx.xx (xx.x)

**Usage**

```
trim_perc(x, output)
```

**Arguments**

x	input array
output	output handle

**Value**

formatted values

---

trim_perc1	<i>Format of xx.xx (xx.xx)</i>
------------	--------------------------------

---

**Description**

Format of xx.xx (xx.xx)

**Usage**

```
trim_perc1(x, output)
```

**Arguments**

x	input array
output	output handle

**Value**

formatted values

---

t_aesi_slide	<i>Table of AEs of Special Interest adapted from <a href="https://insightsengineering.github.io/tlg-catalog/stable/tables/adverse-events/aet01_aesi.html">https://insightsengineering.github.io/tlg-catalog/stable/tables/adverse-events/aet01_aesi.html</a></i>
--------------	--

---

## Description

Table of AEs of Special Interest adapted from [https://insightsengineering.github.io/tlg-catalog/stable/tables/adverse-events/aet01\\_aesi.html](https://insightsengineering.github.io/tlg-catalog/stable/tables/adverse-events/aet01_aesi.html)

## Usage

```
t_aesi_slide(adsl, adae, aesi, arm = "ACTARM", grad_var = "AETOXGR")
```

## Arguments

adsl	ADSL data set, dataframe
adae	ADAE data set, dataframe.
aesi	AESI variable which will act as a filter to select the rows required to create the table. An example of AESI variable is CQ01NAM.
arm	Arm variable, character, "ACTARM" by default.
grad_var	Grading variable, character, "AETOXGR" by default.

## Value

rtables object

## Author(s)

Kai Xiang Lim ('limk43')

## Examples

```
library(dplyr)
adsl <- eg_adsl
adae <- eg_adae
adae_atoxgr <- adae %>% dplyr::mutate(ATOXGR = AETOXGR)
t_aesi_slide(adsl, adae, aesi = "CQ01NAM")
t_aesi_slide(adsl, adae, aesi = "CQ01NAM", arm = "ARM", grad_var = "AESEV")
t_aesi_slide(adsl, adae_atoxgr, aesi = "CQ01NAM", grad_var = "ATOXGR")
```

---

t_ae_pt_diff_slide	<i>Adverse event table</i>
--------------------	----------------------------

---

**Description**

Adverse event table

**Usage**

```
t_ae_pt_diff_slide(
  adsl,
  adae,
  arm = "TRT01A",
  cutoff = NA,
  split_by_study = FALSE,
  side_by_side = NULL
)
```

**Arguments**

adsl	ADSL data set, dataframe
adae	ADAE data set, dataframe
arm	Arm variable, character, "TRT01A" by default.
cutoff	Cutoff threshold
split_by_study	Split by study, building structured header for tables
side_by_side	"GlobalAsia" or "GlobalAsiaChina" to define the side by side requirement

**Value**

rtables object

**Note**

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

**Examples**

```
library(dplyr)
adsl <- eg_adsl %>%
  dplyr::mutate(TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")))
adae <- eg_adae %>%
  dplyr::mutate(
    TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")),
    ATOXGR = AETOXGR
  )
out <- t_ae_pt_diff_slide(adsl, adae, "TRT01A", 2)
print(out)
generate_slides(out, paste0(tempdir(), "/ae_diff.pptx"))
```

---

t_ae_pt_slide	<i>Adverse event table</i>
---------------	----------------------------

---

### Description

Adverse event table

### Usage

```
t_ae_pt_slide(
  adsl,
  adae,
  arm = "TRT01A",
  cutoff = NA,
  prune_by_total = FALSE,
  split_by_study = FALSE,
  side_by_side = NULL
)
```

### Arguments

adsl	ADSL data set, dataframe
adae	ADAE data set, dataframe
arm	Arm variable, character, "TRT01A" by default.
cutoff	Cutoff threshold
prune_by_total	Prune according total column
split_by_study	Split by study, building structured header for tables
side_by_side	A logical value indicating whether to display the data side by side.

### Value

rtables object

### Note

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

### Examples

```
library(dplyr)
# Example 1
adsl <- eg_adsl %>%
  dplyr::mutate(TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")))
adae <- eg_adae %>%
  dplyr::mutate(
    TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")),
```

```

      ATOXGR = AETOXGR
    )
  out <- t_ae_pt_slide(adsl, adae, "TRT01A", 2)
  print(out)
  generate_slides(out, paste0(tempdir(), "/ae.pptx"))

  # Example 2, prune by total column
  out2 <- t_ae_pt_slide(adsl, adae, "TRT01A", 25, prune_by_total = TRUE)
  print(out2)
  generate_slides(out, paste0(tempdir(), "/ae2.pptx"))

```

---

t\_ae\_pt\_soc\_diff\_slide

*Adverse event table*


---

## Description

Adverse event table

## Usage

```

t_ae_pt_soc_diff_slide(
  adsl,
  adae,
  arm = "TRT01A",
  cutoff = NA,
  split_by_study = FALSE,
  side_by_side = NULL
)

```

## Arguments

adsl	ADSL data set, dataframe
adae	ADAE data set, dataframe
arm	Arm variable, character, "TRT01A" by default.
cutoff	Cutoff threshold
split_by_study	Split by study, building structured header for tables
side_by_side	"GlobalAsia" or "GlobalAsiaChina" to define the side by side requirement

## Value

rtables object

## Note

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

**Examples**

```

library(dplyr)
adsl <- eg_adsl %>%
  dplyr::mutate(TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")))
adae <- eg_adae %>%
  dplyr::mutate(
    TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")),
    ATOXGR = AETOXGR
  )
out <- t_ae_pt_soc_diff_slide(adsl, adae, "TRT01A", 2)
print(out)
generate_slides(out, paste0(tempdir(), "/ae_diff.pptx"))

```

---

t_ae_pt_soc_slide	<i>Adverse event table</i>
-------------------	----------------------------

---

**Description**

Adverse event table

**Usage**

```

t_ae_pt_soc_slide(
  adsl,
  adae,
  arm,
  cutoff = NA,
  prune_by_total = FALSE,
  split_by_study = FALSE,
  side_by_side = NULL
)

```

**Arguments**

adsl	ADSL data set, dataframe
adae	ADAE data set, dataframe
arm	Arm variable, character
cutoff	Cutoff threshold
prune_by_total	Prune according total column
split_by_study	Split by study, building structured header for tables
side_by_side	"GlobalAsia" or "GlobalAsiaChina" to define the side by side requirement

**Value**

rtables object



**Examples**

```

library(dplyr)
# Example 1
adsl <- eg_adsl %>%
  dplyr::mutate(TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")))
adae <- eg_adae %>%
  dplyr::mutate(
    TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")),
    ATOXGR = AETOXGR
  )
out <- t_ae_pt_soc_slide(adsl, adae, "TRT01A", 2)
print(out)
generate_slides(out, paste0(tempdir(), "/ae.pptx"))

# Example 2, prune by total column
out2 <- t_ae_pt_soc_slide(adsl, adae, "TRT01A", 25, prune_by_total = TRUE)
print(out2)
generate_slides(out2, paste0(tempdir(), "/ae2.pptx"))

```

t\_ae\_slide

*Adverse event table***Description**

Adverse event table

**Usage**

```

t_ae_slide(
  adsl,
  adae,
  arm = "TRT01A",
  split_by_study = FALSE,
  side_by_side = NULL
)

```

**Arguments**

adsl	ADSL data set, dataframe
adae	ADAE data set, dataframe
arm	Arm variable, character, "TRT01A" by default.
split_by_study	Split by study, building structured header for tables
side_by_side	should table be displayed side by side

**Value**

rtables object

**Note**

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

**Examples**

```
library(dplyr)
adsl <- eg_adsl %>%
  dplyr::mutate(TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")))
adae <- eg_adae %>%
  dplyr::mutate(
    TRT01A = factor(TRT01A, levels = c("A: Drug X", "B: Placebo")),
    ATOXGR = AETOXGR
  )
out <- t_ae_slide(adsl, adae, "TRT01A")
print(out)
generate_slides(out, paste0(tempdir(), "/ae.pptx"))
```

---

t_ae_summ_slide	<i>Adverse event summary table</i>
-----------------	------------------------------------

---

**Description**

Adverse event summary table

**Usage**

```
t_ae_summ_slide(
  adsl,
  adae,
  arm = "TRT01A",
  dose_adjust_flags = NA,
  dose_adjust_labels = NA,
  gr34_highest_grade_only = TRUE
)
```

**Arguments**

adsl	ADSL dataset, dataframe
adae	ADAE dataset, dataframe
arm	Arm variable, character, "TRT01A" by default.
dose_adjust_flags	Character or a vector of characters. Each character is a variable name in adae dataset. These variables are Logical vectors which flag AEs leading to dose adjustment, such as drug discontinuation, dose interruption and reduction. The flag can be related to any drug, or a specific drug.

**dose\_adjust\_labels**

Character or a vector of characters. Each character represents a label displayed in the AE summary table (e.g. AE leading to discontinuation from drug X). The order of the labels should match the order of variable names in dose\_adjust\_flags.

**gr34\_highest\_grade\_only**

A logical value. Default is TRUE, such that only patients with the highest AE grade as 3 or 4 are included for the count of the "Grade 3-4 AE" and "Treatment-related Grade 3-4 AE" ; set it to FALSE if you want to include patients with the highest AE grade as 5.

**Value**

an rtables object

**Examples**

```
library(dplyr)
ADSL <- eg_adsl
ADAE <- eg_adae

ADAE <- ADAE %>%
  dplyr::mutate(ATOXGR = AETOXGR)
t_ae_summ_slide(adsl = ADSL, adae = ADAE)

# add flag for ae leading to dose reduction
ADAE$reduce_flg <- ifelse(ADAE$AEACN == "DOSE REDUCED", TRUE, FALSE)
t_ae_summ_slide(
  adsl = ADSL, adae = ADAE,
  dose_adjust_flags = c("reduce_flg"),
  dose_adjust_labels = c("AE leading to dose reduction of drug X")
)
# add flgs for ae leading to dose reduction, drug withdraw and drug interruption
ADAE$withdraw_flg <- ifelse(ADAE$AEACN == "DRUG WITHDRAWN", TRUE, FALSE)
ADAE$interrup_flg <- ifelse(ADAE$AEACN == "DRUG INTERRUPTED", TRUE, FALSE)
out <- t_ae_summ_slide(
  adsl = ADSL, adae = ADAE, arm = "TRT01A",
  dose_adjust_flags = c("withdraw_flg", "reduce_flg", "interrup_flg"),
  dose_adjust_labels = c(
    "AE leading to discontinuation from drug X",
    "AE leading to drug X reduction",
    "AE leading to drug X interruption"
  )
)
print(out)
generate_slides(out, paste0(tempdir(), "/ae_summary.pptx"))
```

**Description**

Death table

**Usage**

```
t_dd_slide(adsl, arm = "TRT01A", split_by_study = FALSE, side_by_side = NULL)
```

**Arguments**

adsl	ADSL data set, dataframe
arm	Arm variable, character, "TRT01A" by default.
split_by_study	Split by study, building structured header for tables
side_by_side	used for studies in China. "GlobalAsia" or "GlobalAsiaChina" to define the side by side requirement.

**Value**

rtables object

**Note**

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

**Examples**

```
library(dplyr)
adsl <- eg_adsl %>% preprocess_t_dd()
out1 <- t_dd_slide(adsl, "TRT01A")
print(out1)
generate_slides(out1, paste0(tempdir(), "/dd.pptx"))

out2 <- t_dd_slide(adsl, "TRT01A", split_by_study = TRUE)
print(out2)
```

---

t\_dm\_slide

*Demographic table*

---

**Description**

Demographic table

**Usage**

```
t_dm_slide(  
  adsl,  
  arm = "TRT01P",  
  vars = c("AGE", "SEX", "RACE"),  
  stats = c("median", "range", "count_fraction"),  
  split_by_study = FALSE,  
  side_by_side = NULL  
)
```

**Arguments**

adsl	ADSL data set, dataframe
arm	Arm variable, character, "TRT01P" by default.
vars	Characters of variables
stats	see '.stats' from [tern::analyze_vars()]
split_by_study	Split by study, building structured header for tables
side_by_side	"GlobalAsia" or "GlobalAsiaChina" to define the side by side requirement

**Value**

rtables object

**Note**

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

**Examples**

```
library(dplyr)  
adsl <- eg_adsl  
out1 <- t_dm_slide(adsl, "TRT01P", c("SEX", "AGE", "RACE", "ETHNIC", "COUNTRY"))  
print(out1)  
generate_slides(out1, paste0(tempdir(), "/dm.pptx"))  
  
out2 <- t_dm_slide(adsl, "TRT01P", c("SEX", "AGE", "RACE", "ETHNIC", "COUNTRY"),  
  split_by_study = TRUE  
)  
print(out2)
```

---

t_dor_slide	<i>DOR table</i>
-------------	------------------

---

**Description**

DOR table

**Usage**

```
t_dor_slide(adsl, adtte, arm = "TRT01P", refgroup = NULL)
```

**Arguments**

adsl	ADSL dataset
adtte	ADTTE dataset
arm	Arm variable, character, "TRT01P" by default.
refgroup	Reference group

**Value**

An 'rtables' object

**Note**

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

**Examples**

```
library(dplyr)
adsl <- eg_adsl %>%
  dplyr::mutate(TRT01P = factor(TRT01P, levels = c("A: Drug X", "B: Placebo", "C: Combination")))
adtte <- eg_adtte %>%
  dplyr::filter(PARAMCD == "OS") %>%
  dplyr::mutate(TRT01P = factor(TRT01P, levels = c("A: Drug X", "B: Placebo", "C: Combination")))
out <- t_dor_slide(adsl, adtte)
print(out)
generate_slides(out, paste0(tempdir(), "/dor.pptx"))
```

---

t_ds_slide	<i>Discontinue table</i>
------------	--------------------------

---

**Description**

Discontinue table

**Usage**

```
t_ds_slide(adsl, arm = "TRT01P", split_by_study = FALSE, side_by_side = NULL)
```

**Arguments**

adsl	ADSL data
arm	Arm variable, character, "TRT01P" by default.
split_by_study	Split by study, building structured header for tables
side_by_side	"GlobalAsia" or "GlobalAsiaChina" to define the side by side requirement

**Note**

\* Default arm variables are set to "TRT01A" for safety output, and "TRT01P" for efficacy output

**Examples**

```
library(dplyr)
adsl <- eg_adsl %>%
  mutate(DISTRIFL = sample(c("Y", "N"), size = nrow(eg_adsl), replace = TRUE, prob = c(.1, .9))) %>%
  preprocess_t_ds()
out1 <- t_ds_slide(adsl, "TRT01P")
print(out1)
generate_slides(out1, paste0(tempdir(), "/ds.pptx"))

out2 <- t_ds_slide(adsl, "TRT01P", split_by_study = TRUE)
print(out2)
```

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