Package 'autocogs'

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Title Automatic Cognostic Summaries		
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add_cog_group

Add a cognostic group

Description

Add a new cognostic to be used when calculating automatic cognostics.

Usage

```
add_cog_group(name, fields, description = NA, fn, ...)
```

Arguments

name	Name of cognostic group
fields	data.frame of 'dimension' and 'type' columns. dplyr::bind_rows() of field_info outputs for convenience
description	Description of cognostic group
fn	function to calculate a cognostic group. May return a named list or a single row tibble. Each value of the return data should be the output of cog_desc
	ignored

add_layer_cogs 3

Description

Add a new set of cognostic groups for a given plot layer. If the plot layer is found, the corresponding cognostic groups will be calculated.

Usage

```
add_layer_cogs(name, description, cog_groups, kind = "ggplot", ...)
```

Arguments

name	Name of plot layer. This should match the output of the "name" values of layer_info
description	Description of cognostic group
cog_groups	A data.frame (or tibble) containing the columns: "cog_group", "cols", "name". "cog_group" column should contain a string value of a known cognostic group. "cols" should be a single value or vector of column names to use from the data supplied by layer_info during calculations. "name" should contain the final storage name of the cognostic group.
kind	String value that will match the output of plot_class of the desired plot object
• • •	ignored
autocog	Auto cognostic function

Description

Calculate an auto cognostic function given a name

Usage

```
autocog(.name, ..., .fn_only = FALSE)
```

Arguments

```
.name name of a known cognostic
... arguments passed onto the found function
.fn_only boolean that determines if the function should be returned
```

Examples

```
autocog("univariate_continuous", iris$Sepal.Length)
fn <- autocog("univariate_continuous", .fn_only = TRUE)
fn(iris$Sepal.Length)</pre>
```

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autocog_

Default Cognostic Group Functions

Description

These set of functions comprise the default cognostic groups. Each function produces it's own cognostic information given the required pieces of data.

The functions' print method will display the description. autocog_* functions will take the known_cog_groups() functions and format the output into a single row tibble. Any new known cognostic group function, NAME, will create a function called autocog_NAME, which may be called.

Default Cognostic Group Functions:

- autocog_bivariate_continuous
- autocog_bivariate_counts
- autocog_bivariate_step
- autocog_boxplot
- · autocog_density_2d_continuous
- autocog_density_continuous
- autocog_grouped_counts
- autocog_grouped_testing
- autocog_hex_counts
- autocog_histogram_counts
- autocog_linear_model
- · autocog_loess_model
- autocog_pairwise_counts
- autocog_quantile_quantile
- autocog_scagnostics
- autocog_smooth_line
- · autocog_square_counts
- autocog_univariate_continuous
- autocog_univariate_counts
- autocog_univariate_discrete

Arguments

x data that should appear on an x axis y data that should appear on an y axis

... ignored

direction step direction. Defaults to "hv"

na.rm should NA points be removed when performing calculations

h, n, bins, binwidth, clusters, bw, adjust, kernel, trim, group, groups, center, boundary, closed, pad, bre parameters usually set by corresponding "geoms" to be used within ggplot2

Stat* methods

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See Also

```
known_cog_groups()
```

Examples

```
autocog_bivariate_continuous
autocog_bivariate_continuous(iris$Sepal.Length, iris$Sepal.Width)
```

cog_desc

Cognostic

Description

Add a description to a cognostic (subset metric)

Usage

```
cog_desc(x, desc = NULL)
```

Arguments

x univariate scalardesc description of x

Examples

```
cog_desc(mean(1:10), "mean of 10 numbers")
```

cog_group

Cog group data frame

Description

Make a cog group data frame to be passed into add_layer_cogs

Usage

```
cog\_group(...)
```

Arguments

sets of three values to fill in 'cog_group', 'cols', and 'name'

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Examples

```
cog_group(
  "univariate_discrete", "x", "_x",
  "univariate_counts", "x", "_n"
)
cog_group(
  "univariate_continuous", "x", "_x",
  "univariate_continuous", "y", "_y",
  "bivariate_continuous", c("x", "y"), "_bivar",
  "scagnostics", c("x", "y"), "_scagnostic",
  "bivariate_counts", c("x", "y"), "_n"
)
```

cog_spec

Cognostic Specification

Description

Cognostic Specification

Usage

```
cog_spec(
  bivariate_continuous = TRUE,
 bivariate_counts = TRUE,
 bivariate_step = TRUE,
  boxplot = TRUE,
  density_2d_continuous = TRUE,
  density_continuous = TRUE,
  grouped_counts = TRUE,
  grouped_testing = TRUE,
  hex_counts = TRUE,
 histogram_counts = TRUE,
  linear_model = TRUE,
  loess_model = TRUE,
  pairwise_counts = TRUE,
  quantile_quantile = TRUE,
  scagnostics = TRUE,
  smooth_line = TRUE,
  square_counts = TRUE,
  univariate_continuous = TRUE,
  univariate_counts = TRUE,
  univariate_discrete = TRUE,
  .keep\_layer = TRUE
)
as_cog_specs(p, specs)
```

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Arguments

bivariate_continuous, bivariate_counts, bivariate_step, boxplot, density_2d_continuous, density_continuous, density_continuous

Value

cognostic specification that determines which cogs are added or removed if possible

Examples

```
# example cog specifications
# display like normal
cog_spec(); TRUE
# remove scagnostics
cog_spec(scagnostics = FALSE)
# remove layer
cog_spec(.keep_layer = FALSE); FALSE
# set up data
p <- ggplot2::qplot(Sepal.Length, Sepal.Width, data = iris, geom = c("point", "smooth"))</pre>
dt <- tibble::tibble(panel = list(p))</pre>
# compute cognostics like normal
add_panel_cogs(dt)
# do not compute scagnostics for geom_point cognostics
# compute geom_smooth cognostics
add_panel_cogs(dt, spec = list(cog_spec(scagnostics = FALSE), TRUE))
# do not compute scagnostics for geom_point cognostics
# do not compute geom_smooth cognostics
add_panel_cogs(dt, spec = list(cog_spec(scagnostics = FALSE), FALSE))
```

field_info

Field Type Information

Description

Field Type Information

8 known_layer_cogs

Usage

```
field_info(
  dimension = c("x", "y", "z", "group", "any"),
  type = c("continuous", "discrete", "date", "any")
)
```

Arguments

dimension field name. Use one of the listed options provided type field type. Use one of the listed options provided

known_cog_groups

Cognostic Group information

Description

To add more cognostic groups, please see add_cog_group()

Usage

```
known_cog_groups()
```

Examples

known_cog_groups()

known_layer_cogs

Layer Cognostic groups

Description

Display all layer cognostic information to be paired with information from known_cog_groups().

Usage

```
known_layer_cogs()
```

Examples

```
known_layer_cogs()
```

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layer_count

Plot layer count

Description

Retrieves the number of layers in a given plot

Usage

```
layer_count(p)
## Default S3 method:
layer_count(p)
## S3 method for class 'ggplot'
layer_count(p)
```

Arguments

р

plot object

Value

number

Examples

```
library(ggplot2)
p <- ggplot(iris, aes(Sepal.Length, Sepal.Width)) + geom_point()
layer_count(p) # 1
layer_count(p + geom_smooth(method = "lm") + geom_density_2d()) # 3</pre>
```

layer_info

Data List

Description

Data List

Usage

```
layer_info(p, keep = TRUE, ...)
## Default S3 method:
layer_info(p, keep = TRUE, ...)
## S3 method for class 'ggplot'
layer_info(p, keep = TRUE, ...)
```

plot_class

Arguments

p plot object

keep boolean vector (size = 1 or length(plot\$layers)). Determines if that layer should

have cognostics calculated

... parameters passed on to corresponding layer_info

Examples

```
require(ggplot2)
p <- ggplot(iris, aes(Sepal.Length, Sepal.Width)) +
  geom_point(data = mpg, mapping = aes(cty, hwy))
layer_info(p)</pre>
```

panel_cogs

Panel cognostics

Description

Return or concatenate panel cognostics. For each panel (plot) in the panel column, cognostics will be calculated for each panel. The result will be returned in a nested tibble.

Usage

```
panel_cogs(dt, panel_col = "panel", ...)
add_panel_cogs(dt, panel_col = "panel", ...)
```

Arguments

dt data to be used

panel_col panel column to be used in dt

... parameters passed to layer_info

plot_class

Plot class

Description

First class of the plot object. Exception is ggplot2 as many objects are of class 'gg'

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Usage

```
plot_class(p)
## Default S3 method:
plot_class(p)
## S3 method for class 'gg'
plot_class(p)
## S3 method for class 'ggplot'
plot_class(p)
```

Arguments

p plot object to retrieve class from

Examples

```
library(ggplot2)
p <- qplot(Sepal.Length, Sepal.Width, data = iris)
plot_class(p)</pre>
```

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