Package ‘ggtrace’

Type Package

Title Trace and Highlight Groups of Data Points

Version 0.2.0

Description Provides 'ggplot2' geoms that allow groups of data points to be outlined or highlighted for emphasis. This is particularly useful when working with dense datasets that are prone to overplotting.

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URL https://github.com/rnabioco/ggtrace

BugReports https://github.com/rnabioco/ggtrace/issues

Depends R (>= 4.0.0)

Imports ggplot2, grid, rlang

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R topics documented:

clusters ...................................................... 2
draw_key ...................................................... 2
GeomPathTrace .............................................. 3
gem_path_trace ............................................. 3
gem_point_trace ........................................... 7
stocks ......................................................... 10

Index

<table>
<thead>
<tr>
<th>clusters</th>
<th>Mock clusters</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Description

Mock clusters

Usage

clusters

draw_key

Format

A tibble with 14282 rows and 3 variables

draw_key

Key glyphs for legends

Description

Each geom has an associated function that draws the key when the geom needs to be displayed in a legend. These functions are called draw_key_*(), where * stands for the name of the respective key glyph. The key glyphs can be customized for individual geoms by providing a geom with the key_glyph argument.

Usage

draw_key_point_trace(data, params, size)

draw_key_path_trace(data, params, size)

Arguments

data A single row data frame containing the scaled aesthetics to display in this key

params A list of additional parameters supplied to the geom.

size Width and height of key in mm.
**Value**

A grid grob

**Examples**

```r
p <- ggplot2::ggplot(stocks, ggplot2::aes(day, value, color = name))

# key glyphs can be specified by their name
p + ggplot2::geom_line(key_glyph = "point_trace")

# key glyphs can be specified via their drawing function
p + ggplot2::geom_line(key_glyph = ggplot2::draw_key_rect)
```

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**Description**

GeomPathTrace

**Value**

ggproto object

**See Also**

GeomPath

---

**geom_path_trace**

Trace lines

---

**Description**

These geoms are similar to `ggplot2::geom_path()`, `ggplot2::geom_line()`, and `ggplot2::geom_step()`, but also include the ability to highlight line segments of interest. These geoms accept normal `ggplot2` graphical parameters with some modifications. `fill` controls the color of the center line, `color` controls the outline color, and `stroke` controls outline width, similar to how filled shapes are modified for other `ggplot2` geoms. Additional parameters including `size`, `alpha`, `linetype`, `linejoin`, `lineend`, and `linemitre` are also accepted.
Usage

```r
geom_path_trace(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  ...,  
  trace_position = "all",
  background_params = list(color = NA),
  lineend = "butt",
  linejoin = "round",
  linemitre = 10,
  arrow = NULL,
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE
)
```

```r
geom_line_trace(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  na.rm = FALSE,
  orientation = NA,
  show.legend = NA,
  inherit.aes = TRUE,
  trace_position = "all",
  background_params = list(color = NA),
  ...
)
```

```r
geom_step_trace(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  direction = "hv",
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE,
  trace_position = "all",
  background_params = list(color = NA),
  ...
)
```
**Arguments**

**mapping**
Set of aesthetic mappings created by `aes()` or `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, as a string.

**position**
Position adjustment, either as a string, or the result of a call to a position adjustment function.

**...**
Other arguments passed on to `layer()`. These are often aesthetics, used to set an aesthetic to a fixed value, like `colour = "red"` or `size = 3`. They may also be parameters to the paired geom/stat.

**trace_position**
Specifies which data points to outline, can be one of:
- "all" to outline every group plotted
- A predicate specifying which data points to outline. This must evaluate to `TRUE` or `FALSE` within the context of the input data. e.g. `value > 100`

**background_params**
Named list specifying aesthetic parameters to use for background data points when a predicate is passed to `trace_position`, e.g. `list(color = "red")`

**lineend**
Line end style (round, butt, square).

**linejoin**
Line join style (round, mitre, bevel).

**linemitre**
Line mitre limit (number greater than 1).

**arrow**
Arrow specification, as created by `grid::arrow()`.

**na.rm**
If `FALSE`, the default, missing values are removed with a warning. If `TRUE`, missing values are silently removed.

**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()`.

**orientation**
The orientation of the layer. The default (NA) automatically determines the orientation from the aesthetic mapping. In the rare event that this fails it can be given explicitly by setting orientation to either "x" or "y". See the `Orientation` section for more detail.

**direction**
direction of stairs: 'vh' for vertical then horizontal, 'hv' for horizontal then vertical, or 'mid' for step half-way between adjacent x-values.
Value

ggplot object

Aesthetics

geom_path_trace() understands the following aesthetics (required aesthetics are in bold):

- x
- y
- alpha
- colour
- fill
- group
- linetype
- size
- stroke

Learn more about setting these aesthetics in vignette("ggplot2-specs").

See Also

geom_path; geom_line; geom_step

Examples

# Modify line color for each group
ggplot2::ggplot(
  stocks,
  ggplot2::aes(day, value, fill = name)
) +
  geom_line_trace() +
  ggplot2::theme_minimal()

# Modify outline color for each group
ggplot2::ggplot(
  stocks,
  ggplot2::aes(day, value, color = name)
) +
  geom_line_trace() +
  ggplot2::theme_minimal()

# Specify outline color for each group
clrs <- c(
  CAC = "#E69F00",
  DAX = "#0072B2",
  FTSE = "#009E73",
  SMI = "#56B4E9"
)
geom_point_trace

```r
ggplot2::ggplot(
  stocks,
  ggplot2::aes(day, value, color = name)
) +
  geom_line_trace(stroke = 1) +
  ggplot2::scale_color_manual(values = clrs) +
  ggplot2::theme_minimal()

# Outline a subset of data points

```r
ggplot2::ggplot(
  stocks,
  ggplot2::aes(day, value, color = name)
) +
  geom_line_trace(trace_position = day > 1500, stroke = 1) +
  ggplot2::theme_minimal()

# Modify appearance of background data points

```r
ggplot2::ggplot(
  stocks,
  ggplot2::aes(day, value, color = name)
) +
  geom_line_trace(
    trace_position = day > 1500,
    background_params = list(color = NA, fill = "grey75"),
    stroke = 1
  ) +
  ggplot2::theme_minimal()

# Remove outline

```r
ggplot2::ggplot(
  stocks,
  ggplot2::aes(day, value, fill = name)
) +
  geom_line_trace(
    trace_position = day > 1500,
    background_params = list(fill = "grey75"),
    color = NA
  ) +
  ggplot2::theme_minimal()
```

**Description**

This geom is similar to `ggplot2::geom_point()`, but also includes the ability to outline points of interest. `geom_point_trace()` accepts normal `ggplot2` graphical parameters with some modifications. `fill` controls the color of each point, `color` controls the outline color, and `stroke` controls outline width, similar to how filled shapes are modified for other `ggplot2` geoms. Additional parameters including `size`, `linetype`, and `alpha` are also accepted.
Usage

domain

data

test

eval

Arguments

mapping  Set of aesthetic mappings created by aes() or 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:

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, as a string.

position  Position adjustment, either as a string, or the result of a call to a position adjustment function.

...  Other arguments passed on to layer(). These are often aesthetics, used to set an aesthetic to a fixed value, like color = "red" or size = 3. They may also be parameters to the paired geom/stat.

trace_position  Specifies which data points to outline, can be one of:

- "all" to outline every group plotted
- "bottom" to only outline the bottom layer of data points
- A predicate specifying which data points to outline. This must evaluate to TRUE or FALSE within the context of the input data. e.g. value > 100

background_params  Named list specifying aesthetic parameters to use for background data points when a predicate is passed to trace_position, e.g. list(color = "red")

na.rm  If FALSE, the default, missing values are removed with a warning. If TRUE, missing values are silently removed.
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()`.

Value

ggplot object

Aesthetics

day_point_trace() understands the following aesthetics (required aesthetics are in bold):

• x
• y
• alpha
• colour
• fill
• group
• linetype
• shape
• size
• stroke

Learn more about setting these aesthetics in `vignette("ggplot2-specs")`.

See Also

geom_point

Examples

# Modify outline color for each group
```r
ggplot2::ggplot(
groups,
  ggplot2::aes(UMAP_1, UMAP_2, color = cluster)
) +
geom_point_trace() +
ggplot2::theme_minimal()
```

# Outline a subset of points
```r
ggplot2::ggplot(
groups,
  ggplot2::aes(UMAP_1, UMAP_2, fill = cluster)
) +
geom_point_trace(trace_position = signal < 0 | signal > 17) +
```


```
ggplot2::theme_minimal()

# Modify appearance of background points
ggplot2::ggplot(
  clusters,
  ggplot2::aes(UMAP_1, UMAP_2, fill = cluster)
) +
  geom_point_trace(
    trace_position = signal < 0 | signal > 17,
    background_params = list(color = NA, fill = "grey85")
  ) +
  ggplot2::theme_minimal()
```

---

**stocks**  
*EuStockMarkets in long format*

**Description**

EuStockMarkets in long format

**Usage**

```r
stocks
```

**Format**

A tibble with 74440 rows and 3 variables
Index

* datasets
  clusters, 2
  GeomPathTrace, 3
  stocks, 10

aes(), 5,8
aes_(), 5,8

borders(), 5,9

clusters, 2

draw_key, 2
draw_key_path_trace (draw_key), 2
draw_key_point_trace (draw_key), 2

fortify(), 5,8

geom_line, 6
geom_line_trace (geom_path_trace), 3
geom_path, 6
geom_path_trace, 3
geom_point, 9
geom_point_trace, 7
geom_step, 6
geom_step_trace (geom_path_trace), 3
GeomLineTrace (GeomPathTrace), 3
GeomPath, 3
GeomPathTrace, 3
GeomPointTrace (GeomPathTrace), 3
GeomStepTrace (GeomPathTrace), 3

ggplot(), 5,8
grid::arrow(), 5

layer(), 5,8

stocks, 10