Package 'ggplot2.utils'

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Type Package

```
Title Selected Utilities Extending 'ggplot2'
Version 0.3.3
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Description Selected utilities, in particular 'geoms' and 'stats'
      functions, extending the 'ggplot2' package. This package imports
      functions from 'EnvStats' <doi:10.1007/978-1-4614-8456-1> by Millard
      (2013), 'ggpp' <a href="mailto://CRAN.R-project.org/package=ggpp">thttps://CRAN.R-project.org/package=ggpp</a> by Aphalo et
      al. (2023) and 'ggstats' <doi:10.5281/zenodo.10183964> by Larmarange
      (2023), and then exports them. This package also contains modified
      code from 'ggquickeda' <https://CRAN.R-project.org/package=ggquickeda>
      by Mouksassi et al. (2023) for Kaplan-Meier lines and ticks additions
      to plots. All functions are tested to make sure that they work
      reliably.
License Apache License 2.0
URL https://insightsengineering.github.io/ggplot2.utils/
BugReports https://github.com/insightsengineering/ggplot2.utils/issues
Depends ggplot2 (>= 3.3.0), R (>= 3.6)
Imports checkmate, EnvStats, ggpp, ggstats, survival
Suggests dplyr, lifecycle, testthat (>= 3.0.0), tibble, vdiffr
Config/testthat/edition 3
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      'stat_n_text.R' 'stat_prop.R' 'table_themes.R' 'ttheme_set.R'
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```

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ggplot2.utils-package ggplot2.utils Package

Description

ggplot2.utils provides simple access to utility functions extending ggplot2.

Details

Currently all of the functions are imported from other extension packages:

- ggpp: geom_table() and associated functions.
- EnvStats: stat_n_text() and associated functions.
- ggstats: stat_prop() and associated functions.

Author(s)

Maintainer: Daniel Sabanés Bové <daniel.sabanes_bove@rconis.com>
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- Samer Mouksassi (wrote original Kaplan-Meier code)
- Michael Sachs (wrote original Kaplan-Meier code)

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• F. Hoffmann-La Roche AG [copyright holder, funder]

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See Also

Useful links:

- https://insightsengineering.github.io/ggplot2.utils/
- Report bugs at https://github.com/insightsengineering/ggplot2.utils/issues

geom_km

Add a Kaplan-Meier Survival Curve

Description

[Experimental] Adds the Kaplan-Meier survival curve.

Usage

```
geom_km(
  mapping = NULL,
  data = NULL,
  stat = "km",
  position = "identity",
  show.legend = NA,
  inherit.aes = TRUE,
  na.rm = TRUE,
  ...
)
```

Arguments

mapping

Set of aesthetic mappings created by aes(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.

data

The data to be displayed in this layer. There are three options:

If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().

A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

stat

The statistical transformation to use on the data for this layer. When using a geom_*() function to construct a layer, the stat argument can be used the override the default coupling between geoms and stats. The stat argument accepts the following:

• A Stat ggproto subclass, for example StatCount.

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• A string naming the stat. To give the stat as a string, strip the function name of the stat_ prefix. For example, to use stat_count(), give the stat as "count".

 For more information and other ways to specify the stat, see the layer stat documentation.

position

A position adjustment to use on the data for this layer. This can be used in various ways, including to prevent overplotting and improving the display. The position argument accepts the following:

- The result of calling a position function, such as position_jitter(). This method allows for passing extra arguments to the position.
- A string naming the position adjustment. To give the position as a string, strip the function name of the position_ prefix. For example, to use position_jitter(), give the position as "jitter".
- For more information and other ways to specify the position, see the layer position documentation.

show.legend

logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes. It can also be a named logical vector to finely select the aesthetics to display.

inherit.aes

If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. borders().

na.rm

If FALSE, the default, missing values are removed with a warning. If TRUE, missing values are silently removed.

. . .

Other arguments passed on to layer()'s params argument. These arguments broadly fall into one of 4 categories below. Notably, further arguments to the position argument, or aesthetics that are required can *not* be passed through Unknown arguments that are not part of the 4 categories below are ignored.

- Static aesthetics that are not mapped to a scale, but are at a fixed value and apply to the layer as a whole. For example, colour = "red" or linewidth = 3. The geom's documentation has an **Aesthetics** section that lists the available options. The 'required' aesthetics cannot be passed on to the params. Please note that while passing unmapped aesthetics as vectors is technically possible, the order and required length is not guaranteed to be parallel to the input data.
- When constructing a layer using a stat_*() function, the ... argument can be used to pass on parameters to the geom part of the layer. An example of this is stat_density(geom = "area", outline.type = "both"). The geom's documentation lists which parameters it can accept.
- Inversely, when constructing a layer using a geom_*() function, the ... argument can be used to pass on parameters to the stat part of the layer.
 An example of this is geom_area(stat = "density", adjust = 0.5). The stat's documentation lists which parameters it can accept.
- The key_glyph argument of layer() may also be passed on through This can be one of the functions described as key glyphs, to change the display of the layer in the legend.

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Aesthetics

geom_km() understands the following aesthetics (required aesthetics in bold):

- x: the survival/censoring times, automatically mapped by stat_km().
- y: the survival probability estimates, automatically mapped by stat_km().
- alpha
- color
- linetype
- linewidth

Author(s)

Inspired by geom_km written by Michael Sachs (in ggkm) and Samer Mouksassi (in ggquickeda). Here we directly use ggplot2::geom_step() instead of the more general ggplot2::geom_path().

See Also

The default stat for this geom is stat_km().

Examples

```
library(ggplot2)
sex <- rbinom(250, 1, .5)
df <- data.frame(
   time = exp(rnorm(250, mean = sex)),
   status = rbinom(250, 1, .75),
   sex = sex
)
ggplot(df, aes(time = time, status = status, color = factor(sex))) +
   geom_km()</pre>
```

geom_km_ticks

Add Tick Marks to a Kaplan-Meier Survival Curve

Description

[Experimental] Adds tickmarks at the times when there are censored observations but no events.

Usage

```
geom_km_ticks(
  mapping = NULL,
  data = NULL,
  stat = "km_ticks",
  position = "identity",
  show.legend = NA,
  inherit.aes = TRUE,
```

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```
na.rm = TRUE,
...
)
```

Arguments

mapping

Set of aesthetic mappings created by aes(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.

data

The data to be displayed in this layer. There are three options:

If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().

A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula $(e.g. \sim head(.x, 10))$.

stat

The statistical transformation to use on the data for this layer. When using a geom_*() function to construct a layer, the stat argument can be used the override the default coupling between geoms and stats. The stat argument accepts the following:

- A Stat ggproto subclass, for example StatCount.
- A string naming the stat. To give the stat as a string, strip the function name of the stat_ prefix. For example, to use stat_count(), give the stat as "count".
- For more information and other ways to specify the stat, see the layer stat documentation.

position

A position adjustment to use on the data for this layer. This can be used in various ways, including to prevent overplotting and improving the display. The position argument accepts the following:

- The result of calling a position function, such as position_jitter(). This method allows for passing extra arguments to the position.
- A string naming the position adjustment. To give the position as a string, strip the function name of the position_ prefix. For example, to use position_jitter(), give the position as "jitter".
- For more information and other ways to specify the position, see the layer position documentation.

show.legend

logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes. It can also be a named logical vector to finely select the aesthetics to display.

inherit.aes

If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. borders().

na.rm

If FALSE, the default, missing values are removed with a warning. If TRUE, missing values are silently removed.

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. . .

Other arguments passed on to layer()'s params argument. These arguments broadly fall into one of 4 categories below. Notably, further arguments to the position argument, or aesthetics that are required can *not* be passed through Unknown arguments that are not part of the 4 categories below are ignored.

- Static aesthetics that are not mapped to a scale, but are at a fixed value and apply to the layer as a whole. For example, colour = "red" or linewidth = 3. The geom's documentation has an **Aesthetics** section that lists the available options. The 'required' aesthetics cannot be passed on to the params. Please note that while passing unmapped aesthetics as vectors is technically possible, the order and required length is not guaranteed to be parallel to the input data.
- When constructing a layer using a stat_*() function, the ... argument can be used to pass on parameters to the geom part of the layer. An example of this is stat_density(geom = "area", outline.type = "both"). The geom's documentation lists which parameters it can accept.
- Inversely, when constructing a layer using a geom_*() function, the ... argument can be used to pass on parameters to the stat part of the layer. An example of this is geom_area(stat = "density", adjust = 0.5). The stat's documentation lists which parameters it can accept.
- The key_glyph argument of layer() may also be passed on through This can be one of the functions described as key glyphs, to change the display of the layer in the legend.

Aesthetics

geom_km_ticks() understands the following aesthetics (required aesthetics in bold):

- x: the survival/censoring times, automatically mapped by stat_km_ticks().
- y: the survival probability estimates, automatically mapped by stat_km_ticks().
- alpha
- color
- shape
- size
- stroke
- fill

Author(s)

Michael Sachs (in ggkm), Samer Mouksassi (in ggquickeda).

See Also

The default stat for this geom is stat_km_ticks().

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Examples

```
library(ggplot2)
sex <- rbinom(250, 1, .5)
df <- data.frame(
   time = exp(rnorm(250, mean = sex)),
   status = rbinom(250, 1, .75),
   sex = sex
)
ggplot(df, aes(time = time, status = status, color = factor(sex), group = factor(sex))) +
   geom_km() +
   geom_km_ticks(col = "black")</pre>
```

geom_table

Inset tables

Description

[Experimental]

See ggpp::geom_table() for details.

Value

A plot layer instance.

stat_km

Adds a Kaplan-Meier Estimate of Survival Statistic

Description

[Experimental] This stat is for computing the Kaplan-Meier survival estimate for right-censored data. It requires the aesthetic mapping time for the observation times and status which indicates the event status, either 0 for alive and 1 for dead, or 1 for alive and 2 for dead.

Usage

```
stat_km(
  mapping = NULL,
  data = NULL,
  geom = "km",
  position = "identity",
  show.legend = NA,
  inherit.aes = TRUE,
  ...
)
```

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Arguments

mapping

Set of aesthetic mappings created by aes(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.

data

The data to be displayed in this layer. There are three options:

If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().

A data. frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a data. frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

geom

The geometric object to use to display the data for this layer. When using a stat_*() function to construct a layer, the geom argument can be used to override the default coupling between stats and geoms. The geom argument accepts the following:

- A Geom ggproto subclass, for example GeomPoint.
- A string naming the geom. To give the geom as a string, strip the function name of the geom_ prefix. For example, to use geom_point(), give the geom as "point".
- For more information and other ways to specify the geom, see the layer geom documentation.

position

A position adjustment to use on the data for this layer. This can be used in various ways, including to prevent overplotting and improving the display. The position argument accepts the following:

- The result of calling a position function, such as position_jitter(). This
 method allows for passing extra arguments to the position.
- A string naming the position adjustment. To give the position as a string, strip the function name of the position_ prefix. For example, to use position_jitter(), give the position as "jitter".
- For more information and other ways to specify the position, see the layer position documentation.

show.legend

logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes. It can also be a named logical vector to finely select the aesthetics to display.

inherit.aes

If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. borders().

. . .

Other arguments passed on to layer()'s params argument. These arguments broadly fall into one of 4 categories below. Notably, further arguments to the position argument, or aesthetics that are required can *not* be passed through Unknown arguments that are not part of the 4 categories below are ignored.

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• Static aesthetics that are not mapped to a scale, but are at a fixed value and apply to the layer as a whole. For example, colour = "red" or linewidth = 3. The geom's documentation has an **Aesthetics** section that lists the available options. The 'required' aesthetics cannot be passed on to the params. Please note that while passing unmapped aesthetics as vectors is technically possible, the order and required length is not guaranteed to be parallel to the input data.

- When constructing a layer using a stat_*() function, the ... argument can be used to pass on parameters to the geom part of the layer. An example of this is stat_density(geom = "area", outline.type = "both"). The geom's documentation lists which parameters it can accept.
- Inversely, when constructing a layer using a geom_*() function, the ... argument can be used to pass on parameters to the stat part of the layer.
 An example of this is geom_area(stat = "density", adjust = 0.5). The stat's documentation lists which parameters it can accept.
- The key_glyph argument of layer() may also be passed on through This can be one of the functions described as key glyphs, to change the display of the layer in the legend.

Value

A data.frame with columns:

- time: time in data.
- survival: survival estimate at time.

Note

Logical status is not supported.

Author(s)

Michael Sachs (in ggkm), Samer Mouksassi (in ggquickeda).

Examples

```
library(ggplot2)
sex <- rbinom(250, 1, .5)
df <- data.frame(
   time = exp(rnorm(250, mean = sex)),
   status = rbinom(250, 1, .75),
   sex = sex
)
ggplot(df, aes(time = time, status = status, color = factor(sex))) +
   stat_km()</pre>
```

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stat_km_ticks

Adds Tick Marks to a Kaplan-Meier Estimate of Survival Statistic

Description

[Experimental] This stat is for computing the location of the tick marks for the Kaplan-Meier survival estimate for right-censored data. It requires the aesthetic mapping time for the observation times and status which indicates the event status, either 0 for alive and 1 for dead, or 1 for alive and 2 for dead.

Usage

```
stat_km_ticks(
  mapping = NULL,
  data = NULL,
  geom = "km_ticks",
  position = "identity",
  show.legend = NA,
  inherit.aes = TRUE,
   ...
)
```

Arguments

mapping

Set of aesthetic mappings created by aes(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.

data

The data to be displayed in this layer. There are three options:

If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().

A data. frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a data. frame, and will be used as the layer data. A function can be created from a formula (e.g. \sim head(.x, 10)).

geom

The geometric object to use to display the data for this layer. When using a stat_*() function to construct a layer, the geom argument can be used to override the default coupling between stats and geoms. The geom argument accepts the following:

- A Geom ggproto subclass, for example GeomPoint.
- A string naming the geom. To give the geom as a string, strip the function name of the geom_ prefix. For example, to use geom_point(), give the geom as "point".
- For more information and other ways to specify the geom, see the layer geom documentation.

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position

A position adjustment to use on the data for this layer. This can be used in various ways, including to prevent overplotting and improving the display. The position argument accepts the following:

- The result of calling a position function, such as position_jitter(). This
 method allows for passing extra arguments to the position.
- A string naming the position adjustment. To give the position as a string, strip the function name of the position_ prefix. For example, to use position_jitter(), give the position as "jitter".
- For more information and other ways to specify the position, see the layer position documentation.

show.legend

logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes. It can also be a named logical vector to finely select the aesthetics to display.

inherit.aes

If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. borders().

. . .

Other arguments passed on to layer()'s params argument. These arguments broadly fall into one of 4 categories below. Notably, further arguments to the position argument, or aesthetics that are required can *not* be passed through Unknown arguments that are not part of the 4 categories below are ignored.

- Static aesthetics that are not mapped to a scale, but are at a fixed value and apply to the layer as a whole. For example, colour = "red" or linewidth = 3. The geom's documentation has an Aesthetics section that lists the available options. The 'required' aesthetics cannot be passed on to the params. Please note that while passing unmapped aesthetics as vectors is technically possible, the order and required length is not guaranteed to be parallel to the input data.
- When constructing a layer using a stat_*() function, the ... argument can be used to pass on parameters to the geom part of the layer. An example of this is stat_density(geom = "area", outline.type = "both"). The geom's documentation lists which parameters it can accept.
- Inversely, when constructing a layer using a geom_*() function, the ... argument can be used to pass on parameters to the stat part of the layer.
 An example of this is geom_area(stat = "density", adjust = 0.5). The stat's documentation lists which parameters it can accept.
- The key_glyph argument of layer() may also be passed on through This can be one of the functions described as key glyphs, to change the display of the layer in the legend.

Value

A data.frame with columns:

- time: time in data.
- survival: survival estimate at time.
- n.risk: number of patients at risk.

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- n. censor: number of patients censored.
- n.event: number of patients with event.

Note

Logical status is not supported.

Author(s)

Michael Sachs (in ggkm), Samer Mouksassi (in ggquickeda).

Examples

```
library(ggplot2)
sex <- rbinom(250, 1, .5)
df <- data.frame(
   time = exp(rnorm(250, mean = sex)),
   status = rbinom(250, 1, .75),
   sex = sex
)
ggplot(df, aes(time = time, status = status, color = factor(sex))) +
   stat_km() +
   stat_km_ticks()</pre>
```

stat_n_text

Add Text Indicating the Sample Size to a ggplot2 Plot

Description

[Experimental]

```
See EnvStats::stat_n_text() for details.
```

Value

A plot layer including the sample size text.

stat_prop

Compute Proportions According to Custom Denominator

Description

[Experimental]

```
See ggstats::stat_prop() for details.
```

Value

A plot layer containing the custom proportions.

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ttheme_gtdefault

Table themes

Description

[Experimental]

See ggpp::ttheme_gtdefault() for details.

Value

A list object that can be used as ttheme in the construction of tables with functions from package 'gridExtra'.

ttheme_set

Set default table theme

Description

[Experimental]

See ggpp::ttheme_set() for details.

Value

A named list with the previous value of the option.

Note

When testing this function, we found that in contrast to the original documentation, the theme is not fixed when the plot object is constructed. Instead, the option setting affects the rendering of ready built plot objects.

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