

# Package ‘ggraph’

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**Type** Package

**Title** An Implementation of Grammar of Graphics for Graphs and Networks

**Version** 2.2.1

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**Description** The grammar of graphics as implemented in ggplot2 is a poor fit for graph and network visualizations due to its reliance on tabular data input. ggraph is an extension of the ggplot2 API tailored to graph visualizations and provides the same flexible approach to building up plots layer by layer.

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**Encoding** UTF-8

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autograph

*Quickplot wrapper for networks***Description**

This function is intended to quickly show an overview of your network data. While it returns a `ggraph` object that layers etc can be added to it is limited in use and should not be used as a foundation for more complicated plots. It allows colour, labeling and sizing of nodes and edges, and the exact combination of layout and layers will depend on these as well as the features of the network. The output of this function may be fine-tuned at any release and should not be considered stable. If a plot should be reproducible it should be created manually.

**Usage**

```
autograph(graph, ...)

## Default S3 method:
autograph(
  graph,
  ...,
  node_colour = NULL,
  edge_colour = NULL,
  node_size = NULL,
  edge_width = NULL,
  node_label = NULL,
  edge_label = NULL
)
```

**Arguments**

```
graph          An object coercible to a tbl_graph
...            arguments passed on to methods
node_colour, edge_colour
                Colour mapping for nodes and edges
node_size, edge_width
                Size/width mapping for nodes and edges
node_label, edge_label
                Label mapping for nodes and edges
```

**Examples**

```
library(tidygraph)
gr <- create_notable('herschel') %>%
  mutate(class = sample(letters[1:3], n(), TRUE)) %E>%
  mutate(weight = runif(n()))

# Standard graph
autograph(gr)

# Adding node labels will cap edges
autograph(gr, node_label = class)

# Use tidygraph calls for mapping
autograph(gr, node_size = centrality_pagerank())

# Trees are plotted as dendrograms
iris_tree <- hclust(dist(iris[1:4]), method = 'euclidean'), method = 'ward.D2')
autograph(iris_tree)
```

---

`facet_edges`*Create small multiples based on edge attributes*

---

## Description

This function is equivalent to `ggplot2::facet_wrap()` but only facets edges. Nodes are repeated in every panel.

## Usage

```
facet_edges(  
  facets,  
  nrow = NULL,  
  ncol = NULL,  
  scales = "fixed",  
  shrink = TRUE,  
  labeller = "label_value",  
  as.table = TRUE,  
  switch = deprecated(),  
  drop = TRUE,  
  dir = "h",  
  strip.position = "top"  
)
```

## Arguments

<code>facets</code>	A set of variables or expressions quoted by <code>vars()</code> and defining faceting groups on the rows or columns dimension. The variables can be named (the names are passed to <code>labeller</code> ). For compatibility with the classic interface, can also be a formula or character vector. Use either a one sided formula, <code>~a + b</code> , or a character vector, <code>c("a", "b")</code> .
<code>nrow, ncol</code>	Number of rows and columns.
<code>scales</code>	Should scales be fixed ("fixed", the default), free ("free"), or free in one dimension ("free_x", "free_y")?
<code>shrink</code>	If TRUE, will shrink scales to fit output of statistics, not raw data. If FALSE, will be range of raw data before statistical summary.
<code>labeller</code>	A function that takes one data frame of labels and returns a list or data frame of character vectors. Each input column corresponds to one factor. Thus there will be more than one with <code>vars(cyl, am)</code> . Each output column gets displayed as one separate line in the strip label. This function should inherit from the "labeller" S3 class for compatibility with <code>labeller()</code> . You can use different labeling functions for different kind of labels, for example use <code>label_parsed()</code> for formatting facet labels. <code>label_value()</code> is used by default, check it for more details and pointers to other options.

<code>as.table</code>	If TRUE, the default, the facets are laid out like a table with highest values at the bottom-right. If FALSE, the facets are laid out like a plot with the highest value at the top-right.
<code>switch</code>	By default, the labels are displayed on the top and right of the plot. If "x", the top labels will be displayed to the bottom. If "y", the right-hand side labels will be displayed to the left. Can also be set to "both".
<code>drop</code>	If TRUE, the default, all factor levels not used in the data will automatically be dropped. If FALSE, all factor levels will be shown, regardless of whether or not they appear in the data.
<code>dir</code>	Direction: either "h" for horizontal, the default, or "v", for vertical.
<code>strip.position</code>	By default, the labels are displayed on the top of the plot. Using <code>strip.position</code> it is possible to place the labels on either of the four sides by setting <code>strip.position = c("top", "bottom", "left", "right")</code>

### See Also

Other ggraph-facets: [facet\\_graph\(\)](#), [facet\\_nodes\(\)](#)

### Examples

```
gr <- tidygraph::as_tbl_graph(highschool)

ggraph(gr) +
  geom_edge_link() +
  geom_node_point() +
  facet_edges(~year)
```

---

facet\_graph

*Create a grid of small multiples by node and/or edge attributes*

---

### Description

This function is equivalent to `ggplot2::facet_grid()` in that it allows for building a grid of small multiples where rows and columns correspond to a specific data value. While `ggplot2::facet_grid()` could be used it would lead to unexpected results as it is not possible to specify whether you are referring to a node or an edge attribute. Furthermore `ggplot2::facet_grid()` will draw edges in panels even though the panel does not contain both terminal nodes. `facet_graph` takes care of all of these issues, allowing you to define which data type the rows and columns are referencing as well as filtering the edges based on the nodes in each panel (even when nodes are not drawn).

### Usage

```
facet_graph(
  facets,
  row_type = "edge",
  col_type = "node",
  margins = FALSE,
```

```

scales = "fixed",
space = "fixed",
shrink = TRUE,
labeller = "label_value",
as.table = TRUE,
switch = NULL,
drop = TRUE
)

```

## Arguments

facets	<b>[Deprecated]</b> Please use rows and cols instead.
row_type, col_type	Either 'node' or 'edge'. Which data type is being faceted in the rows and columns. Default is to facet on nodes column wise and on edges row wise.
margins	Either a logical value or a character vector. Margins are additional facets which contain all the data for each of the possible values of the faceting variables. If FALSE, no additional facets are included (the default). If TRUE, margins are included for all faceting variables. If specified as a character vector, it is the names of variables for which margins are to be created.
scales	Are scales shared across all facets (the default, "fixed"), or do they vary across rows ("free_x"), columns ("free_y"), or both rows and columns ("free")?
space	If "fixed", the default, all panels have the same size. If "free_y" their height will be proportional to the length of the y scale; if "free_x" their width will be proportional to the length of the x scale; or if "free" both height and width will vary. This setting has no effect unless the appropriate scales also vary.
shrink	If TRUE, will shrink scales to fit output of statistics, not raw data. If FALSE, will be range of raw data before statistical summary.
labeller	A function that takes one data frame of labels and returns a list or data frame of character vectors. Each input column corresponds to one factor. Thus there will be more than one with <code>vars(cyl, am)</code> . Each output column gets displayed as one separate line in the strip label. This function should inherit from the "labeller" S3 class for compatibility with <code>labeller()</code> . You can use different labeling functions for different kind of labels, for example use <code>label_parsed()</code> for formatting facet labels. <code>label_value()</code> is used by default, check it for more details and pointers to other options.
as.table	If TRUE, the default, the facets are laid out like a table with highest values at the bottom-right. If FALSE, the facets are laid out like a plot with the highest value at the top-right.
switch	By default, the labels are displayed on the top and right of the plot. If "x", the top labels will be displayed to the bottom. If "y", the right-hand side labels will be displayed to the left. Can also be set to "both".
drop	If TRUE, the default, all factor levels not used in the data will automatically be dropped. If FALSE, all factor levels will be shown, regardless of whether or not they appear in the data.

**See Also**

Other ggraph-facets: [facet\\_edges\(\)](#), [facet\\_nodes\(\)](#)

**Examples**

```
library(tidygraph)
gr <- as_tbl_graph(highschool) %>%
  mutate(popularity = as.character(cut(centrality_degree(mode = 'in'),
    breaks = 3,
    labels = c('low', 'medium', 'high')
  )))
ggraph(gr) +
  geom_edge_link() +
  geom_node_point() +
  facet_graph(year ~ popularity)
```

---

facet\_nodes

*Create small multiples based on node attributes*

---

**Description**

This function is equivalent to `ggplot2::facet_wrap()` but only facets nodes. Edges are drawn if their terminal nodes are both present in a panel.

**Usage**

```
facet_nodes(
  facets,
  nrow = NULL,
  ncol = NULL,
  scales = "fixed",
  shrink = TRUE,
  labeller = "label_value",
  as.table = TRUE,
  switch = deprecated(),
  drop = TRUE,
  dir = "h",
  strip.position = "top"
)
```

**Arguments**

**facets** A set of variables or expressions quoted by `vars()` and defining faceting groups on the rows or columns dimension. The variables can be named (the names are passed to `labeller`). For compatibility with the classic interface, can also be a formula or character vector. Use either a one sided formula, `~a + b`, or a character vector, `c("a", "b")`.



nrow, ncol	Number of rows and columns.
scales	Should scales be fixed ("fixed", the default), free ("free"), or free in one dimension ("free_x", "free_y")?
shrink	If TRUE, will shrink scales to fit output of statistics, not raw data. If FALSE, will be range of raw data before statistical summary.
labeller	A function that takes one data frame of labels and returns a list or data frame of character vectors. Each input column corresponds to one factor. Thus there will be more than one with <code>vars(cyl, am)</code> . Each output column gets displayed as one separate line in the strip label. This function should inherit from the "labeller" S3 class for compatibility with <code>labeller()</code> . You can use different labeling functions for different kind of labels, for example use <code>label_parsed()</code> for formatting facet labels. <code>label_value()</code> is used by default, check it for more details and pointers to other options.
as.table	If TRUE, the default, the facets are laid out like a table with highest values at the bottom-right. If FALSE, the facets are laid out like a plot with the highest value at the top-right.
switch	By default, the labels are displayed on the top and right of the plot. If "x", the top labels will be displayed to the bottom. If "y", the right-hand side labels will be displayed to the left. Can also be set to "both".
drop	If TRUE, the default, all factor levels not used in the data will automatically be dropped. If FALSE, all factor levels will be shown, regardless of whether or not they appear in the data.
dir	Direction: either "h" for horizontal, the default, or "v", for vertical.
strip.position	By default, the labels are displayed on the top of the plot. Using <code>strip.position</code> it is possible to place the labels on either of the four sides by setting <code>strip.position = c("top", "bottom", "left", "right")</code>

### See Also

Other `ggraph`-facets: [facet\\_edges\(\)](#), [facet\\_graph\(\)](#)

### Examples

```
library(tidygraph)
gr <- as_tbl_graph(highschool) %>%
  mutate(popularity = as.character(cut(centrality_degree(mode = 'in'),
    breaks = 3,
    labels = c('low', 'medium', 'high')
  )))
ggraph(gr) +
  geom_edge_link() +
  geom_node_point() +
  facet_nodes(~popularity)
```

---

flare

*The class hierarchy of the flare visualization library*

---

### Description

This dataset contains the graph that describes the class hierarchy for the **Flare** ActionScript visualization library. It contains both the class hierarchy as well as the import connections between classes. This dataset has been used extensively in the D3.js documentation and examples and are included here to make it easy to redo the examples in ggraph.

### Usage

flare

### Format

A list of three data.frames describing the software structure of flare:

**edges** This data.frame maps the hierarchical structure of the class hierarchy as an edgelist, with the class in from being the superclass of the class in to.

**vertices** This data.frame gives additional information on the classes. It contains the full name, size and short name of each class.

**imports** This data.frame contains the class imports for each class implementation. The from column gives the importing class and the to column gives the import.

### Source

The data have been adapted from the JSON downloaded from <https://gist.github.com/mbostock/1044242#file-readme-flare-imports-json> courtesy of Mike Bostock. The Flare framework is the work of the **UC Berkeley Visualization Lab**.

---

geometry

*Define simple shapes for line capping*

---

### Description

This set of functions makes it easy to define shapes at the terminal points of edges that are used to shorten the edges. The shapes themselves are not drawn, but the edges will end at the boundary of the shape rather than at the node position. This is especially relevant when drawing arrows at the edges as the arrows will be partly obscured by the node unless the edge is shortened. Edge shortening is dynamic and will update as the plot is resized, making sure that the capping remains at an absolute distance to the end point.

**Usage**

```

geometry(
  type = "circle",
  width = 1,
  height = width,
  width_unit = "cm",
  height_unit = width_unit
)

circle(radius = 1, unit = "cm")

square(length = 1, unit = "cm")

ellipsis(a = 1, b = 1, a_unit = "cm", b_unit = a_unit)

rectangle(width = 1, height = 1, width_unit = "cm", height_unit = width_unit)

label_rect(label, padding = margin(1, 1, 1.5, 1, "mm"), ...)

is.geometry(x)

```

**Arguments**

type	The type of geometry to use. Currently 'circle' and 'rect' is supported.
width, height, length, radius, a, b	The dimensions of the shape.
unit, width_unit, height_unit, a_unit, b_unit	The unit for the numbers given.
label	The text to be enclosed
padding	extra size to be added around the text using the <code>ggplot2::margin()</code> function
...	Passed on to <code>grid::gpar()</code>
x	An object to test for geometry inheritance

**Details**

geometry is the base constructor, while the rest are helpers to save typing. circle creates circles with a given radius, square creates squares at a given side length, ellipsis creates ellipses with given a and b values (width and height radii), and rectangle makes rectangles of a given width and height. label\_rect is a helper that, given a list of strings and potentially formatting options creates a rectangle that encloses the string.

**Value**

A geometry object encoding the specified shape.

**Examples**

```

geometry(c('circle', 'rect', 'rect'), 1:3, 3:1)

circle(1:4, 'mm')

label_rect(c('some', 'different', 'words'), fontsize = 18)

```

---

geom\_axis\_hive

*Draw rectangular bars and labels on hive axes*


---

**Description**

This function lets you annotate the axes in a hive plot with labels and color coded bars.

**Usage**

```

geom_axis_hive(
  mapping = NULL,
  data = NULL,
  position = "identity",
  label = TRUE,
  axis = TRUE,
  show.legend = NA,
  ...
)

```

**Arguments**

mapping	Set of aesthetic mappings created by <a href="#">aes()</a> . If specified and <code>inherit.aes = TRUE</code> (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 <code>NULL</code> , the default, the data is inherited from the plot data as specified in the call to <a href="#">ggplot()</a> . A <code>data.frame</code> , or other object, will override the plot data. All objects will be fortified to produce a data frame. See <a href="#">fortify()</a> for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <a href="#">position_jitter</a> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
label	Should the axes be labelled. Defaults to <code>TRUE</code>
axis	Should a rectangle be drawn along the axis. Defaults to <code>TRUE</code>

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.

...                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.

## Aesthetics

`geom_axis_hive` understand the following aesthetics.

- alpha
- colour
- fill
- size
- linetype
- label\_size
- family
- fontface
- lineheight

## Author(s)

Thomas Lin Pedersen

## Examples

```
# Plot the flare import graph as a hive plot
library(tidygraph)
flareGr <- as_tbl_graph(flare$imports) %>%
  mutate(
    type = dplyr::case_when(
      centrality_degree(mode = 'in') == 0 ~ 'Source',
      centrality_degree(mode = 'out') == 0 ~ 'Sink',
      TRUE ~ 'Both'
    )
  ) %>%
  activate(edges) %>%
  mutate(
    type = dplyr::case_when(
      grepl('flare.analytics', paste(.N()$name[from], .N()$name[to])) ~ 'Analytics',
      TRUE ~ 'Other'
    )
  )
ggraph(flareGr, 'hive', axis = type) +
  geom_edge_hive(aes(colour = type), edge_alpha = 0.1) +
  geom_axis_hive(aes(colour = type)) +
  coord_fixed()
```

---

`geom_conn_bundle`*Create hierarchical edge bundles between node connections*

---

### Description

Hierarchical edge bundling is a technique to introduce some order into the hairball structure that can appear when there's a lot of overplotting and edge crossing in a network plot. The concept requires that the network has an intrinsic hierarchical structure that defines the layout but is not shown. Connections between points (that is, not edges) are then drawn so that they loosely follow the underlying hierarchical structure. This results in a flow-like structure where lines that partly move in the same direction will be bundled together.

### Usage

```
geom_conn_bundle(  
  mapping = NULL,  
  data = get_con(),  
  position = "identity",  
  arrow = NULL,  
  lineend = "butt",  
  show.legend = NA,  
  n = 100,  
  tension = 0.8,  
  ...  
)
```

```
geom_conn_bundle2(  
  mapping = NULL,  
  data = get_con(),  
  position = "identity",  
  arrow = NULL,  
  lineend = "butt",  
  show.legend = NA,  
  n = 100,  
  tension = 0.8,  
  ...  
)
```

```
geom_conn_bundle0(  
  mapping = NULL,  
  data = get_con(),  
  position = "identity",  
  arrow = NULL,  
  lineend = "butt",  
  show.legend = NA,  
  tension = 0.8,  
  ...  
)
```

)

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The result of a call to <code>get_con()</code>
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
lineend	Line end style (round, butt, square).
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.
n	The number of points to create along the path.
tension	How "loose" should the bundles be. 1 will give very tight bundles, while 0 will turn of bundling completely and give straight lines. Defaults to 0.8
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

**Aesthetics**

`geom_conn_bundle*` understands the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- **circular**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

**Computed variables**

**index** The position along the path (not computed for the \*0 version)

**Note**

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

**Author(s)**

Thomas Lin Pedersen

**References**

Holten, D. (2006). *Hierarchical edge bundles: visualization of adjacency relations in hierarchical data*. IEEE Transactions on Visualization and Computer Graphics, **12**(5), 741-748. doi:10.1109/TVCG.2006.147

**Examples**

```
# Create a graph of the flare class system
library(tidygraph)
flareGraph <- tbl_graph(flare$vertices, flare$edges) %>%
  mutate(
    class = map_bfs_chr(node_is_root(), .f = function(node, dist, path, ...) {
      if (dist <= 1) {
        return(shortName[node])
      }
      path$result[[nrow(path)]]
    })
  )
importFrom <- match(flare$imports$from, flare$vertices$name)
importTo <- match(flare$imports$to, flare$vertices$name)

# Use class inheritance for layout but plot class imports as bundles
ggraph(flareGraph, 'dendrogram', circular = TRUE) +
  geom_conn_bundle(aes(colour = after_stat(index)),
    data = get_con(importFrom, importTo),
    edge_alpha = 0.25
  ) +
  geom_node_point(aes(filter = leaf, colour = class)) +
  scale_edge_colour_distiller('', direction = 1, guide = 'edge_direction') +
  coord_fixed() +
  ggforce::theme_no_axes()
```



## Description

This geom is mainly intended for arc linear and circular diagrams (i.e. used together with `layout_tbl_graph_linear()`), though it can be used elsewhere. It draws edges as arcs with a height proportional to the distance between the nodes. Arcs are calculated as beziers. For linear layout the placement of control points are related to the curvature argument and the distance between the two nodes. For circular layout the control points are placed on the same angle as the start and end node at a distance related to the distance between the nodes.

## Usage

```
geom_edge_arc(  
  mapping = NULL,  
  data = get_edges(),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  n = 100,  
  fold = FALSE,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...,  
  curvature  
)  
  
geom_edge_arc2(  
  mapping = NULL,  
  data = get_edges("long"),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  n = 100,  
  fold = FALSE,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,
```

```

    check_overlap = FALSE,
    angle_calc = "rot",
    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...,
    curvature
)

geom_edge_arc0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  strength = 1,
  lineend = "butt",
  show.legend = NA,
  fold = fold,
  ...,
  curvature
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
strength	The bend of the curve. 1 approximates a halfcircle while 0 will give a straight line. Negative number will change the direction of the curve. Only used if <code>circular = FALSE</code> .
n	The number of points to create along the path.
fold	Logical. Should arcs appear on the same side of the nodes despite different directions. Default to FALSE.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.

label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.
curvature	Deprecated. Use <code>strength</code> instead.

## Aesthetics

`geom_edge_arc` and `geom_edge_arc0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- **circular**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_arc2` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**

- **group**
- **circular**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_arc and geom\_edge\_arc2 furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Edge variants

Many geom\_edge\_\* layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. colour = after\_stat(index). The version postfixed with a "2" uses the "long" edge format (see [get\\_edges\(\)](#)) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerably less performant so should only be used if this is needed. The version postfixed with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. geom\_edge\_\* and geom\_edge\_\*2 supports this through the start\_cap and end\_cap aesthetics that takes a [geometry\(\)](#) specification and dynamically caps the termini of the edges based on the given specifications. This means that if end\_cap = circle(1, 'cm') the edges will end at a distance of 1cm even during resizing of the plot window.

All `geom_edge_*` and `geom_edge_*2` have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal\(\)](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
# Make a graph with different directions of edges
gr <- create_notable('Meredith') %>%
  convert(to_directed) %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE)) %>%
  activate(edges) %>%
  mutate(
    class = sample(letters[1:3], n(), replace = TRUE),
    switch = sample(c(TRUE, FALSE), n(), replace = TRUE)
  ) %>%
  reroute(from = to, to = from, subset = switch)

ggraph(gr, 'linear') +
  geom_edge_arc(aes(alpha = after_stat(index)))

ggraph(gr, 'linear') +
  geom_edge_arc2(aes(colour = node.class), strength = 0.6)

ggraph(gr, 'linear', circular = TRUE) +
  geom_edge_arc0(aes(colour = class))
```

---

geom\_edge\_bend      *Draw edges as diagonals*

---

### Description

This geom draws edges as cubic bezier curves with the control points positioned along the elbow edge. It has the appearance of a softened elbow edge with the hard angle substituted by a tapered bend.

### Usage

```
geom_edge_bend(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  strength = 1,
  flipped = FALSE,
  n = 100,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  label_colour = "black",
  label_alpha = 1,
  label_parse = FALSE,
  check_overlap = FALSE,
  angle_calc = "rot",
  force_flip = TRUE,
  label_dodge = NULL,
  label_push = NULL,
  show.legend = NA,
  ...
)

geom_edge_bend2(
  mapping = NULL,
  data = get_edges("long"),
  position = "identity",
  arrow = NULL,
  strength = 1,
  flipped = FALSE,
  n = 100,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  label_colour = "black",
  label_alpha = 1,
```

```

    label_parse = FALSE,
    check_overlap = FALSE,
    angle_calc = "rot",
    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...
  )

geom_edge_bend0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  strength = 1,
  flipped = FALSE,
  lineend = "butt",
  show.legend = NA,
  ...
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
strength	The strength of the curvature of the bend. 0 will result in a straight line while 1 will give a strong arc.
flipped	Logical, Has the layout been flipped by reassigning the mapping of x, y etc?
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .

check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. check_overlap happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling geom_text(). Note that this argument is not supported by geom_label().
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If angle_calc is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of angle_calc is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of angle_calc is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Aesthetics

geom\_edge\_bend and geom\_edge\_bend0 understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- **circular**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_bend2 understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- **circular**
- edge\_colour



- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_bend and geom\_edge\_bend2 furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Edge variants

Many geom\_edge\_\* layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. `colour = after_stat(index)`. The version postfixed with a "2" uses the "long" edge format (see [get\\_edges\(\)](#)) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfixed with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. geom\_edge\_\* and geom\_edge\_\*2 supports this through the start\_cap and end\_cap aesthetics that takes a [geometry\(\)](#) specification and dynamically caps the termini of the edges based on the given specifications. This means that if `end_cap = circle(1, 'cm')` the edges will end at a distance of 1cm even during resizing of the plot window.

All geom\_edge\_\* and geom\_edge\_\*2 have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The label\_pos can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The label\_size aesthetic can be used to control the size of the label. Often it is needed to have the

label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal\(\)](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
gr <- create_tree(20, 4) %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE)) %>%
  activate(edges) %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE))

ggraph(gr, 'tree') +
  geom_edge_bend(aes(alpha = after_stat(index)))

ggraph(gr, 'tree') +
  geom_edge_bend2(aes(colour = node.class))

ggraph(gr, 'tree') +
  geom_edge_bend0(aes(colour = class))
```

---

geom\_edge\_bundle\_force

*Bundle edges using force directed edge bundling*

---

### Description

This geom performs force directed edge bundling to reduce visual clutter. It uses a self-organizing approach to bundling in which edges are modeled as flexible springs that can attract each other without the need of a hierarchy. Be aware that this bundling technique works globally and thus may bundle edges that is otherwise unrelated together. Care should be taken when interpreting the

resulting visual. An alternative approach to edge bundling that uses the graph topology is provided by `geom_edge_bundle_path()`.

### Usage

```
geom_edge_bundle_force(  
  mapping = NULL,  
  data = get_edges(),  
  position = "identity",  
  arrow = NULL,  
  n = 100,  
  force = 1,  
  n_cycle = 6,  
  cuts_start = 1,  
  step = 0.04,  
  cuts_new = 2,  
  n_iter = 50,  
  iter_new = 2/3,  
  threshold = 0.6,  
  eps = 1e-08,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...  
)  
  
geom_edge_bundle_force2(  
  mapping = NULL,  
  data = get_edges("long"),  
  position = "identity",  
  arrow = NULL,  
  n = 100,  
  force = 1,  
  n_cycle = 6,  
  cuts_start = 1,  
  step = 0.04,  
  cuts_new = 2,  
  n_iter = 50,  
  iter_new = 2/3,  
  threshold = 0.6,
```

```

    eps = 1e-08,
    lineend = "butt",
    linejoin = "round",
    linemitre = 1,
    label_colour = "black",
    label_alpha = 1,
    label_parse = FALSE,
    check_overlap = FALSE,
    angle_calc = "rot",
    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...
  )

geom_edge_bundle_force0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  force = 1,
  n_cycle = 6,
  cuts_start = 1,
  step = 0.04,
  cuts_new = 2,
  n_iter = 50,
  iter_new = 2/3,
  threshold = 0.6,
  eps = 1e-08,
  lineend = "butt",
  show.legend = NA,
  ...
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
n	The number of points to create along the path.

force	The spring force during bundling
n_cycle	number of iteration cycles
cuts_start	initial number of edge divisions
step	initial step size
cuts_new	factor for how many new division points to add after a cycle
n_iter	number of iteration steps per cycle
iter_new	factor of how to decrease the number of iterations per cycle
threshold	threshold for considering two edges to be interacting
eps	tolerance
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <a href="#">grDevices::plotmath()</a> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <a href="#">grid::unit()</a> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
label_push	A <a href="#">grid::unit()</a> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
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.
...	Other arguments passed on to <a href="#">layer()</a> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Aesthetics

`geom_edge_bundle_force` and `geom_edge_bundle_force0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**

- **y**
- **xend**
- **yend**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_bundle_force2` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_bundle_force` and `geom_edge_bundle_force2` furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

## Edge variants

Many `geom_edge_*` layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. `colour = after_stat(index)`. The version postfixed with a "2" uses the "long" edge format (see `get_edges()`) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfixed with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. `geom_edge_*` and `geom_edge_*2` supports this through the `start_cap` and `end_cap` aesthetics that takes a `geometry()` specification and dynamically caps the termini of the edges based on the given specifications. This means that if `end_cap = circle(1, 'cm')` the edges will end at a distance of 1cm even during resizing of the plot window.

All `geom_edge_*` and `geom_edge_*2` have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

## Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

## Author(s)

David Schoch

## References

Holten, D. and Wijk, J.J.V. (2009). *Force-Directed Edge Bundling for Graph Visualization*. Computer Graphics Forum (Blackwell Publishing Ltd) 28, no. 3: 983-990. <https://doi.org/10.1111/j.1467-8659.2009.01450.x>

## See Also

Other `geom_edge_*`: `geom_edge_arc()`, `geom_edge_bend()`, `geom_edge_bundle_minimal()`, `geom_edge_bundle_path()`, `geom_edge_density()`, `geom_edge_diagonal()`, `geom_edge_elbow()`, `geom_edge_fan()`, `geom_edge_hive()`, `geom_edge_link()`, `geom_edge_loop()`, `geom_edge_parallel()`, `geom_edge_point()`, `geom_edge_sf()`, `geom_edge_span()`, `geom_edge_tile()`

**Examples**

```
# (not necessarily an insightful use)
ggraph(highschool) +
  geom_edge_bundle_force(n_cycle = 2, threshold = 0.4)
```

---

```
geom_edge_bundle_minimal
```

*Bundle edges along the minimal spanning tree*

---

**Description**

This geom performs edge bundling by letting edges follow the shortest path along the minimal spanning tree of the graph. Due to its simplicity it is very fast but does enforce a tree-like appearance to the bundling. Adjusting the `max_distortion` and `tension` parameters may alleviate this to some extent.

**Usage**

```
geom_edge_bundle_minimal(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  n = 100,
  max_distortion = 2,
  weight_fac = 2,
  tension = 1,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  label_colour = "black",
  label_alpha = 1,
  label_parse = FALSE,
  check_overlap = FALSE,
  angle_calc = "rot",
  force_flip = TRUE,
  label_dodge = NULL,
  label_push = NULL,
  show.legend = NA,
  ...
)

geom_edge_bundle_minimal2(
  mapping = NULL,
  data = get_edges("long"),
  position = "identity",
```



```

    arrow = NULL,
    n = 100,
    max_distortion = 2,
    weight_fac = 2,
    tension = 1,
    lineend = "butt",
    linejoin = "round",
    linemitre = 1,
    label_colour = "black",
    label_alpha = 1,
    label_parse = FALSE,
    check_overlap = FALSE,
    angle_calc = "rot",
    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...
)

geom_edge_bundle_minimal0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  max_distortion = 2,
  weight_fac = 2,
  tension = 1,
  lineend = "butt",
  show.legend = NA,
  ...
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
n	The number of points to create along the path.
max_distortion	A multiplication factor to determine the maximum allowed distortion of the path

	during bundling. If the new edge is longer than <code>max_distortion</code> times the old length it is rejected.
<code>weight_fac</code>	The exponent used to assign weights to the graph when calculating the shortest path. The final weights are given as <code>edge_length ^ weight_fac</code> meaning that shorter edges are prioritised when calculating the weights.
<code>tension</code>	A loosening factor when calculating the b-spline of the edge based on the shortest path. Will move control points closer and closer to the direct line as it approaches 0
<code>lineend</code>	Line end style (round, butt, square).
<code>linejoin</code>	Line join style (round, mitre, bevel).
<code>linemitre</code>	Line mitre limit (number greater than 1).
<code>label_colour</code>	The colour of the edge label. If NA it will use the colour of the edge.
<code>label_alpha</code>	The opacity of the edge label. If NA it will use the opacity of the edge.
<code>label_parse</code>	If TRUE, the labels will be parsed into expressions and displayed as described in <a href="#">grDevices::plotmath()</a> .
<code>check_overlap</code>	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
<code>angle_calc</code>	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
<code>force_flip</code>	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
<code>label_dodge</code>	A <a href="#">grid::unit()</a> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
<code>label_push</code>	A <a href="#">grid::unit()</a> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
<code>show.legend</code>	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.
...	Other arguments passed on to <a href="#">layer()</a> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

## Aesthetics

`geom_edge_force_minimal` and `geom_edge_force_minimal0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**

- **edge\_id** (should not be overwritten)
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

*geom\_edge\_force\_minimal2* understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- **edge\_id** (should not be overwritten)
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

*geom\_edge\_force\_minimal* and *geom\_edge\_force\_minimal2* furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### **Computed variables**

**index** The position along the path (not computed for the \*0 version)

## Edge variants

Many `geom_edge_*` layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. `colour = after_stat(index)`. The version postfix with a "2" uses the "long" edge format (see `get_edges()`) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfix with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. `geom_edge_*` and `geom_edge_*2` supports this through the `start_cap` and `end_cap` aesthetics that takes a `geometry()` specification and dynamically caps the termini of the edges based on the given specifications. This means that if `end_cap = circle(1, 'cm')` the edges will end at a distance of 1cm even during resizing of the plot window.

All `geom_edge_*` and `geom_edge_*2` have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

## Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

## Author(s)

Thomas Lin Pedersen

## See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

## Examples

```
ggraph(highschool) +
  geom_edge_bundle_minimal()
```

```
# Allow more edges to bundle
ggraph(highschool) +
  geom_edge_bundle_minimal(max_distortion = 5, tension = 0.9)
```

---

geom\_edge\_bundle\_path *Bundle edges using edge path bundling*

---

## Description

This geom performs edge bundling using the edge path algorithm. This approach uses the underlying graph structure to find shortest paths for each edge in a graph that is gradually removed of its edges. Since it is based on the topology of the graph it should lead to less spurious bundling of unrelated edges compared to `geom_edge_bundle_force()` and also has a simpler parameter space.

## Usage

```
geom_edge_bundle_path(  
  mapping = NULL,  
  data = get_edges(),  
  position = "identity",  
  arrow = NULL,  
  n = 100,  
  directed = NULL,  
  max_distortion = 2,  
  weight_fac = 2,  
  tension = 1,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...  
)  
  
geom_edge_bundle_path2(  
  mapping = NULL,  
  data = get_edges("long"),  
  position = "identity",  
  arrow = NULL,
```

```

n = 100,
directed = NULL,
max_distortion = 2,
weight_fac = 2,
tension = 1,
lineend = "butt",
linejoin = "round",
linemitre = 1,
label_colour = "black",
label_alpha = 1,
label_parse = FALSE,
check_overlap = FALSE,
angle_calc = "rot",
force_flip = TRUE,
label_dodge = NULL,
label_push = NULL,
show.legend = NA,
...
)

geom_edge_bundle_path0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  directed = NULL,
  max_distortion = 2,
  weight_fac = 2,
  tension = 1,
  lineend = "butt",
  show.legend = NA,
  ...
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
n	The number of points to create along the path.

directed	Logical. Should the shortest paths be calculated using direction information of the graph. Setting this to TRUE can help split up bundles that flows in opposite directions. Ignored for undirected graphs
max_distortion	A multiplication factor to determine the maximum allowed distortion of the path during bundling. If the new edge is longer than max_distortion times the old length it is rejected.
weight_fac	The exponent used to assign weights to the graph when calculating the shortest path. The final weights are given as $\text{edge\_length}^{\text{weight\_fac}}$ meaning that shorter edges are prioritised when calculating the weights.
tension	A loosening factor when calculating the b-spline of the edge based on the shortest path. Will move control points closer and closer to the direct line as it approaches 0
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. check_overlap happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If angle_calc is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of angle_calc is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of angle_calc is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Aesthetics

`geom_edge_force_path` and `geom_edge_force_path0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- **edge\_id** (should not be overwritten)
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_force_path2` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- **edge\_id** (should not be overwritten)
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_force_path` and `geom_edge_force_path2` furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)



## Edge variants

Many `geom_edge_*` layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. `colour = after_stat(index)`. The version postfix with a "2" uses the "long" edge format (see `get_edges()`) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfix with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. `geom_edge_*` and `geom_edge_*2` supports this through the `start_cap` and `end_cap` aesthetics that takes a `geometry()` specification and dynamically caps the termini of the edges based on the given specifications. This means that if `end_cap = circle(1, 'cm')` the edges will end at a distance of 1cm even during resizing of the plot window.

All `geom_edge_*` and `geom_edge_*2` have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

## Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

## Author(s)

Thomas Lin Pedersen and David Schoch

## References

Wallinger, M., Archambault, D., Auber, D., Nöllenburg, M., and Peltonen, J. (2022). *Edge-Path Bundling: A Less Ambiguous Edge Bundling Approach*. IEEE Transactions on Visualization and Computer Graphics 28(1) 313-323. <https://doi.org/10.1109/TVCG.2021.3114795>

## See Also

Other `geom_edge_*`: `geom_edge_arc()`, `geom_edge_bend()`, `geom_edge_bundle_force()`, `geom_edge_bundle_minimal`, `geom_edge_density()`, `geom_edge_diagonal()`, `geom_edge_elbow()`, `geom_edge_fan()`, `geom_edge_hive()`, `geom_edge_link()`, `geom_edge_loop()`, `geom_edge_parallel()`, `geom_edge_point()`, `geom_edge_sf()`, `geom_edge_span()`, `geom_edge_tile()`

**Examples**

```

ggraph(highschool) +
  geom_edge_bundle_path()

# Use tension to lessen the effect
ggraph(highschool) +
  geom_edge_bundle_path(tension = 0.8)

```

---

geom\_edge\_density      *Show edges as a density map*

---

**Description**

This geom makes it possible to add a layer showing edge presence as a density map. Each edge is converted to  $n$  points along the line and a jitter is applied. Based on this dataset a two-dimensional kernel density estimation is applied and plotted as a raster image. The density is mapped to the alpha level, making it possible to map a variable to the fill.

**Usage**

```

geom_edge_density(
  mapping = NULL,
  data = get_edges("short"),
  position = "identity",
  show.legend = NA,
  n = 100,
  ...
)

```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default <code>x</code> , <code>y</code> , <code>xend</code> , <code>yend</code> , <code>group</code> and <code>circular</code> are mapped to <code>x</code> , <code>y</code> , <code>xend</code> , <code>yend</code> , <code>edge.id</code> and <code>circular</code> in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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.
n	The number of points to estimate in the <code>x</code> and <code>y</code> direction, i.e. the resolution of the raster.

... 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.

### Aesthetics

`geom_edge_density` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

**x y xend yend** `edge_fill` filter

### Computed variables

**x, y** The coordinates for each pixel in the raster

**density** The density associated with the pixel

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
gr <- create_notable('bull') %>%
  activate(edges) %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE))

ggraph(gr, 'stress') +
  geom_edge_density(aes(fill = class)) +
  geom_edge_link() + geom_node_point()
```

---

geom\_edge\_diagonal      *Draw edges as diagonals*

---

### Description

This geom draws edges as diagonal bezier curves. The name comes from D3.js where this shape was called diagonals until it was renamed to [links](#). A diagonal in this context is a quadratic bezier with the control points positioned halfway between the start and end points but on the same axis. This produces a pleasing fan-in, fan-out line that is mostly relevant for hierarchical layouts as it implies an overall directionality in the plot.

### Usage

```
geom_edge_diagonal(  
  mapping = NULL,  
  data = get_edges(),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  flipped = FALSE,  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...  
)  
  
geom_edge_diagonal2(  
  mapping = NULL,  
  data = get_edges("long"),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  flipped = FALSE,  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  ...  
)
```

```

    label_colour = "black",
    label_alpha = 1,
    label_parse = FALSE,
    check_overlap = FALSE,
    angle_calc = "rot",
    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...
)

geom_edge_diagonal0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  strength = 1,
  flipped = FALSE,
  lineend = "butt",
  show.legend = NA,
  ...
)

```

## Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
strength	The strength of the curvature of the diagonal. 0 will result in a straight line while 1 will give the familiar S-shape.
flipped	Logical, Has the layout been flipped by reassigning the mapping of x, y etc?
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.

label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Aesthetics

`geom_edge_diagonal` and `geom_edge_diagonal0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- **circular**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_diagonal2` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**

- **circular**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_diagonal and geom\_edge\_diagonal2 furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Edge variants

Many geom\_edge\_\* layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. colour = after\_stat(index). The version postfix with a "2" uses the "long" edge format (see [get\\_edges\(\)](#)) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfix with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. geom\_edge\_\* and geom\_edge\_\*2 supports this through the start\_cap and end\_cap aesthetics that takes a [geometry\(\)](#) specification and dynamically caps the termini of the edges based on the given specifications. This means that if end\_cap = circle(1, 'cm') the edges will end at a distance of 1cm even during resizing of the plot window.

All geom\_edge\_\* and geom\_edge\_\*2 have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs

to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
gr <- create_tree(20, 4) %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE)) %>%
  activate(edges) %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE))

ggraph(gr, 'tree') +
  geom_edge_diagonal(aes(alpha = after_stat(index)))

ggraph(gr, 'tree') +
  geom_edge_diagonal2(aes(colour = node.class))

ggraph(gr, 'tree') +
  geom_edge_diagonal0(aes(colour = class))
```



## Description

This geom draws edges as an angle in the same manner as known from classic dendrogram plots of hierarchical clustering results. In case a circular transformation has been applied the first line segment will be drawn as an arc as expected. This geom is only applicable to layouts that return a direction for the edges (currently `layout_tbl_graph_dendrogram()`, `layout_tbl_graph_partition()` and `layout_tbl_graph_igraph()` with the "tree" algorithm).

## Usage

```
geom_edge_elbow(  
  mapping = NULL,  
  data = get_edges(),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  flipped = FALSE,  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...  
)  
  
geom_edge_elbow2(  
  mapping = NULL,  
  data = get_edges("long"),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  flipped = FALSE,  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",
```

```

    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...
  )

geom_edge_elbow(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  flipped = FALSE,
  lineend = "butt",
  show.legend = NA,
  ...
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
strength	How bend the elbow should be. 1 will give a right angle, while 0 will give a straight line. Ignored for circular layouts
flipped	Logical, Has the layout been flipped by reassigning the mapping of x, y etc?
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .

angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If angle_calc is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of angle_calc is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of angle_calc is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

## Aesthetics

geom\_edge\_elbow and geom\_edge\_elbow0 understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- **circular**
- **direction**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_elbow2 understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- **circular**
- **direction**
- edge\_colour
- edge\_width

- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_elbow and geom\_edge\_elbow2 furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Edge variants

Many geom\_edge\_\* layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. colour = after\_stat(index). The version postfix with a "2" uses the "long" edge format (see [get\\_edges\(\)](#)) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfix with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. geom\_edge\_\* and geom\_edge\_\*2 supports this through the start\_cap and end\_cap aesthetics that takes a [geometry\(\)](#) specification and dynamically caps the termini of the edges based on the given specifications. This means that if end\_cap = circle(1, 'cm') the edges will end at a distance of 1cm even during resizing of the plot window.

All geom\_edge\_\* and geom\_edge\_\*2 have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The label\_pos can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The label\_size aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot

dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
irisDen <- hclust(dist(iris[1:4], method = 'euclidean'), method = 'ward.D2') %>%
  as_tbl_graph() %>%
  mutate(class = sample(letters[1:3], n(), TRUE)) %>%
  activate(edges) %>%
  mutate(class = sample(letters[1:3], n(), TRUE))

ggraph(irisDen, 'dendrogram', circular = TRUE) +
  geom_edge_elbow(aes(alpha = after_stat(index)))

ggraph(irisDen, 'dendrogram') +
  geom_edge_elbow2(aes(colour = node.class))

ggraph(irisDen, 'dendrogram', height = height) +
  geom_edge_elbow0(aes(colour = class))
```

---

geom\_edge\_fan

*Draw edges as curves of different curvature*

---

### Description

This geom draws edges as cubic beziers with the control point positioned half-way between the nodes and at an angle dependent on the presence of parallel edges. This results in parallel edges being drawn in a non-overlapping fashion resembling the standard approach used in [igraph::plot.igraph\(\)](#). Before calculating the curvature the edges are sorted by direction so that edges going the same way will be adjacent. This geom is currently the only choice for non-simple graphs if edges should not be overplotted.

**Usage**

```
geom_edge_fan(  
  mapping = NULL,  
  data = get_edges(),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...,  
  spread  
)  
  
geom_edge_fan2(  
  mapping = NULL,  
  data = get_edges("long"),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...,  
  spread  
)  
  
geom_edge_fan0(  
  mapping = NULL,  
  data = get_edges(),  
  position = "identity",  
  arrow = NULL,  
  strength = 1,  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...,  
  spread  
)
```

```

mapping = NULL,
data = get_edges(),
position = "identity",
arrow = NULL,
strength = 1,
lineend = "butt",
show.legend = NA,
...,
spread
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
strength	Modify the width of the fans <code>strength &gt; 1</code> will create wider fans while the reverse will make them more narrow.
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'

label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.
spread	Deprecated. Use <code>strength</code> instead.

### Aesthetics

`geom_edge_fan` and `geom_edge_fan0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- **from**
- **to**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_fan2` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- **from**
- **to**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_fan` and `geom_edge_fan2` furthermore takes the following aesthetics.

- start\_cap



- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Edge variants

Many `geom_edge_*` layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. `colour = after_stat(index)`. The version postfix with a "2" uses the "long" edge format (see `get_edges()`) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfix with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. `geom_edge_*` and `geom_edge_*2` supports this through the `start_cap` and `end_cap` aesthetics that takes a `geometry()` specification and dynamically caps the termini of the edges based on the given specifications. This means that if `end_cap = circle(1, 'cm')` the edges will end at a distance of 1cm even during resizing of the plot window.

All `geom_edge_*` and `geom_edge_*2` have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

**Author(s)**

Thomas Lin Pedersen

**See Also**

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

**Examples**

```
require(tidygraph)
gr <- create_notable('bull') %>%
  convert(to_directed) %>%
  bind_edges(data.frame(from = c(1, 2, 2, 3), to = c(2, 1, 3, 2))) %E>%
  mutate(class = sample(letters[1:3], 9, TRUE)) %N>%
  mutate(class = sample(c('x', 'y'), 5, TRUE))

ggraph(gr, 'stress') +
  geom_edge_fan(aes(alpha = after_stat(index)))

gggraph(gr, 'stress') +
  geom_edge_fan2(aes(colour = node.class))

gggraph(gr, 'stress') +
  geom_edge_fan0(aes(colour = class))
```

---

 geom\_edge\_hive

---

*Draw edges in hive plots*


---

**Description**

This geom is only intended for use together with the hive layout. It draws edges between nodes as bezier curves, with the control points positioned at the same radii as the start or end point, and at a distance defined by the curvature argument.

**Usage**

```
geom_edge_hive(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  strength = 1,
  n = 100,
  lineend = "butt",
  linejoin = "round",
```

```
    linemitre = 1,
    label_colour = "black",
    label_alpha = 1,
    label_parse = FALSE,
    check_overlap = FALSE,
    angle_calc = "rot",
    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...,
    curvature
)

geom_edge_hive2(
  mapping = NULL,
  data = get_edges("long"),
  position = "identity",
  arrow = NULL,
  strength = 1,
  n = 100,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  label_colour = "black",
  label_alpha = 1,
  label_parse = FALSE,
  check_overlap = FALSE,
  angle_calc = "rot",
  force_flip = TRUE,
  label_dodge = NULL,
  label_push = NULL,
  show.legend = NA,
  ...,
  curvature
)

geom_edge_hive0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  strength = 1,
  lineend = "butt",
  show.legend = NA,
  ...,
  curvature
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default <code>x</code> , <code>y</code> , <code>xend</code> , <code>yend</code> , <code>group</code> and <code>circular</code> are mapped to <code>x</code> , <code>y</code> , <code>xend</code> , <code>yend</code> , <code>edge.id</code> and <code>circular</code> in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
strength	The curvature of the bezier. Defines the distance from the control points to the midpoint between the start and end node. 1 means the control points are positioned halfway between the nodes and the middle of the two axes, while 0 means it coincide with the nodes (resulting in straight lines)
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.
curvature	Deprecated. Use <code>strength</code> instead.

**Aesthetics**

geom\_edge\_hive and geom\_edge\_hive0 understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_hive2 understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_hive and geom\_edge\_hive2 furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

**Computed variables**

**index** The position along the path (not computed for the \*0 version)

## Edge variants

Many `geom_edge_*` layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. `colour = after_stat(index)`. The version postfixed with a "2" uses the "long" edge format (see `get_edges()`) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfixed with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. `geom_edge_*` and `geom_edge_*2` supports this through the `start_cap` and `end_cap` aesthetics that takes a `geometry()` specification and dynamically caps the termini of the edges based on the given specifications. This means that if `end_cap = circle(1, 'cm')` the edges will end at a distance of 1cm even during resizing of the plot window.

All `geom_edge_*` and `geom_edge_*2` have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

## Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

## Author(s)

Thomas Lin Pedersen

## See Also

Other `geom_edge_*`: `geom_edge_arc()`, `geom_edge_bend()`, `geom_edge_bundle_force()`, `geom_edge_bundle_minimal`, `geom_edge_bundle_path()`, `geom_edge_density()`, `geom_edge_diagonal()`, `geom_edge_elbow()`, `geom_edge_fan()`, `geom_edge_link()`, `geom_edge_loop()`, `geom_edge_parallel()`, `geom_edge_point()`, `geom_edge_sf()`, `geom_edge_span()`, `geom_edge_tile()`

## Examples

```
# Plot the flare import graph as a hive plot
library(tidygraph)
flareGr <- as_tbl_graph(flare$imports) %>%
```

```

mutate(
  type = dplyr::case_when(
    centrality_degree(mode = 'in') == 0 ~ 'Source',
    centrality_degree(mode = 'out') == 0 ~ 'Sink',
    TRUE ~ 'Both'
  )
) %>%
activate(edges) %>%
mutate(
  type = dplyr::case_when(
    grepl('flare.analytics', paste(.N())$name[from], .N())$name[to])) ~ 'Analytics',
    TRUE ~ 'Other'
  )
)

ggraph(flareGr, 'hive', axis = type) +
  geom_edge_hive(aes(colour = type), edge_alpha = 0.1) +
  coord_fixed()

```

---

geom\_edge\_link

*Draw edges as straight lines between nodes*


---

## Description

This geom draws edges in the simplest way - as straight lines between the start and end nodes. Not much more to say about that...

## Usage

```

geom_edge_link(
  mapping = NULL,
  data = get_edges("short"),
  position = "identity",
  arrow = NULL,
  n = 100,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  label_colour = "black",
  label_alpha = 1,
  label_parse = FALSE,
  check_overlap = FALSE,
  angle_calc = "rot",
  force_flip = TRUE,
  label_dodge = NULL,
  label_push = NULL,
  show.legend = NA,
  ...

```

```

)

geom_edge_link2(
  mapping = NULL,
  data = get_edges("long"),
  position = "identity",
  arrow = NULL,
  n = 100,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  label_colour = "black",
  label_alpha = 1,
  label_parse = FALSE,
  check_overlap = FALSE,
  angle_calc = "rot",
  force_flip = TRUE,
  label_dodge = NULL,
  label_push = NULL,
  show.legend = NA,
  ...
)

geom_edge_link0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  lineend = "butt",
  show.legend = NA,
  ...
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).



linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <a href="#">grDevices::plotmath()</a> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <a href="#">grid::unit()</a> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
label_push	A <a href="#">grid::unit()</a> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
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.
...	Other arguments passed on to <a href="#">layer()</a> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Edge variants

Many `geom_edge_*` layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. `colour = after_stat(index)`. The version postfix with a "2" uses the "long" edge format (see [get\\_edges\(\)](#)) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfix with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. `geom_edge_*` and `geom_edge_*2` supports this through the `start_cap` and `end_cap` aesthetics that takes a [geometry\(\)](#) specification and dynamically caps the termini of the edges based on the given specifications. This means that if `end_cap = circle(1, 'cm')` the edges will end at a distance of 1cm even during resizing of the plot window.

All `geom_edge_*` and `geom_edge_*2` have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs

to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Aesthetics

`geom_edge_link` and `geom_edge_link0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- `edge_colour`
- `edge_width`
- `edge_linetype`
- `edge_alpha`
- `filter`

`geom_edge_link2` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- `edge_colour`
- `edge_width`
- `edge_linetype`
- `edge_alpha`
- `filter`

`geom_edge_link` and `geom_edge_link2` furthermore takes the following aesthetics.

- `start_cap`
- `end_cap`

- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Author(s)

Thomas Lin Pedersen

### See Also

Other geom\_edge\_\*: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
gr <- create_notable('bull') %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE)) %>%
  activate(edges) %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE))

ggraph(gr, 'stress') +
  geom_edge_link(aes(alpha = after_stat(index)))

ggraph(gr, 'stress') +
  geom_edge_link2(aes(colour = node.class))

ggraph(gr, 'stress') +
  geom_edge_link0(aes(colour = class))
```

---

geom_edge_loop	<i>Draw edges as diagonals</i>
----------------	--------------------------------

---

## Description

This geom draws edge loops (edges starting and ending at the same node). Loops are drawn as bezier curves starting and ending at the position of the node and with control points protruding at an angle and in a direction specified in the call. As the start and end node is always the same no \*2 method is provided. Loops can severely clutter up your visualization which is why they are decoupled from the other edge drawings. Only plot them if they are of importance. If the graph doesn't contain any loops the geom adds nothing silently.

## Usage

```
geom_edge_loop(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  n = 100,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  label_colour = "black",
  label_alpha = 1,
  label_parse = FALSE,
  check_overlap = FALSE,
  angle_calc = "rot",
  force_flip = TRUE,
  label_dodge = NULL,
  label_push = NULL,
  show.legend = NA,
  ...
)

geom_edge_loop0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  lineend = "butt",
  show.legend = NA,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

**Aesthetics**

`geom_edge_loop` and `geom_edge_loop0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **from**
- **to**
- **span** 90
- **direction** 45
- **strength** 1
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_loop furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Edge variants

Many geom\_edge\_\* layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. colour = after\_stat(index). The version postfix with a "2" uses the "long" edge format (see [get\\_edges\(\)](#)) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfix with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. `geom_edge_*` and `geom_edge_*2` supports this through the `start_cap` and `end_cap` aesthetics that takes a `geometry()` specification and dynamically caps the termini of the edges based on the given specifications. This means that if `end_cap = circle(1, 'cm')` the edges will end at a distance of 1cm even during resizing of the plot window.

All `geom_edge_*` and `geom_edge_*2` have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
gr <- as_tbl_graph(
  data.frame(from = c(1, 1, 2, 2, 3, 3, 3), to = c(1, 2, 2, 3, 3, 1, 2))
)

ggraph(gr, 'stress') +
  geom_edge_loop(aes(alpha = after_stat(index))) +
  geom_edge_fan(aes(alpha = after_stat(index)))

ggraph(gr, 'stress') +
  geom_edge_loop0() +
  geom_edge_fan0()
```

---

geom\_edge\_parallel     *Draw multi edges as parallel lines*

---

### Description

This geom draws multi edges as parallel lines. The edges are first sorted by direction and then shifted a fixed amount so that all edges are visible.

### Usage

```
geom_edge_parallel(  
  mapping = NULL,  
  data = get_edges(),  
  position = "identity",  
  arrow = NULL,  
  sep = unit(2, "mm"),  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",  
  force_flip = TRUE,  
  label_dodge = NULL,  
  label_push = NULL,  
  show.legend = NA,  
  ...  
)  
  
geom_edge_parallel2(  
  mapping = NULL,  
  data = get_edges("long"),  
  position = "identity",  
  arrow = NULL,  
  sep = unit(2, "mm"),  
  n = 100,  
  lineend = "butt",  
  linejoin = "round",  
  linemitre = 1,  
  label_colour = "black",  
  label_alpha = 1,  
  label_parse = FALSE,  
  check_overlap = FALSE,  
  angle_calc = "rot",
```



```

    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...
)

geom_edge_parallel0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  arrow = NULL,
  sep = unit(2, "mm"),
  lineend = "butt",
  show.legend = NA,
  ...
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
sep	The separation between parallel edges, given as a <code>grid::unit()</code>
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).
label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.

force_flip	Logical. If angle_calc is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of angle_calc is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of angle_calc is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Aesthetics

`geom_edge_parallel` and `geom_edge_parallel0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- **from**
- **to**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_parallel2` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **group**
- **from**
- **to**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_parallel and geom\_edge\_parallel2 furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Edge variants

Many geom\_edge\_\* layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. colour = after\_stat(index). The version postfixed with a "2" uses the "long" edge format (see [get\\_edges\(\)](#)) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfixed with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. geom\_edge\_\* and geom\_edge\_\*2 supports this through the start\_cap and end\_cap aesthetics that takes a [geometry\(\)](#) specification and dynamically caps the termini of the edges based on the given specifications. This means that if end\_cap = circle(1, 'cm') the edges will end at a distance of 1cm even during resizing of the plot window.

All geom\_edge\_\* and geom\_edge\_\*2 have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs to know the edge type etc. Labels are drawn by providing a label aesthetic. The label\_pos can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The label\_size aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the angle\_calc argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

**Edge aesthetic name expansion**

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

**Author(s)**

David Schoch and Thomas Lin Pedersen

**See Also**

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

**Examples**

```
require(tidygraph)
gr <- create_notable('bull') %>%
  convert(to_directed) %>%
  bind_edges(data.frame(from = c(1, 2, 2, 3), to = c(2, 1, 3, 2))) %E>%
  mutate(class = sample(letters[1:3], 9, TRUE)) %N>%
  mutate(class = sample(c('x', 'y'), 5, TRUE))

ggraph(gr, 'stress') +
  geom_edge_parallel(aes(alpha = after_stat(index)))

ggraph(gr, 'stress') +
  geom_edge_parallel2(aes(colour = node.class))

ggraph(gr, 'stress') +
  geom_edge_parallel0(aes(colour = class))

# Use capping and sep to fine tune the look
ggraph(gr, 'stress') +
  geom_edge_parallel(start_cap = circle(1), end_cap = circle(1),
                    arrow = arrow(length = unit(2, 'mm')), sep = unit(4, 'mm')) +
  geom_node_point(size = 12)
```

---

geom\_edge\_point

*Draw edges as glyphs*

---

**Description**

This geom draws edges as glyphs with their x-position defined by the x-position of the start node, and the y-position defined by the y-position of the end node. As such it will result in a matrix layout when used in conjunction with [layout\\_tbl\\_graph\\_matrix\(\)](#)

**Usage**

```
geom_edge_point(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  mirror = FALSE,
  show.legend = NA,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
mirror	Logical. Should edge points be duplicated on both sides of the diagonal. Intended for undirected graphs. Default to FALSE
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

**Aesthetics**

`geom_edge_point` understands the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- edge\_shape
- edge\_colour
- edge\_size
- edge\_alpha
- filter

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
gr <- create_notable('zachary') %>%
  mutate(group = group_infomap()) %>%
  morph(to_split, group) %>%
  activate(edges) %>%
  mutate(edge_group = as.character(.N())$group[1]) %>%
  unmorph()

ggraph(gr, 'matrix', sort.by = node_rank_hclust()) +
  geom_edge_point(aes(colour = edge_group), mirror = TRUE, edge_size = 3) +
  scale_y_reverse() +
  coord_fixed() +
  labs(edge_colour = 'Infomap Cluster') +
  ggtitle("Zachary' Karate Club")
```

---

geom\_edge\_sf

*Draw edges as LINESTRINGs in geographical space*

---

### Description

This geom is equivalent in functionality to `ggplot2::geom_sf()` for `LINESTRING` geometries and allows for plotting of edges in their geographical space in different colours, linetypes and widths.

### Usage

```
geom_edge_sf(
  mapping = NULL,
  data = get_sf_edges(),
  position = "identity",
  show.legend = NA,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default geometry is mapped to the geometry in the edge data.
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 <code>ggplot()</code> . A <code>data.frame</code> , or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter()</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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. You can also set this to one of "polygon", "line", and "point" to override the default legend.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

**Aesthetics**

`geom_edge_sf` understand the following aesthetics.

- alpha
- colour
- linetype
- filter

**Author(s)**

Lorena Abad

**See Also**

Other `geom_edge_*`: `geom_edge_arc()`, `geom_edge_bend()`, `geom_edge_bundle_force()`, `geom_edge_bundle_minimal`, `geom_edge_bundle_path()`, `geom_edge_density()`, `geom_edge_diagonal()`, `geom_edge_elbow()`, `geom_edge_fan()`, `geom_edge_hive()`, `geom_edge_link()`, `geom_edge_loop()`, `geom_edge_parallel()`, `geom_edge_point()`, `geom_edge_span()`, `geom_edge_tile()`

**Examples**

```
if (require("sfnetworks", quietly = TRUE)) {
  gr <- sfnetworks::as_sfnetwork(roxel)
  ggraph(gr, 'sf') + geom_edge_sf()
}
```

---

 geom\_edge\_span

*Draw edges as vertical spans*


---

**Description**

This edge geom is mainly intended for use with [fabric](#) layouts. It draws edges as vertical segments with an optional end shape adornment. Due to the special nature of fabric layouts where nodes are not a single point in space but a line, this geom doesn't derive the x position from the location of the terminal nodes, but defaults to using the edge\_x variable calculated by the fabric layout. If this geom is used with other layouts x and xend must be given explicitly.

**Usage**

```
geom_edge_span(
  mapping = NULL,
  data = get_edges("short"),
  position = "identity",
  end_shape = NA,
  arrow = NULL,
  n = 100,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  label_colour = "black",
  label_alpha = 1,
  label_parse = FALSE,
  check_overlap = FALSE,
  angle_calc = "rot",
  force_flip = TRUE,
  label_dodge = NULL,
  label_push = NULL,
  show.legend = NA,
  ...
)

geom_edge_span2(
  mapping = NULL,
  data = get_edges("long"),
  position = "identity",
  end_shape = NA,
```



```

    arrow = NULL,
    n = 100,
    lineend = "butt",
    linejoin = "round",
    linemitre = 1,
    label_colour = "black",
    label_alpha = 1,
    label_parse = FALSE,
    check_overlap = FALSE,
    angle_calc = "rot",
    force_flip = TRUE,
    label_dodge = NULL,
    label_push = NULL,
    show.legend = NA,
    ...
)

geom_edge_span0(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  end_shape = NA,
  arrow = NULL,
  lineend = "butt",
  show.legend = NA,
  ...
)

```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <code>get_edges()</code> or a data.frame giving edges in correct format (see details for for guidance on the format). See <code>get_edges()</code> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
end_shape	The adornment to put at the ends of the span. The naming follows the conventions of the shape aesthetic in <code>ggplot2::geom_point()</code>
arrow	Arrow specification, as created by <code>grid::arrow()</code> .
n	The number of points to create along the path.
lineend	Line end style (round, butt, square).
linejoin	Line join style (round, mitre, bevel).
linemitre	Line mitre limit (number greater than 1).

label_colour	The colour of the edge label. If NA it will use the colour of the edge.
label_alpha	The opacity of the edge label. If NA it will use the opacity of the edge.
label_parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>grDevices::plotmath()</code> .
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
angle_calc	Either 'none', 'along', or 'across'. If 'none' the label will use the angle aesthetic of the geom. If 'along' The label will be written along the edge direction. If 'across' the label will be written across the edge direction.
force_flip	Logical. If <code>angle_calc</code> is either 'along' or 'across' should the label be flipped if it is on it's head. Default to TRUE.
label_dodge	A <code>grid::unit()</code> giving a fixed vertical shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
label_push	A <code>grid::unit()</code> giving a fixed horizontal shift to add to the label in case of <code>angle_calc</code> is either 'along' or 'across'
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Aesthetics

`geom_edge_span` and `geom_edge_span0` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- **xend**
- **yend**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

`geom_edge_span2` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**

- **group**
- edge\_colour
- edge\_width
- edge\_linetype
- edge\_alpha
- filter

geom\_edge\_span and geom\_edge\_span2 furthermore takes the following aesthetics.

- start\_cap
- end\_cap
- label
- label\_pos
- label\_size
- angle
- hjust
- vjust
- family
- fontface
- lineheight

### Computed variables

**index** The position along the path (not computed for the \*0 version)

### Edge variants

Many geom\_edge\_\* layers comes in 3 flavors depending on the level of control needed over the drawing. The default (no numeric postfix) generate a number of points (n) along the edge and draws it as a path. Each point along the line has a numeric value associated with it giving the position along the path, and it is therefore possible to show the direction of the edge by mapping to this e.g. colour = after\_stat(index). The version postfix with a "2" uses the "long" edge format (see [get\\_edges\(\)](#)) and makes it possible to interpolate node parameter between the start and end node along the edge. It is considerable less performant so should only be used if this is needed. The version postfix with a "0" draws the edge in the most performant way, often directly using an appropriate grob from the grid package, but does not allow for gradients along the edge.

Often it is beneficial to stop the drawing of the edge before it reaches the node, for instance in cases where an arrow should be drawn and the arrowhead shouldn't lay on top or below the node point. geom\_edge\_\* and geom\_edge\_\*2 supports this through the start\_cap and end\_cap aesthetics that takes a [geometry\(\)](#) specification and dynamically caps the termini of the edges based on the given specifications. This means that if end\_cap = circle(1, 'cm') the edges will end at a distance of 1cm even during resizing of the plot window.

All geom\_edge\_\* and geom\_edge\_\*2 have the ability to draw a label along the edge. The reason this is not a separate geom is that in order for the label to know the location of the edge it needs

to know the edge type etc. Labels are drawn by providing a label aesthetic. The `label_pos` can be used to specify where along the edge it should be drawn by supplying a number between 0 and 1. The `label_size` aesthetic can be used to control the size of the label. Often it is needed to have the label written along the direction of the edge, but since the actual angle is dependent on the plot dimensions this cannot be calculated beforehand. Using the `angle_calc` argument allows you to specify whether to use the supplied angle aesthetic or whether to draw the label along or across the edge.

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_tile\(\)](#)

### Examples

```
require(tidygraph)
gr <- play_smallworld(n_dim = 3, dim_size = 3, order = 1, p_rewire = 0.6)

# Standard use
ggraph(gr, 'fabric', sort.by = node_rank_fabric()) +
  geom_node_range(colour = 'grey80') +
  geom_edge_span()

# Add end shapes
ggraph(gr, 'fabric', sort.by = node_rank_fabric()) +
  geom_node_range(colour = 'grey80') +
  geom_edge_span(end_shape = 'circle')

# If the layout include shadow edges these can be styled differently
ggraph(gr, 'fabric', sort.by = node_rank_fabric(), shadow.edges = TRUE) +
  geom_node_range(colour = 'grey80') +
  geom_edge_span(aes(colour = shadow_edge), end_shape = 'square') +
  scale_edge_colour_manual(values = c('FALSE' = 'black', 'TRUE' = 'grey'))
```

---

geom_edge_tile	<i>Draw edges as glyphs</i>
----------------	-----------------------------

---

### Description

This geom draws edges as tiles with their x-position defined by the x-position of the start node, and the y-position defined by the y-position of the end node. As such it will result in a matrix layout when used in conjunction with [layout\\_tbl\\_graph\\_matrix\(\)](#)

### Usage

```
geom_edge_tile(
  mapping = NULL,
  data = get_edges(),
  position = "identity",
  mirror = FALSE,
  show.legend = NA,
  ...
)
```

### Arguments

mapping	Set of aesthetic mappings created by <a href="#">ggplot2::aes()</a> or <a href="#">ggplot2::aes_()</a> . By default x, y, xend, yend, group and circular are mapped to x, y, xend, yend, edge.id and circular in the edge data.
data	The return of a call to <a href="#">get_edges()</a> or a data.frame giving edges in correct format (see details for for guidance on the format). See <a href="#">get_edges()</a> for more details on edge extraction.
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <a href="#">position_jitter</a> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
mirror	Logical. Should edge points be duplicated on both sides of the diagonal. Intended for undirected graphs. Default to FALSE
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.
...	Other arguments passed on to <a href="#">layer()</a> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Aesthetics

`geom_edge_tile` understands the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**

- `y`
- `edge_fill`
- `edge_colour`
- `edge_size`
- `edge_alpha`
- `filter`

### Edge aesthetic name expansion

In order to avoid excessive typing edge aesthetic names are automatically expanded. Because of this it is not necessary to write `edge_colour` within the `aes()` call as `colour` will automatically be renamed appropriately.

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_edge_*`: [geom\\_edge\\_arc\(\)](#), [geom\\_edge\\_bend\(\)](#), [geom\\_edge\\_bundle\\_force\(\)](#), [geom\\_edge\\_bundle\\_minimal](#), [geom\\_edge\\_bundle\\_path\(\)](#), [geom\\_edge\\_density\(\)](#), [geom\\_edge\\_diagonal\(\)](#), [geom\\_edge\\_elbow\(\)](#), [geom\\_edge\\_fan\(\)](#), [geom\\_edge\\_hive\(\)](#), [geom\\_edge\\_link\(\)](#), [geom\\_edge\\_loop\(\)](#), [geom\\_edge\\_parallel\(\)](#), [geom\\_edge\\_point\(\)](#), [geom\\_edge\\_sf\(\)](#), [geom\\_edge\\_span\(\)](#)

### Examples

```
require(tidygraph)
gr <- create_notable('zachary') %>%
  mutate(group = group_infomap()) %>%
  morph(to_split, group) %>%
  activate(edges) %>%
  mutate(edge_group = as.character(.N())$group[1]) %>%
  unmorph()

ggraph(gr, 'matrix', sort.by = node_rank_hclust()) +
  geom_edge_tile(aes(fill = edge_group), mirror = TRUE) +
  scale_y_reverse() +
  coord_fixed() +
  labs(edge_colour = 'Infomap Cluster') +
  ggtitle("Zachary' Karate Club")
```

---

geom_node_arc_bar	<i>Show nodes as thick arcs</i>
-------------------	---------------------------------

---

## Description

This geom is equivalent in functionality to `ggforce::geom_arc_bar()` and allows for plotting of nodes as arcs with an inner and outer radius scaled by the coordinate system. Its main use is currently in sunburst plots as created with circular partition layouts

## Usage

```
geom_node_arc_bar(
  mapping = NULL,
  data = NULL,
  position = "identity",
  show.legend = NA,
  ...
)
```

## Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x and y are mapped to x0 and y0 in the node data.
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 <code>ggplot()</code> . A <code>data.frame</code> , or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

### Aesthetics

geom\_node\_point understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x0**
- **y0**
- **r0**
- **r**
- **start**
- **end**
- alpha
- colour
- fill
- shape
- size
- stroke
- filter

### Author(s)

Thomas Lin Pedersen

### See Also

Other geom\_node\_\*: [geom\\_node\\_circle\(\)](#), [geom\\_node\\_point\(\)](#), [geom\\_node\\_range\(\)](#), [geom\\_node\\_sf\(\)](#), [geom\\_node\\_text\(\)](#), [geom\\_node\\_tile\(\)](#), [geom\\_node\\_voronoi\(\)](#)

### Examples

```
require(tidygraph)
gr <- tbl_graph(flare$vertices, flare$edges)
ggraph(gr, 'partition', circular = TRUE, weight = size) +
  geom_node_arc_bar()
```

---

geom\_node\_circle

*Show nodes as circles*

---

### Description

This geom is equivalent in functionality to [ggforce::geom\\_circle\(\)](#) and allows for plotting of nodes as circles with a radius scaled by the coordinate system. Because of the geoms reliance on the coordinate system it will only produce true circles when combined with [ggplot2::coord\\_fixed\(\)](#)



**Usage**

```
geom_node_circle(
  mapping = NULL,
  data = NULL,
  position = "identity",
  show.legend = NA,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x and y are mapped to <code>x0</code> and <code>y0</code> in the node data.
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 <code>ggplot()</code> . A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> 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. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

**Aesthetics**

`geom_node_circle` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x0**
- **y0**
- **r**
- alpha
- colour
- fill
- shape
- size

- stroke
- filter

**Author(s)**

Thomas Lin Pedersen

**See Also**

Other geom\_node\_\*: [geom\\_node\\_arc\\_bar\(\)](#), [geom\\_node\\_point\(\)](#), [geom\\_node\\_range\(\)](#), [geom\\_node\\_sf\(\)](#), [geom\\_node\\_text\(\)](#), [geom\\_node\\_tile\(\)](#), [geom\\_node\\_voronoi\(\)](#)

**Examples**

```
require(tidygraph)
gr <- tbl_graph(flare$vertices, flare$edges)
ggraph(gr, 'circlepack', weight = size) +
  geom_node_circle() +
  coord_fixed()
```

---

geom_node_point	<i>Show nodes as points</i>
-----------------	-----------------------------

---

**Description**

This geom is equivalent in functionality to [ggplot2::geom\\_point\(\)](#) and allows for simple plotting of nodes in different shapes, colours and sizes.

**Usage**

```
geom_node_point(
  mapping = NULL,
  data = NULL,
  position = "identity",
  show.legend = NA,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <a href="#">ggplot2::aes()</a> or <a href="#">ggplot2::aes_()</a> . By default x and y are mapped to x and y in the node data.
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 <a href="#">ggplot()</a> . A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See <a href="#">fortify()</a> for which variables will be created.

	A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired <code>geom/stat</code> .

### Aesthetics

`geom_node_point` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- alpha
- colour
- fill
- shape
- size
- stroke
- filter

### Author(s)

Thomas Lin Pedersen

### See Also

Other `geom_node_*`: [geom\\_node\\_arc\\_bar\(\)](#), [geom\\_node\\_circle\(\)](#), [geom\\_node\\_range\(\)](#), [geom\\_node\\_sf\(\)](#), [geom\\_node\\_text\(\)](#), [geom\\_node\\_tile\(\)](#), [geom\\_node\\_voronoi\(\)](#)

### Examples

```
require(tidygraph)
gr <- create_notable('bull') %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE))

ggraph(gr, 'stress') + geom_node_point()
```

---

geom_node_range	<i>Show nodes as a line spanning a horizontal range</i>
-----------------	---

---

## Description

This geom is most useful together with the [fabric](#) layout for showing the horizontal span of each node.

## Usage

```
geom_node_range(
  mapping = NULL,
  data = NULL,
  position = "identity",
  show.legend = NA,
  ...
)
```

## Arguments

mapping	Set of aesthetic mappings created by <a href="#">ggplot2::aes()</a> or <a href="#">ggplot2::aes_()</a> . By default x is mapped to xmin, xend is mapped to xmax and y and yend are mapped to y in the node data.
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 <a href="#">ggplot()</a> . A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See <a href="#">fortify()</a> 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. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <a href="#">position_jitter</a> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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.
...	Other arguments passed on to <a href="#">layer()</a> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

## Aesthetics

geom\_node\_point understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **xend**
- **y**
- **yend**
- alpha
- colour
- linetype
- size
- filter

## Author(s)

Thomas Lin Pedersen

## See Also

Other geom\_node\_\*: [geom\\_node\\_arc\\_bar\(\)](#), [geom\\_node\\_circle\(\)](#), [geom\\_node\\_point\(\)](#), [geom\\_node\\_sf\(\)](#), [geom\\_node\\_text\(\)](#), [geom\\_node\\_tile\(\)](#), [geom\\_node\\_voronoi\(\)](#)

## Examples

```
require(tidygraph)
gr <- as_tbl_graph(highschool)

ggraph(gr, layout = 'fabric') +
  geom_node_range()
```

---

geom\_node\_sf

*Show nodes as POINTs in geographical space*

---

## Description

This geom is equivalent in functionality to `ggplot2::geom_sf()` for POINT geometries and allows for plotting of nodes in their geographical space in different shapes, colours and sizes.

## Usage

```
geom_node_sf(
  mapping = NULL,
  data = get_sf_nodes(),
  position = "identity",
  show.legend = NA,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default geometry is mapped to the geometry in the node data.
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 <code>ggplot()</code> . A <code>data.frame</code> , or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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. You can also set this to one of "polygon", "line", and "point" to override the default legend.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

**Aesthetics**

`geom_node_sf` understand the following aesthetics.

- alpha
- colour
- shape
- size
- filter

**Author(s)**

Lorena Abad

**See Also**

Other `geom_node_*`: `geom_node_arc_bar()`, `geom_node_circle()`, `geom_node_point()`, `geom_node_range()`, `geom_node_text()`, `geom_node_tile()`, `geom_node_voronoi()`

## Examples

```
library(tidygraph)

if (require("sfnetworks", quietly = TRUE)) {
  gr <- sfnetworks::as_sfnetwork(roxel)
  ggraph(gr, 'sf') +
    geom_node_sf(aes(color = centrality_betweenness()))
}
```

---

geom_node_text	<i>Annotate nodes with text</i>
----------------	---------------------------------

---

## Description

These geoms are equivalent in functionality to `ggplot2::geom_text()` and `ggplot2::geom_label()` and allows for simple annotation of nodes.

## Usage

```
geom_node_text(
  mapping = NULL,
  data = NULL,
  position = "identity",
  parse = FALSE,
  nudge_x = 0,
  nudge_y = 0,
  check_overlap = FALSE,
  show.legend = NA,
  repel = FALSE,
  ...
)

geom_node_label(
  mapping = NULL,
  data = NULL,
  position = "identity",
  parse = FALSE,
  nudge_x = 0,
  nudge_y = 0,
  label.padding = unit(0.25, "lines"),
  label.r = unit(0.15, "lines"),
  label.size = 0.25,
  show.legend = NA,
  repel = FALSE,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x and y are mapped to x and y in the node data.
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 <code>ggplot()</code> . A <code>data.frame</code> , or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string, or the result of a call to a position adjustment function. Cannot be jointly specified with <code>nudge_x</code> or <code>nudge_y</code> .
parse	If TRUE, the labels will be parsed into expressions and displayed as described in <code>?plotmath</code> .
nudge_x, nudge_y	Horizontal and vertical adjustment to nudge labels by. Useful for offsetting text from points, particularly on discrete scales.
check_overlap	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
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.
repel	If TRUE, text labels will be repelled from each other to avoid overlapping, using the <code>GeomTextRepel</code> geom from the <code>ggrepel</code> package.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.
label.padding	Amount of padding around label. Defaults to 0.25 lines.
label.r	Radius of rounded corners. Defaults to 0.15 lines.
label.size	Size of label border, in mm.

**Aesthetics**

`geom_node_text` understands the following aesthetics. Bold aesthetics are automatically set, but can be overwritten. Italic aesthetics are required but not set by default

- **x**
- **y**
- *label*
- alpha



- angle
- colour
- family
- fontface
- hjust
- lineheight
- size
- vjust

**Author(s)**

Thomas Lin Pedersen

**See Also**

Other geom\_node\_\*: [geom\\_node\\_arc\\_bar\(\)](#), [geom\\_node\\_circle\(\)](#), [geom\\_node\\_point\(\)](#), [geom\\_node\\_range\(\)](#), [geom\\_node\\_sf\(\)](#), [geom\\_node\\_tile\(\)](#), [geom\\_node\\_voronoi\(\)](#)

**Examples**

```
require(tidygraph)
gr <- create_notable('bull') %>%
  mutate(class = sample(letters[1:3], n(), replace = TRUE))

ggraph(gr, 'stress') +
  geom_node_point(aes(label = class))

ggraph(gr, 'stress') +
  geom_node_label(aes(label = class), repel = TRUE)
```

---

geom\_node\_tile      *Draw the rectangles in a treemap*

---

**Description**

A treemap is a space filling layout that recursively divides a rectangle to the children of the node. Often only the leaf nodes are drawn as nodes higher up in the hierarchy would obscure what is below. `geom_treemap` is a shorthand for `geom_node_treemap` as node is implicit in the case of treemap drawing

**Usage**

```
geom_node_tile(
  mapping = NULL,
  data = NULL,
  position = "identity",
  show.legend = NA,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x, y, width and height are mapped to x, y, width and height in the node data.
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 <code>ggplot()</code> . A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> 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. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

**Aesthetics**

`geom_treemap` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- x
- y
- **width**
- **height**
- alpha
- colour
- fill
- size
- stroke
- filter

**Author(s)**

Thomas Lin Pedersen

**See Also**

Other geom\_node\_\*: [geom\\_node\\_arc\\_bar\(\)](#), [geom\\_node\\_circle\(\)](#), [geom\\_node\\_point\(\)](#), [geom\\_node\\_range\(\)](#), [geom\\_node\\_sf\(\)](#), [geom\\_node\\_text\(\)](#), [geom\\_node\\_voronoi\(\)](#)

**Examples**

```
# Create a graph of the flare class system
library(tidygraph)
flareGraph <- tbl_graph(flare$vertices, flare$edges) %>%
  mutate(
    class = map_bfs_chr(node_is_root(), .f = function(node, dist, path, ...) {
      if (dist <= 1) {
        return(shortName[node])
      }
      path$result[[nrow(path)]]
    })
  )

gggraph(flareGraph, 'treemap', weight = size) +
  geom_node_tile(aes(fill = class, filter = leaf, alpha = depth), colour = NA) +
  geom_node_tile(aes(linewidth = depth), colour = 'white') +
  scale_alpha(range = c(1, 0.5), guide = 'none') +
  scale_size(range = c(4, 0.2), guide = 'none')
```

---

geom\_node\_voronoi      *Show nodes as voronoi tiles*

---

**Description**

This geom is equivalent in functionality to [ggforce::geom\\_voronoi\\_tile\(\)](#) and allows for plotting of nodes as tiles from a voronoi tessellation. As with [ggforce::geom\\_voronoi\\_tile\(\)](#) it is possible to restrict the size of the tile to a fixed radius, as well as round corners and expand/contract the tile.

**Usage**

```
geom_node_voronoi(
  mapping = NULL,
  data = NULL,
  position = "identity",
  show.legend = NA,
  bound = NULL,
  eps = 1e-09,
  max.radius = NULL,
  normalize = FALSE,
  asp.ratio = 1,
  expand = 0,
  radius = 0,
```

```
    ...
  )
```

### Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> or <code>ggplot2::aes_()</code> . By default x and y are mapped to x and y in the node data and group set to -1.
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 <code>ggplot()</code> . A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> 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. <code>~ head(.x, 10)</code> ).
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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.
bound	The bounding rectangle for the tessellation or a custom polygon to clip the tessellation to. Defaults to NULL which creates a rectangle expanded 10\ vector giving the bounds in the following order: xmin, xmax, ymin, ymax. If supplied as a polygon it should either be a 2-column matrix or a data.frame containing an x and y column.
eps	A value of epsilon used in testing whether a quantity is zero, mainly in the context of whether points are collinear. If anomalous errors arise, it is possible that these may averted by adjusting the value of eps upward or downward.
max.radius	The maximum distance a tile can extend from the point of origin. Will in effect clip each tile to a circle centered at the point with the given radius. If <code>normalize = TRUE</code> the radius will be given relative to the normalized values
normalize	Should coordinates be normalized prior to calculations. If x and y are in wildly different ranges it can lead to tessellation and triangulation that seems off when plotted without <code>ggplot2::coord_fixed()</code> . Normalization of coordinates solves this. The coordinates are transformed back after calculations.
asp.ratio	If <code>normalize = TRUE</code> the x values will be multiplied by this amount after normalization.
expand	A numeric or unit vector of length one, specifying the expansion amount. Negative values will result in contraction instead. If the value is given as a numeric it will be understood as a proportion of the plot area width.
radius	As <code>expand</code> but specifying the corner radius.
...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

## Aesthetics

`geom_node_voronoi` understand the following aesthetics. Bold aesthetics are automatically set, but can be overwritten.

- **x**
- **y**
- alpha
- colour
- fill
- shape
- size
- stroke
- filter

## Author(s)

Thomas Lin Pedersen

## See Also

Other `geom_node_*`: [geom\\_node\\_arc\\_bar\(\)](#), [geom\\_node\\_circle\(\)](#), [geom\\_node\\_point\(\)](#), [geom\\_node\\_range\(\)](#), [geom\\_node\\_sf\(\)](#), [geom\\_node\\_text\(\)](#), [geom\\_node\\_tile\(\)](#)

## Examples

```
require(tidygraph)
gr <- create_notable('meredith') %>%
  mutate(group = sample(letters[1:4], n(), TRUE))

ggraph(gr) +
  geom_node_voronoi(aes(fill = group, colour = group), alpha = 0.3) +
  geom_edge_link(alpha = 0.3) +
  geom_node_point()

# Use max.radius to make the tessellation more "node"-like
ggraph(gr) +
  geom_node_voronoi(aes(fill = group, colour = group), alpha = 0.3, max.radius = 1) +
  geom_edge_link(alpha = 0.3) +
  geom_node_point()
```

get\_con

*Create a connection extractor function***Description**

Connections within the ggraph terminology are links between nodes that are not part of the network structure itself. In that sense connections do not affect the layout calculation in any way and will not be drawn by the standard `geom_edge_*` functions. A connection does not need to only be defined by a start and end node, but can include intermediary nodes. `get_con` helps in creating connection data by letting you specify start and end nodes and automatically finds the shortest path within the graph structure that connects the given points. If this is not what is needed it is also possible to supply a list of vectors giving node indices that define a connection.

**Usage**

```
get_con(
  from = integer(),
  to = integer(),
  paths = NULL,
  ...,
  weight = NULL,
  mode = "all"
)
```

**Arguments**

<code>from, to</code>	The index of the start and end nodes for the connections
<code>paths</code>	A list of integer vectors giving the index of nodes defining connections
<code>...</code>	Additional information to be added to the final data output. Accepts expressions that will be evaluated on the node data in it's original order (irrespective of any reordering by the layout)
<code>weight</code>	An expression to be evaluated on the edge data to provide weights for the shortest path calculations
<code>mode</code>	Character constant, gives whether the shortest paths to or from the given vertices should be calculated for directed graphs. If <code>out</code> then the shortest paths <i>from</i> the vertex, if <code>in</code> then <i>to</i> it will be considered. If <code>all</code> , the default, then the corresponding undirected graph will be used, i.e. not directed paths are searched. This argument is ignored for undirected graphs.

**Value**

A function that takes a `layout_ggraph` object and returns the given connections

**See Also**

Other extractors: [get\\_edges\(\)](#), [get\\_sf\\_nodes\(\)](#)

---

get\_edges

*Create edge extractor function*


---

### Description

This function returns another function that can extract edges from a `ggraph_layout` object. The functionality of the returned function is decided by the arguments to `get_edges`. The need for `get_edges` is mainly to pass to the data argument of the different `geom_edge_*` functions in order to present them with the right kind of data. In general each `geom_edge_*` has the default set correctly so there is only need to modify the data argument if parallel edges should be collapsed.

### Usage

```
get_edges(format = "short", collapse = "none", ...)
```

```
get_sf_edges()
```

### Arguments

format	Either 'short' (the default) or 'long'. See details for a descriptions of the differences
collapse	Either 'none' (the default), 'all' or 'direction'. Specifies whether parallel edges should be merged. See details for more information
...	Additional data that will be <code>cbind</code> 'ed together with the returned edge data. Accepts expressions that will be evaluated on the edge data

### Details

There are two types of return formats possible for the result of the returned function:

**short** In this format each edge is described in one line in the format expected for `ggplot2::geom_segment()`, that is, the start node position is encoded in the `x` and `y` column and the end node position is encoded in the `xend` and `yend` column. If node parameters are added to the edge the name of the parameters will be prefixed with `node1.` for the start node and `node2.` for the end node.

**long** In this format each edge consists of two rows with matching `edge_id` value. The start and end position are both encoded in the `x` and `y` column. The relative position of the rows determines which is the start and end node, the first occurring being the start node. If node parameters are added to the edge data the name of the parameters will be prefixed with `node.`

Node parameters are automatically added so it is possible to format edge aesthetics according to start or end node parameters, or interpolate edge aesthetics between start and end node parameters. Node parameters will be prefixed to avoid name clash with edge parameters. The prefix depends on the format (see above).

If the graph is not simple (it contains at most one edge between each node pair) it can be collapsed so either all edges between two nodes or all edges of the same direction between two nodes are merged. The edge parameters are taken from the first occurring edge, so if some more sophisticated summary is needed it is suggested that the graph be tidied up before plotting with `ggraph`.

**Value**

A data.frame with columns dependent on format as well as the graph type. In addition to the columns discussed in the details section, the data.frame will always contain the columns `from`, `to` and `circular`, the two former giving the indexes of the start and end node and the latter if the layout is circular (needed for correct formatting of some `geom_edge_*`). The graph dependent information is:

**dendrogram** A `label` column will hold the value of the `edgetext` attribute. In addition any value stored in the `edgePar` attribute will be added. Lastly a `direction` column will hold the relative position between the start and end nodes (needed for correct formatting of `geom_edge_elbow()`).

**igraph** All edge attributes of the original graph object is added as columns to the data.frame

**See Also**

Other extractors: [get\\_con\(\)](#), [get\\_sf\\_nodes\(\)](#)

---

<code>get_sf_nodes</code>	<i>Create a node extractor function</i>
---------------------------	---

---

**Description**

This function returns another function that can extract nodes from a `ggraph_layout` object. As a `ggraph_layout` object is essentially a data.frame of nodes it might seem useless to provide this function, but since the node data is not necessarily available until after the `ggraph()` call it can be beneficial to be able to add information to the node data on a per-layer basis. Unlike [get\\_edges\(\)](#) the use of `get_nodes` is not mandatory and is only required if additional data should be added to selected node layers.

**Usage**

```
get_sf_nodes()
```

```
get_nodes(...)
```

**Arguments**

`...` Additional data that should be `cbind`'ed together with the node data. Accepts expressions that will be evaluated on the node data in it's original order (irrespective of any reordering by the layout)

**Value**

A data.frame with the node data as well of any additional data supplied through `...`

**See Also**

Other extractors: [get\\_con\(\)](#), [get\\_edges\(\)](#)



ggraph

*Create a ggraph plot***Description**

This function is the equivalent of `ggplot2::ggplot()` in `ggplot2`. It takes care of setting up the plot object along with creating the layout for the plot based on the graph and the specification passed in. Alternatively a layout can be prepared in advance using `create_layout` and passed as the data argument. See *Details* for a description of all available layouts.

**Usage**

```
ggraph(graph, layout = "auto", ...)

create_layout(graph, layout, circular, ...)

## Default S3 method:
create_layout(graph, layout, ...)

## S3 method for class 'layout_ggraph'
create_layout(graph, ...)

## S3 method for class 'tbl_graph'
create_layout(graph, layout, circular = FALSE, ...)
```

**Arguments**

<code>graph</code>	The object containing the graph. See <i>Details</i> for a list of supported classes. Or a <code>layout_ggraph</code> object as returned from <code>create_layout</code> in which case all subsequent arguments is ignored.
<code>layout</code>	The type of layout to create. Either a valid string, a function, a matrix, or a <code>data.frame</code> (see <i>Details</i> )
<code>...</code>	Arguments passed on to the layout function.
<code>circular</code>	Should the layout be transformed into a radial representation. Only possible for some layouts. Defaults to <code>FALSE</code>

**Details**

Following is a short description of the different layout types available in `ggraph`. Each layout is further described in its own help pages. Any type of regular graph/network data can be represented as a `tbl_graph` object. Because of this the different layouts that can be applied to `tbl_graph` objects are quite diverse, but not all layouts makes sense to all types of graphs. It is up to the user to understand their data and choose an appropriate layout. For standard node-edge diagrams `igraph` defines a long range of different layout functions that are all available through the `igraph` layout where the specific layout is specified using the `algorithm` argument. In order to minimize typing all `igraph` algorithms can also be passed directly into the `layout` argument.

Any object that has an appropriate `as_tbl_graph` method can be passed into `ggraph()` and will automatically be converted underneath.

- `auto` The default layout. See [layout\\_tbl\\_graph\\_auto\(\)](#) for further details
- `igraph` Use one of the internal igraph layout algorithms. The algorithm is specified using the `algorithm` argument. All strings accepted by the `algorithm` argument can also be supplied directly into `layout`. See [layout\\_tbl\\_graph\\_igraph\(\)](#) for further details
- `dendrogram` Lays out the nodes in a tree-like graph as a dendrogram with leaves set at 0 and parents 1 unit above its tallest child. See [layout\\_tbl\\_graph\\_dendrogram\(\)](#) for further details
- `manual` Lets the user manually specify the location of each node. See [layout\\_tbl\\_graph\\_manual\(\)](#) for further details
- `linear` Arranges the nodes linearly or circularly in order to make an arc diagram. See [layout\\_tbl\\_graph\\_linear\(\)](#) for further details
- `matrix` Arranges nodes on a diagonal thus preparing it for use with [geom\\_edge\\_point\(\)](#) to make a matrix plot. See [layout\\_tbl\\_graph\\_matrix\(\)](#) for further details
- `treemap` Creates a treemap from the graph, that is, a space-filing subdivision of rectangles showing a weighted hierarchy. See [layout\\_tbl\\_graph\\_treemap\(\)](#) for further details
- `circlepack` Creates a layout showing a hierarchy as circles within circles. Conceptually equal to treemaps. See [layout\\_tbl\\_graph\\_circlepack\(\)](#) for further details
- `partition` Create icicle or sunburst charts, where each layer subdivides the division given by the preceding layer. See [layout\\_tbl\\_graph\\_partition\(\)](#) for further details
- `hive` Positions nodes on axes spreading out from the center based on node attributes. See [layout\\_tbl\\_graph\\_hive\(\)](#) for further details
- `cactustree` Positions nodes as circles on the periphery of their parent circle. See [layout\\_tbl\\_graph\\_cactustree\(\)](#) for further details
- `backbone` Layout optimised for highly connected small-world graphs such as social networks. See [layout\\_tbl\\_graph\\_backbone\(\)](#) for further details
- `centrality` Place nodes around origin based on their centrality. See [layout\\_tbl\\_graph\\_centrality\(\)](#) for further details
- `eigen` Spectral layout based on the eigenvector of a matrix representation of the graph. See [layout\\_tbl\\_graph\\_eigen\(\)](#) for further details
- `fabric` Draw nodes as horizontal lines and connect them with vertical lines if an edge exists between them. See [layout\\_tbl\\_graph\\_fabric\(\)](#) for further details
- `focus` Place nodes around a focus node based on their distance to that node. See [layout\\_tbl\\_graph\\_focus\(\)](#) for further details
- `pmds` Layout based on multidimensional scaling of a set of pivot nodes, allowing MDS layout to be used on larger graphs. See [layout\\_tbl\\_graph\\_pmds\(\)](#) for further details
- `stress` Layout based on stress minimisation with better stability than Kamada-Kawai layout. See [layout\\_tbl\\_graph\\_stress\(\)](#) for further details
- `unrooted` Draws unrooted trees based on equal angle with optional equal daylight modification. See [layout\\_tbl\\_graph\\_unrooted\(\)](#) for further details
- `htree` Draws binary trees as a space filling fractal. See [layout\\_tbl\\_graph\\_htree\(\)](#) for further details

Alternatively a matrix or a data.frame can be provided to the layout argument. In the former case the first column will be used as x coordinates and the second column will be used as y coordinates, further columns are dropped. In the latter case the data.frame is used as the layout table and must thus contain a numeric x and y column.

Lastly a function can be provided to the layout argument. It will be called with the graph object as its first argument and any additional argument passed into `ggraph()/create_layout()`. The function must return either a data.frame or an object coercible to one and have an x and y column, or an object coercible to a `tbl_graph`. In the latter case the node data is extracted and used as layout (and must thus contain an x and y column) and the graph will be added as the graph attribute.

## Value

For `ggraph()` an object of class `gg` onto which layers, scales, etc. can be added. For `create_layout()` an object inheriting from `layout_ggraph`. `layout_ggraph` itself inherits from `data.frame` and can be considered as such. The data.frame contains the node positions in the x and y column along with additional columns generated by the specific layout, as well as node parameters inherited from the graph. Additional information is stored as attributes to the data.frame. The original graph object is stored in the `graph` attribute and the `circular` attribute contains a logical indicating whether the layout has been transformed to a circular representation.

## See Also

[get\\_edges\(\)](#) for extracting edge information from the layout and [get\\_con\(\)](#) for extracting path information.

## Examples

```
require(tidygraph)
gr <- create_notable('bull')
layout <- create_layout(gr, layout = 'igraph', algorithm = 'kk')
```

---

guide\_edge\_colourbar *Colourbar legend for edges*

---

## Description

This function is equivalent to `ggplot2::guide_colourbar()` but works for edge aesthetics.

## Usage

```
guide_edge_colourbar(..., available_aes = c("edge_colour", "edge_fill"))
guide_edge_colorbar(..., available_aes = c("edge_colour", "edge_fill"))
```

**Arguments**

- ... Arguments passed on to `ggplot2::guide_colourbar`
- title** A character string or expression indicating a title of guide. If NULL, the title is not shown. By default (`waiver()`), the name of the scale object or the name specified in `labs()` is used for the title.
- theme** A `theme` object to style the guide individually or differently from the plot's theme settings. The theme argument in the guide overrides, and is combined with, the plot's theme.
- nbin** A numeric specifying the number of bins for drawing the colourbar. A smoother colourbar results from a larger value.
- display** A string indicating a method to display the colourbar. Can be one of the following:
- "raster" to display as a bitmap image.
  - "rectangles" to display as a series of rectangles.
  - "gradient" to display as a linear gradient.
- Note that not all devices are able to render rasters and gradients.
- raster** **[Deprecated]** A logical. If TRUE then the colourbar is rendered as a raster object. If FALSE then the colourbar is rendered as a set of rectangles. Note that not all graphics devices are capable of rendering raster image.
- alpha** A numeric between 0 and 1 setting the colour transparency of the bar. Use NA to preserve the alpha encoded in the colour itself (default).
- draw.ulim** A logical specifying if the upper limit tick marks should be visible.
- draw.llim** A logical specifying if the lower limit tick marks should be visible.
- position** A character string indicating where the legend should be placed relative to the plot panels.
- direction** A character string indicating the direction of the guide. One of "horizontal" or "vertical."
- reverse** logical. If TRUE the colourbar is reversed. By default, the highest value is on the top and the lowest value is on the bottom
- order** positive integer less than 99 that specifies the order of this guide among multiple guides. This controls the order in which multiple guides are displayed, not the contents of the guide itself. If 0 (default), the order is determined by a secret algorithm.
- available\_aes** A vector of character strings listing the aesthetics for which a colourbar can be drawn.

**Value**

A guide object

---

 guide\_edge\_coloursteps

*Coloursteps legend for edges*


---

## Description

This function is equivalent to `ggplot2::guide_coloursteps()` but works for edge aesthetics.

## Usage

```
guide_edge_coloursteps(
  even.steps = TRUE,
  show.limits = NULL,
  ...,
  available_aes = c("edge_colour", "edge_fill")
)

guide_edge_colorsteps(
  even.steps = TRUE,
  show.limits = NULL,
  ...,
  available_aes = c("edge_colour", "edge_fill")
)
```

## Arguments

<code>even.steps</code>	Should the rendered size of the bins be equal, or should they be proportional to their length in the data space? Defaults to TRUE
<code>show.limits</code>	Logical. Should the limits of the scale be shown with labels and ticks. Default is NULL meaning it will take the value from the scale. This argument is ignored if <code>labels</code> is given as a vector of values. If one or both of the limits is also given in <code>breaks</code> it will be shown irrespective of the value of <code>show.limits</code> .
<code>...</code>	Arguments passed on to <code>ggplot2::guide_colourbar</code>
<code>title</code>	A character string or expression indicating a title of guide. If NULL, the title is not shown. By default ( <code>waiver()</code> ), the name of the scale object or the name specified in <code>labs()</code> is used for the title.
<code>theme</code>	A <code>theme</code> object to style the guide individually or differently from the plot's theme settings. The theme argument in the guide overrides, and is combined with, the plot's theme.
<code>nbin</code>	A numeric specifying the number of bins for drawing the colourbar. A smoother colourbar results from a larger value.
<code>display</code>	A string indicating a method to display the colourbar. Can be one of the following: <ul style="list-style-type: none"> <li>• "raster" to display as a bitmap image.</li> <li>• "rectangles" to display as a series of rectangles.</li> </ul>

- "gradient" to display as a linear gradient.

Note that not all devices are able to render rasters and gradients.

**raster** [**Deprecated**] A logical. If TRUE then the colourbar is rendered as a raster object. If FALSE then the colourbar is rendered as a set of rectangles.

Note that not all graphics devices are capable of rendering raster image.

**alpha** A numeric between 0 and 1 setting the colour transparency of the bar.

Use NA to preserve the alpha encoded in the colour itself (default).

**draw.ulim** A logical specifying if the upper limit tick marks should be visible.

**draw.llim** A logical specifying if the lower limit tick marks should be visible.

**position** A character string indicating where the legend should be placed relative to the plot panels.

**direction** A character string indicating the direction of the guide. One of "horizontal" or "vertical."

**reverse** logical. If TRUE the colourbar is reversed. By default, the highest value is on the top and the lowest value is on the bottom

**order** positive integer less than 99 that specifies the order of this guide among multiple guides. This controls the order in which multiple guides are displayed, not the contents of the guide itself. If 0 (default), the order is determined by a secret algorithm.

**available\_aes** A vector of character strings listing the aesthetics for which a colourbar can be drawn.

## Value

A guide object

---

guide\_edge\_direction *Edge direction guide*

---

## Description

This guide is intended to show the direction of edges based on the aesthetics mapped to its progression, such as changing width, colour and opacity.

## Usage

```
guide_edge_direction(
  title = NULL,
  theme = NULL,
  arrow = NULL,
  labels = NULL,
  nbin = 500,
  position = NULL,
  direction = NULL,
  reverse = FALSE,
```

```

  order = 0,
  override.aes = list(),
  ...,
  available_aes = c("edge_colour", "edge_alpha", "edge_width"),
  arrow.position = deprecated()
)

```

## Arguments

title	A character string or expression indicating a title of guide. If NULL, the title is not shown. By default ( <code>waiver()</code> ), the name of the scale object or the name specified in <code>labs()</code> is used for the title.
theme	A <code>theme</code> object to style the guide individually or differently from the plot's theme settings. The theme argument in the guide overrides, and is combined with, the plot's theme.
arrow	Logical. Should an arrow be drawn to illustrate the direction. Defaults to TRUE. The arrow is styled with the <code>legend.axis.line</code> theme element. If FALSE the direction will be indicated by the text given in <code>labels</code>
labels	A vector with two strings giving the labels to place at the start and the end of the legend to indicate direction if <code>arrow = FALSE</code>
nbin	A numeric specifying the number of bins for drawing the colourbar. A smoother colourbar results from a larger value.
position	A character string indicating where the legend should be placed relative to the plot panels.
direction	A character string indicating the direction of the guide. One of "horizontal" or "vertical."
reverse	logical. If TRUE the colourbar is reversed. By default, the highest value is on the top and the lowest value is on the bottom
order	positive integer less than 99 that specifies the order of this guide among multiple guides. This controls the order in which multiple guides are displayed, not the contents of the guide itself. If 0 (default), the order is determined by a secret algorithm.
override.aes	A list specifying aesthetic parameters of legend key. See details and examples.
...	ignored.
available_aes	A vector of character strings listing the aesthetics for which a colourbar can be drawn.
arrow.position	<b>[Deprecated]</b> The position of the arrow relative to the example edge. Use the <code>legend.text.position</code> argument in <code>theme()</code> instead.

## Examples

```

gr <- tidygraph::as_tbl_graph(highschool)
ggraph(gr, layout = 'kk') +
  geom_edge_fan(aes(alpha = after_stat(index))) +
  guides(edge_alpha = guide_edge_direction())

```

```

# Use text labels instead of an arrow
ggraph(gr, layout = 'kk') +
  geom_edge_fan(aes(alpha = after_stat(index))) +
  guides(edge_alpha = guide_edge_direction(labels = c('start', 'end')))

# Style the indicator arrow
arrow_style <- element_line(linewidth = 3, arrow = grid::arrow(type = "closed"))
ggraph(gr, layout = 'kk') +
  geom_edge_fan(aes(alpha = after_stat(index))) +
  guides(
    edge_alpha = guide_edge_direction(
      theme = theme(legend.axis.line = arrow_style)
    )
  )

```

---

highschool

*Friendship among high school boys*


---

### Description

This dataset shows the friendship among high school boys as assessed by the question: "What fellows here in school do you go around with most often?". The question was posed twice, with one year in between (1957 and 1958) and shows the evolution in friendship between the two timepoints.

### Usage

```
highschool
```

### Format

The graph is stored as an unnamed edgelist with a year attribute.

**from** The boy answering the question

**to** The boy being the answer to the question

**year** The year the friendship was reported

### Source

Coleman, J. S. *Introduction to Mathematical Sociology*. New York: Free Press, pp.450-451.



---

layout\_tbl\_graph\_auto *Automatically pick a layout based on graph type*

---

## Description

This function infers the layout from the graph structure and is the default when calling `ggraph()`. If an `x` and `y` argument is passed along, the manual layout is chosen. Otherwise if the graph is either a rooted tree or a rooted forest the layout will be dendrogram if the nodes contains a height variable or tree if not. If the tree is unrooted the unrooted layout will be used. If the tree is a DAG the sygiyama layout will be used. Otherwise the stress layout will be used (or `sparse_tree` if the graph contains more than 2000 nodes).

## Usage

```
layout_tbl_graph_auto(graph, circular, ...)
```

## Arguments

<code>graph</code>	A <code>tbl_graph</code> object
<code>circular</code>	Logical. Should the layout be transformed to a circular representation. Defaults to FALSE. Only applicable if the graph is a tree structure
<code>...</code>	Arguments passed on to the chosen layout

## Value

A `data.frame` with the columns `x`, `y`, `circular` as well as any information stored as node variables in the `tbl_graph` object.

## See Also

Other `layout_tbl_graph_*`: [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustree\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

 layout\_tbl\_graph\_backbone

*Place node to emphasize group structure*


---

### Description

This layout is optimised for drawing small-world types of graphs often found in social networks, where distinct groups are still highly connected to the remaining graph. Typical layouts struggle with this as they attempt to minimise the edge length of all edges equally. The backbone layout is based on weighing edges based on how well they hold together communities. The end result is that communities tend to stick together despite high interconnectivity.

### Usage

```
layout_tbl_graph_backbone(graph, keep = 0.2, circular = FALSE)
```

### Arguments

graph	A <code>tbl_graph</code> object
keep	The fraction of edges to use for creating the backbone
circular	ignored

### Value

A data.frame with the columns `x`, `y`, `circular` as well as any information stored as node variables in the `tbl_graph` object. Further an edge attribute called `backbone` is added giving whether the edge was selected as backbone.

### Author(s)

The underlying algorithm is implemented in the `graphlayouts` package by David Schoch

### References

Nocaj, A., Ortmann, M., & Brandes, U. (2015). *Untangling the hairballs of multi-centered, small-world online social media networks*. *Journal of Graph Algorithms and Applications: JGAA*, 19(2), 595-618.

### See Also

Other `layout_tbl_graph_*`: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_cactustree\(\)](#), [layout\\_tbl\\_graph\\_central\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

`layout_tbl_graph_cactustree`*Calculate nodes as fractal circle buds*

---

## Description

The cactustree layout is a hierarchical layout optimised for use with hierarchical edge bundling (`geom_conn_bundle()`). It is a fractal layout that places node children as circles on the periphery of their parent circle, each circle scaled by the total weight of their children.

## Usage

```
layout_tbl_graph_cactustree(  
  graph,  
  direction = "out",  
  weight = NULL,  
  scale_factor = 0.75,  
  overlap = 0.5,  
  upright = FALSE,  
  circular = FALSE  
)
```

## Arguments

<code>graph</code>	An <code>tbl_graph</code> object
<code>direction</code>	The direction of the tree in the graph. 'out' (default) means that parents point towards their children, while 'in' means that children point towards their parent.
<code>weight</code>	An optional node variable to use as weight. If NULL all leaf nodes will be assigned a weight of 1.
<code>scale_factor</code>	A scaling factor for the circles in the layout. The radius will be calculated as $\text{total\_weight}^{\text{scale\_factor}}$ with <code>total_weight</code> being the weight of the node and all its children.
<code>overlap</code>	How much is the center of child nodes offset from the periphery of their parent as a proportion of their own radius.
<code>upright</code>	Logical. Should the children of the root only be distributed around the top half of the root or all the way around.
<code>circular</code>	Logical. Should the layout be transformed to a circular representation. Ignored.

## Value

A data.frame with the columns `x`, `y`, `r`, `leaf`, `depth`, `circular` as well as any information stored as node variables in the `tbl_graph` object.

**Note**

cactustree is a layout intended for trees, that is, graphs where nodes only have one parent and zero or more children. If the provided graph does not fit this format an attempt to convert it to such a format will be made.

**References**

Dang, T., Forbes, A. (2017). *CactusTree: A Tree Drawing Approach for Hierarchical Edge Bundling*. 2017 IEEE Pacific Visualization Symposium, 210-214. <https://doi.org/10.1109/PACIFICVIS.2017.8031596>

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_centrality

*Place nodes in circles according to centrality measure*

---

**Description**

This layout places nodes in circles with the radii relative to a given centrality measure. Under the hood it use stress majorisation to place nodes optimally given the radius constraint.

**Usage**

```
layout_tbl_graph_centrality(
  graph,
  centrality,
  scale = TRUE,
  niter = 500,
  tolerance = 1e-04,
  tseq = seq(0, 1, 0.2),
  group = NULL,
  shrink = 10,
  circular = FALSE
)
```

**Arguments**

graph	A tbl_graph object
centrality	An expression evaluating to a centrality measure for the nodes. See the different centrality_*() algorithms in tidygraph for a selection.

scale	Should the centrality measure be scaled between 0 and 100
niter	number of iterations during stress optimization
tolerance	stopping criterion for stress optimization
tseq	Transitioning steps
group	An expression evaluating to a grouping of the nodes. If given the layout will keep grouped nodes within an angle range of the origin
shrink	shrink the reserved angle range for a group to increase the gaps between groups
circular	ignored

**Value**

A data.frame with the columns x, y, circular, centrality as well as any information stored as node variables in the tbl\_graph object.

**Author(s)**

The underlying algorithm is implemented in the graphlayouts package by David Schoch

**References**

Brandes, U., & Pich, C. (2011). *More flexible radial layout*. Journal of Graph Algorithms and Applications, 15(1), 157-173.

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustree\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_circlepack

*Calculate nodes as circles packed within their parent circle*

---

**Description**

The circle packing algorithm is basically a treemap using circles instead of rectangles. Due to the nature of circles they cannot be packed as efficiently leading to increased amount of "empty space" as compared to a treemap. This can be beneficial though, as the added empty space can aid in visually showing the hierarchy.

**Usage**

```
layout_tbl_graph_circlepack(
  graph,
  weight = NULL,
  circular = FALSE,
  sort.by = NULL,
  direction = "out"
)
```

**Arguments**

graph	An tbl_graph object
weight	An optional node variable to use as weight. Will only affect the weight of leaf nodes as the weight of non-leaf nodes are derived from their children.
circular	Logical. Should the layout be transformed to a circular representation. Ignored.
sort.by	The name of a node variable to sort the nodes by.
direction	The direction of the tree in the graph. 'out' (default) means that parents point towards their children, while 'in' means that children point towards their parent.

**Details**

The circle packing is based on the algorithm developed by Weixin Wang and collaborators which tries to find the most dense packing of circles as they are added, one by one. This makes the algorithm very dependent on the order in which circles are added and it is possible that layouts could sometimes be optimized by choosing a different ordering. The algorithm for finding the enclosing circle is the randomized incremental algorithm proposed by Emo Welzl. Both of the above algorithms are the same as used in the D3.js implementation of circle packing and their C++ implementation in ggraph is inspired by Mike Bostocks JavaScript implementation.

**Value**

A data.frame with the columns x, y, r, leaf, depth, circular as well as any information stored as node variables in the tbl\_graph object.

**Note**

Circle packing is a layout intended for trees, that is, graphs where nodes only have one parent and zero or more children. If the provided graph does not fit this format an attempt to convert it to such a format will be made.

**References**

- Wang, W., Wang, H. H., Dai, G., & Wang, H. (2006). *Visualization of large hierarchical data by circle packing*. *Chi*, 517-520.
- Welzl, E. (1991). *Smallest enclosing disks (balls and ellipsoids)*. *New Results and New Trends in Computer Science*, 359-370.

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matr](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_dendrogram

*Apply a dendrogram layout to layout\_tbl\_graph*

---

**Description**

This layout mimics the `igraph::layout_as_tree()` algorithm supplied by igraph, but puts all leaves at 0 and builds it up from there, instead of starting from the root and building it from there. The height of branch points are related to the maximum distance to an edge from the branch node, or read from a node variable.

**Usage**

```
layout_tbl_graph_dendrogram(
  graph,
  circular = FALSE,
  offset = pi/2,
  height = NULL,
  length = NULL,
  repel = FALSE,
  ratio = 1,
  direction = "out"
)
```

**Arguments**

graph	A tbl_graph object
circular	Logical. Should the layout be transformed to a circular representation. Defaults to FALSE.
offset	If circular = TRUE, where should it begin. Defaults to pi/2 which is equivalent to 12 o'clock.
height	The node variable holding the height of each node in the dendrogram. If NULL it will be calculated as the maximal distance to a leaf.
length	An edge parameter giving the length of each edge. The node height will be calculated from the maximal length to the root node (ignored if height does not evaluate to NULL)

repel	Should leaves repel each other relative to the height of their common ancestor. Will emphasize clusters
ratio	The strength of repulsion if <code>repel = TRUE</code> . Higher values will give more defined clusters
direction	The direction to the leaves. Defaults to 'out'

**Value**

A data.frame with the columns `x`, `y`, `circular`, `depth` and `leaf` as well as any information stored as node variables on the `tbl_graph`

**Note**

This function is not intended to be used directly but by setting `layout = 'dendrogram'` in `create_layout()`

**See Also**

Other `layout_tbl_graph_*`: `layout_tbl_graph_auto()`, `layout_tbl_graph_backbone()`, `layout_tbl_graph_cactustre`, `layout_tbl_graph_centrality()`, `layout_tbl_graph_circlepack()`, `layout_tbl_graph_eigen()`, `layout_tbl_graph_fabric()`, `layout_tbl_graph_focus()`, `layout_tbl_graph_hive()`, `layout_tbl_graph_htree()`, `layout_tbl_graph_igraph()`, `layout_tbl_graph_linear()`, `layout_tbl_graph_manual()`, `layout_tbl_graph_matr`, `layout_tbl_graph_metro()`, `layout_tbl_graph_partition()`, `layout_tbl_graph_pmds()`, `layout_tbl_graph_sf()`, `layout_tbl_graph_stress()`, `layout_tbl_graph_treemap()`, `layout_tbl_graph_unrooted()`

---

layout\_tbl\_graph\_eigen

*Place nodes according to their eigenvalues*

---

**Description**

This layout is based on the idea of spectral layouts where node coordinates are calculated directly by decomposing a matrix representation of the graph and extracting the eigenvectors.

**Usage**

```
layout_tbl_graph_eigen(
  graph,
  type = "laplacian",
  eigenvector = "smallest",
  circular = FALSE
)
```



**Arguments**

graph	A <code>tbl_graph</code> object
type	The type of matrix to extract the eigenvectors from. Either 'laplacian' or 'adjacency'
eigenvector	The eigenvector to use for coordinates. Either 'smallest' or 'largest'
circular	ignored

**Value**

A `data.frame` with the columns `x`, `y`, `circular` as well as any information stored as node variables in the `tbl_graph` object.

**Author(s)**

The underlying algorithm is implemented in the `graphlayouts` package by David Schoch

**See Also**

Other `layout_tbl_graph_*`: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustreemap\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_fabric

*Create a fabric layout*

---

**Description**

This layout is a bit unusual in that it shows nodes as horizontal line ranges and edges as evenly spaced vertical spans connecting the nodes. As with the matrix layout the strength comes from better scalability but its use requires some experience recognising the patterns that different connectivity features give rise to. As with matrix layouts the ordering of nodes has huge power over the look of the plot. The `node_rank_fabric()` mimics the default ordering from the original BioFabric implementation, but other ranking algorithms from `tidygraph` can be used with the `sort.by` argument as well. Fabric layouts tend to become quite wide as the graph grows which is something that should be handled with care - e.g. by only zooming in on a specific region.

**Usage**

```

layout_tbl_graph_fabric(
  graph,
  circular = FALSE,
  sort.by = NULL,
  shadow.edges = FALSE
)

node_rank_fabric()

```

**Arguments**

graph	An tbl_graph object
circular	Ignored
sort.by	An expression providing the sorting of the nodes. If NULL the nodes will be ordered by their index in the graph.
shadow.edges	Should shadow edges be shown.

**Value**

A data.frame with the columns x, xmin, xmax, y, circular as well as any information stored as node variables in the tbl\_graph object. Further, the edges of the graph will gain a edge\_x variable giving the horizontal position of the edge as well as a shadow\_edge variable denoting whether the edge is a shadow edge added by the layout.

**References**

BioFabric website: <https://biofabric.systemsbiology.net>

Longabaugh, William J.R. (2012). *Combing the hairball with BioFabric: a new approach for visualization of large networks*. BMC Bioinformatics, 13: 275. doi:10.1186/1471210513275

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matr](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

`layout_tbl_graph_focus`*Place nodes in circles based on distance to a specific node*

---

**Description**

This layout constrains node placement to a radius relative to its distance to a given node. It then uses stress majorisation to find an optimal node distribution according to this constraint.

**Usage**

```
layout_tbl_graph_focus(  
  graph,  
  focus,  
  weights = NULL,  
  niter = 500,  
  tolerance = 1e-04,  
  group = NULL,  
  shrink = 10,  
  circular = TRUE  
)
```

**Arguments**

<code>graph</code>	a <code>tbl_graph</code> object
<code>focus</code>	An expression evaluating to a selected node. Can either be a single integer or a logical vector with a single TRUE element.
<code>weights</code>	An expression evaluated on the edge data to provide edge weights for the layout. Currently ignored for the sparse version
<code>niter</code>	number of iterations during stress optimization
<code>tolerance</code>	stopping criterion for stress optimization
<code>group</code>	An expression evaluating to a grouping of the nodes. If given the layout will keep grouped nodes within an angle range of the origin
<code>shrink</code>	shrink the reserved angle range for a group to increase the gaps between groups
<code>circular</code>	ignored

**Value**

A data.frame with the columns `x`, `y`, `circular`, `distance` as well as any information stored as node variables in the `tbl_graph` object.

**Author(s)**

The underlying algorithm is implemented in the `graphlayouts` package by David Schoch

**References**

Brandes, U., & Pich, C. (2011). *More flexible radial layout*. *Journal of Graph Algorithms and Applications*, 15(1), 157-173.

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matr](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_hive *Place nodes in a Hive Plot layout*

---

**Description**

Hive plots were invented by Martin Krzywinski as a perceptually uniform and scalable alternative to standard node-edge layouts. In hive plots nodes are positioned on axes radiating out from a center based on their own information e.g. membership of a class, size of neighborhood, etc. Edges are then drawn between nodes as bezier curves. As the placement of nodes is not governed by convoluted algorithms but directly reflects the qualities of the nodes itself the resulting plot can be easier to interpret as well as compare to other graphs.

**Usage**

```
layout_tbl_graph_hive(  
  graph,  
  axis,  
  axis.pos = NULL,  
  sort.by = NULL,  
  divide.by = NULL,  
  divide.order = NULL,  
  normalize = TRUE,  
  center.size = 0.1,  
  divide.size = 0.05,  
  use.numeric = FALSE,  
  offset = pi/2,  
  split.axes = "none",  
  split.angle = pi/6,  
  circular = FALSE  
)
```

**Arguments**

graph	An tbl_graph object
axis	The node attribute to use for assigning nodes to axes
axis.pos	The relative distance to the prior axis. Default (NULL) places axes equidistant.
sort.by	The node attribute to use for placing nodes along their axis. Defaults (NULL) places nodes sequentially.
divide.by	An optional node attribute to subdivide each axis by.
divide.order	The order the axis subdivisions should appear in
normalize	Logical. Should axis lengths be equal or reflect the number of nodes in each axis. Defaults to TRUE.
center.size	The size of the blank center, that is, the start position of the axes.
divide.size	The distance between subdivided axis segments.
use.numeric	Logical, If the sort.by attribute is numeric, should these values be used directly in positioning the nodes along the axes. Defaults to FALSE which sorts the numeric values and positions them equidistant from each other.
offset	Change the overall rotation of the hive plot by changing the offset of the first axis.
split.axes	Should axes be split to show edges between nodes on the same axis? One of: 'none' Do not split axes and show in-between edges 'loops' Only split axes that contain in-between edges 'all' Split all axes
split.angle	The angular distance between the two axes resulting from a split.
circular	Ignored.

**Details**

In order to be able to draw all edges without edges crossing axes you should not assign nodes to axes based on a variable with more than three levels.

**Value**

A data.frame with the columns x, y, r, center\_size, split, axis, section, angle, circular as well as any information stored as node variables in the tbl\_graph object.

**References**

Krzywinski, M., Birol, I., Jones, SJM., and Marra, MA. (2012). *Hive plots-rational approach to visualizing networks*. *Brief Bioinform* 13 (5): 627-644. <https://doi.org/10.1093/bib/bbr069>

<https://www.hiveplot.net/>

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matr](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_htree

*Layout binary trees in a fractal H formation*

---

**Description**

This is a spac efficient layout only useful for binary trees. It is fractal and works by offsetting child nodes from their parent either horizontally or vertically depending on depth. The offset is decreased at each step by a factor of the square root of 2.

**Usage**

```
layout_tbl_graph_htree(
  graph,
  sort.by = NULL,
  direction = "out",
  circular = FALSE
)
```

**Arguments**

graph	An tbl_graph object
sort.by	The name of a node variable to sort the nodes by.
direction	The direction of the tree in the graph. 'out' (default) means that parents point towards their children, while 'in' means that children point towards their parent.
circular	Logical. Should the layout be transformed to a circular representation. Ignored

**Value**

A data.frame with the columns x, y, leaf, depth, circular as well as any information stored as node variables in the tbl\_graph object.

**Note**

H Tree is a layout intended for trees, that is, graphs where nodes only have one parent and zero or more children. If the provided graph does not fit this format an attempt to convert it to such a format will be made.

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matr](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_igraph

*Use igraph layout algorithms for layout\_tbl\_graph*

---

**Description**

This layout function makes it easy to apply one of the layout algorithms supplied in igraph when plotting with ggraph. Layout names are auto completed so there is no need to write layout\_with\_graphopt or layout\_as\_tree, just graphopt and tree (though the former will also work if you want to be super explicit). Circular layout is only supported for tree-like layout (tree and sugiyama) and will throw an error when applied to other layouts.

**Usage**

```
layout_tbl_graph_igraph(
  graph,
  algorithm,
  circular,
  offset = pi/2,
  use.dummy = FALSE,
  ...
)
```

**Arguments**

graph	A tbl_graph object.
algorithm	The type of layout algorithm to apply. See <i>Details</i> or <a href="#">igraph::layout_()</a> for links to the layouts supplied by igraph.
circular	Logical. Should the layout be transformed to a circular representation. Defaults to FALSE. Only applicable to algorithm = 'tree' and algorithm = 'sugiyama'.
offset	If circular = TRUE, where should it begin. Defaults to pi/2 which is equivalent to 12 o'clock.
use.dummy	Logical. In the case of algorithm = 'sugiyama' should the dummy-infused graph be used rather than the original. Defaults to FALSE.
...	Arguments passed on to the respective layout functions

**Details**

igraph provides a huge amount of possible layouts. They are all briefly described below:

**Hierarchical layouts**

`tree` Uses the *Reingold-Tilford* algorithm to place the nodes below their parent with the parent centered above its children. See `igraph::as_tree()`

`sugiyama` Designed for directed acyclic graphs (that is, hierarchies where multiple parents are allowed) it minimizes the number of crossing edges. See `igraph::with_sugiyama()`

**Standard layouts**

`bipartite` Minimize edge-crossings in a simple two-row (or column) layout for bipartite graphs. See `igraph::as_bipartite()`

`star` Place one node in the center and the rest equidistantly around it. See `igraph::as_star()`

`circle` Place nodes in a circle in the order of their index. Consider using `layout_tbl_graph_linear()` with `circular=TRUE` for more control. See `igraph::in_circle()`

`nicely` Tries to pick an appropriate layout. See `igraph::nicely()` for a description of the simple decision tree it uses

`dh` Uses *Davidson and Harels* simulated annealing algorithm to place nodes. See `igraph::with_dh()`

`gem` Place nodes on the plane using the GEM force-directed layout algorithm. See `igraph::with_gem()`

`graphopt` Uses the Graphopt algorithm based on alternating attraction and repulsion to place nodes. See `igraph::with_graphopt()`

`grid` Place nodes on a rectangular grid. See `igraph::on_grid()`

`mds` Perform a multidimensional scaling of nodes using either the shortest path or a user supplied distance. See `igraph::with_mds()`

`sphere` Place nodes uniformly on a sphere - less relevant for 2D visualizations of networks. See `igraph::on_sphere()`

`randomly` Places nodes uniformly random. See `igraph::randomly()`

`fr` Places nodes according to the force-directed algorithm of Fruchterman and Reingold. See `igraph::with_fr()`

`kk` Uses the spring-based algorithm by Kamada and Kawai to place nodes. See `igraph::with_kk()`

`drl` Uses the force directed algorithm from the DrL toolbox to place nodes. See `igraph::with_drl()`

`lgl` Uses the algorithm from Large Graph Layout to place nodes. See `igraph::with_lgl()`

**Value**

A data.frame with the columns `x`, `y`, `circular` as well as any information stored as node variables in the `tbl_graph` object.

**Note**

This function is not intended to be used directly but by setting `layout = 'igraph'` in `create_layout()`



**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_linear

*Place nodes on a line or circle*

---

**Description**

This layout puts all nodes on a line, possibly sorted by a node attribute. If `circular = TRUE` the nodes will be laid out on the unit circle instead. In the case where the `sort.by` attribute is numeric, the numeric values will be used as the x-position and it is thus possible to have uneven spacing between the nodes.

**Usage**

```
layout_tbl_graph_linear(  
  graph,  
  circular,  
  sort.by = NULL,  
  use.numeric = FALSE,  
  offset = pi/2,  
  weight = NULL  
)
```

**Arguments**

<code>graph</code>	An <code>tbl_graph</code> object
<code>circular</code>	Logical. Should the layout be transformed to a circular representation. Defaults to FALSE.
<code>sort.by</code>	The name of a node variable to sort the nodes by.
<code>use.numeric</code>	Logical. Should a numeric <code>sort.by</code> attribute be used as the actual x-coordinates in the layout. May lead to overlapping nodes. Defaults to FALSE
<code>offset</code>	If <code>circular = TRUE</code> , where should it begin. Defaults to <code>pi/2</code> which is equivalent to 12 o'clock.
<code>weight</code>	A weight for each node. Nodes will be spread out according to their weight so that nodes with heigher weight will have more space around them. Ignored if <code>use.numeric = TRUE</code>

**Value**

A data.frame with the columns `x`, `y`, `circular` as well as any information stored as node variables in the `tbl_graph` object. Further, if `circular = FALSE` a `width` column and if `circular = TRUE` a `start`, `end`, and `r0` column.

**See Also**

Other `layout_tbl_graph_*`: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustree\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

`layout_tbl_graph_manual`

*Manually specify a layout for layout\_tbl\_graph*

---

**Description**

This layout function lets you pass the node positions in manually. The supplied positions must match the order of the nodes in the `tbl_graph`

**Usage**

```
layout_tbl_graph_manual(graph, x, y, circular)
```

**Arguments**

<code>graph</code>	An <code>tbl_graph</code> object
<code>x</code> , <code>y</code>	Expressions with the <code>x</code> and <code>y</code> positions of the nodes
<code>circular</code>	Ignored

**Value**

A data.frame with the columns `x`, `y`, `circular` as well as any information stored as node variables in the `tbl_graph`.

**See Also**

Other `layout_tbl_graph_*`: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustree\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

 layout\_tbl\_graph\_matrix

*Place nodes on a diagonal*


---

## Description

This layout puts all nodes on a diagonal, thus preparing the layout for use with `geom_edge_point()` resulting in a matrix layout. While matrix layouts excel in scalability, the interpretation of the visual is very dependent on the sorting of the nodes. Different sorting algorithms have been implemented in `tidygraph` and these can be used directly. Behrisch *et al.* (2016) have provided a nice overview of some of the different sorting algorithms and what insight they might bring, along with a rundown of different patterns to look out for.

## Usage

```
layout_tbl_graph_matrix(graph, circular = FALSE, sort.by = NULL)
```

## Arguments

<code>graph</code>	An <code>tbl_graph</code> object
<code>circular</code>	Ignored
<code>sort.by</code>	An expression providing the sorting of the nodes. If <code>NULL</code> the nodes will be ordered by their index in the graph.

## Value

A `data.frame` with the columns `x`, `y`, `circular` as well as any information stored as node variables in the `tbl_graph` object.

## References

Behrisch, M., Bach, B., Riche, N. H., Schreck, T., Fekete, J.-D. (2016). *Matrix Reordering Methods for Table and Network Visualization*. *Computer Graphics Forum*, 35: 693–716. doi:[10.1111/cgf.12935](https://doi.org/10.1111/cgf.12935)

## See Also

Other `layout_tbl_graph_*`: `layout_tbl_graph_auto()`, `layout_tbl_graph_backbone()`, `layout_tbl_graph_cactustr`, `layout_tbl_graph_centrality()`, `layout_tbl_graph_circlepack()`, `layout_tbl_graph_dendrogram()`, `layout_tbl_graph_eigen()`, `layout_tbl_graph_fabric()`, `layout_tbl_graph_focus()`, `layout_tbl_graph_hive()`, `layout_tbl_graph_htree()`, `layout_tbl_graph_igraph()`, `layout_tbl_graph_linear()`, `layout_tbl_graph_manua`, `layout_tbl_graph_metro()`, `layout_tbl_graph_partition()`, `layout_tbl_graph_pmds()`, `layout_tbl_graph_sf()`, `layout_tbl_graph_stress()`, `layout_tbl_graph_treemap()`, `layout_tbl_graph_unrooted()`

---

 layout\_tbl\_graph\_metro

*Place nodes according to the standard design of metro maps*


---

### Description

This layout tries to optimise the placement of nodes so that they adhere to the classic look of metro maps. As such it optimizes on the distribution of incident edge angles, conformity of edge length, and edge angles in multiples of 45 degrees. As it works as a refinement of an already existing layout (the physical location of metro stations) it requires an a priori node location. Due to its purpose it probably works best with planar graphs.

### Usage

```
layout_tbl_graph_metro(
  graph,
  x,
  y,
  length = 2,
  grid_space = 0.0025,
  optimization_weights = NULL,
  max_movement = 5,
  circular = FALSE
)
```

### Arguments

graph	A tbl_graph object
x, y	The start location of the nodes
length	Desired multiple of grid point spacing. (length * grid_space determines desired edge length)
grid_space	The distance between consecutive grid points
optimization_weights	The relative weight to be placed on the 5 criteria during optimization as a numeric vector of length 4. The criteria are: <ul style="list-style-type: none"> <li>• Angular Resolution Criterion: The angles of incident edges at each station should be maximized, because if there is only a small angle between any two adjacent edges, then it can become difficult to distinguish between them.</li> <li>• Edge Length Criterion: The edge lengths across the whole map should be approximately equal to ensure regular spacing between stations. It is based on the preferred multiple, l, of the grid spacing, g. The purpose of the criterion is to penalize edges that are longer than or shorter than lg.</li> <li>• Balanced Edge Length Criterion: The length of edges incident to a particular station should be similar.</li> </ul>

- Line Straightness Criterion: (not yet implemented) Edges that form part of a line should, where possible, be co-linear either side of each station that the line passes through.
- Octilinearity Criterion: Each edge should be drawn horizontally, vertically, or diagonally at 45 degree, so we penalize edges that are not at a desired angle. If NULL all criteria are given equal weight.

max\_movement    Number of grid points a station can move away from its original position  
circular            ignored

### Value

A data.frame with the columns x, y, circular as well as any information stored as node variables in the tbl\_graph object.

### Author(s)

The underlying algorithm is implemented in the graphlayouts package by David Schoch

### References

Stott, J., Rodgers, P., Martinez-Ovando, J. C., and Walker, S. G. (2011). *Automatic metro map layout using multicriteria optimization* In IEEE Trans Vis Comput Graph 17(1) pp. 101-114. <https://doi.org/10.1109/tvcg.2010.24>

### See Also

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustree\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_partition

*Calculate nodes as areas dividing their parent*

---

### Description

The partition layout is a way to show hierarchical data in the same way as [layout\\_tbl\\_graph\\_treemap\(\)](#). Instead of subdividing the parent area the partition layout shows the division of a nodes children next to the area of the node itself. As such the node positions will be very reminiscent of a reingold-tilford tree layout but by plotting nodes as areas it better communicate the total weight of a node by summing up all its children. Often partition layouts are called icicle plots or sunburst diagrams (in case a radial transform is applied).

**Usage**

```
layout_tbl_graph_partition(
  graph,
  weight = NULL,
  circular = FALSE,
  height = NULL,
  sort.by = NULL,
  direction = "out",
  offset = pi/2,
  const.area = TRUE
)
```

**Arguments**

graph	An tbl_graph object
weight	An optional node variable to use as weight. Will only affect the weight of leaf nodes as the weight of non-leaf nodes are derived from their children.
circular	Logical. Should the layout be transformed to a circular representation. If TRUE the resulting layout will be a sunburst diagram.
height	An optional node variable to use as height. If NULL all nodes will be given a height of 1.
sort.by	The name of a node variable to sort the nodes by.
direction	The direction of the tree in the graph. 'out' (default) means that parents point towards their children, while 'in' means that children point towards their parent.
offset	If circular = TRUE, where should it begin. Defaults to pi/2 which is equivalent to 12 o'clock.
const.area	Logical. Should 'height' be scaled for area proportionality when using circular = TRUE. Defaults to TRUE.

**Value**

If circular = FALSE A data.frame with the columns x, y, width, height, leaf, depth, circular as well as any information stored as node variables in the tbl\_graph object. If circular = TRUE A data.frame with the columns x, y, r0, r, start, end, leaf, depth, circular as well as any information stored as node variables in the tbl\_graph object.

**Note**

partition is a layout intended for trees, that is, graphs where nodes only have one parent and zero or more children. If the provided graph does not fit this format an attempt to convert it to such a format will be made.

**References**

Kruskal, J. B., Landwehr, J. M. (1983). *Icicle Plots: Better Displays for Hierarchical Clustering*. American Statistician Vol 37(2), 162-168. <https://doi.org/10.2307/2685881>

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manua](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_pmds *Place nodes based on a multidimensional scaling of a set of pivot nodes*

---

**Description**

This layout is similar to the 'mds' layout but uses only a subset of pivot nodes for the mds calculation, making it considerably faster and thus suited for large graphs

**Usage**

```
layout_tbl_graph_pmds(graph, pivots, weights = NULL, circular = FALSE)
```

**Arguments**

graph	A tbl_graph object
pivots	The number of pivot nodes
weights	An expression evaluated on the edge data to provide edge weights for the layout. Currently ignored for the sparse version
circular	ignored

**Value**

A data.frame with the columns x, y, circular as well as any information stored as node variables in the tbl\_graph object.

**Author(s)**

The underlying algorithm is implemented in the graphlayouts package by David Schoch

**References**

Brandes, U. and Pich, C. (2006). *Eigensolver Methods for Progressive Multidimensional Scaling of Large Data*. In International Symposium on Graph Drawing (pp. 42-53). Springer

**See Also**

Other `layout_tbl_graph_*`: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manua](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted](#)

---

`layout_tbl_graph_sf`     *Place nodes on their geographical space*

---

**Description**

This layout is built for objects of class `sfnetwork` and is meant to plot a graph on its geographical space, by extracting its X and Y coordinates

**Usage**

```
layout_tbl_graph_sf(graph, circular = FALSE)
```

**Arguments**

<code>graph</code>	An <code>sfnetwork</code> object
<code>circular</code>	ignored

**Value**

A `data.frame` with the columns `x`, `y`, `circular`.

**See Also**

Other `layout_tbl_graph_*`: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustre](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manua](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted](#)



---

 layout\_tbl\_graph\_stress

*Place nodes using stress majorisation*


---

### Description

This layout is related to the stress-minimization algorithm known as Kamada-Kawai (available as the 'kk' layout), but uses another optimization strategy. It generally have better runtime, quality, and stability compared to the Kamada-Kawai layout and is thus generally preferred. The sparse version of the layout have better performance (especially on larger networks) at the expense of layout quality, but will generally outperform many other algorithms for large graphs in both runtime and quality (e.g. the 'drl' layout from igraph).

### Usage

```
layout_tbl_graph_stress(
  graph,
  weights = NULL,
  niter = 500,
  tolerance = 1e-04,
  mds = TRUE,
  bbox = 50,
  x = NULL,
  y = NULL,
  circular = FALSE
)

layout_tbl_graph_sparse_stress(
  graph,
  pivots,
  weights = NULL,
  niter = 500,
  circular = FALSE
)
```

### Arguments

graph	a tbl_graph object
weights	An expression evaluated on the edge data to provide edge weights for the layout. Currently ignored for the sparse version
niter	number of iterations during stress optimization
tolerance	stopping criterion for stress optimization
mds	should an MDS layout be used as initial layout (default: TRUE)
bbox	constrain dimension of output. Only relevant to determine the placement of disconnected graphs.

x, y	Expressions evaluated on the node data giving coordinates along x and/or y axis to fix nodes to. You can choose to only fix selected nodes by leaving the remaining nodes with NA values.
circular	ignored
pivots	The number of pivot nodes.

**Value**

A data.frame with the columns x, y, circular as well as any information stored as node variables in the tbl\_graph object.

**Author(s)**

The underlying algorithm is implemented in the graphlayouts package by David Schoch

**References**

Gansner, E. R., Koren, Y., & North, S. (2004). *Graph drawing by stress majorization*. In International Symposium on Graph Drawing (pp. 239-250). Springer, Berlin, Heidelberg.

Ortmann, M. and Klimenta, M. and Brandes, U. (2016). *A Sparse Stress Model*. <https://arxiv.org/pdf/1608.08909.pdf>

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustreemap\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

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layout\_tbl\_graph\_treemap

*Calculate nodes as rectangles subdividing that of their parent*

---

**Description**

A treemap is a space filling hierarchical layout that maps nodes to rectangles. The rectangles of the children of a node is packed into the rectangle of the node so that the size of a rectangle is a function of the size of the children. The size of the leaf nodes can be mapped arbitrarily (defaults to 1). Many different algorithms exist for dividing a rectangle into smaller bits, some optimizing the aspect ratio and some focusing on the ordering of the rectangles. See details for more discussions on this. The treemap layout was first developed by Ben Shneiderman for visualizing disk usage in the early '90 and has seen many improvements since.

**Usage**

```
layout_tbl_graph_treemap(
  graph,
  algorithm = "split",
  weight = NULL,
  circular = FALSE,
  sort.by = NULL,
  direction = "out",
  height = 1,
  width = 1
)
```

**Arguments**

graph	A <code>tbl_graph</code> object
algorithm	The name of the tiling algorithm to use. Defaults to 'split'
weight	An optional node variable to use as weight. Will only affect the weight of leaf nodes as the weight of non-leaf nodes are derived from their children.
circular	Logical. Should the layout be transformed to a circular representation. Ignored.
sort.by	The name of a node variables to sort the nodes by.
direction	The direction of the tree in the graph. 'out' (default) means that parents point towards their children, while 'in' means that children point towards their parent.
height	The height of the bounding rectangle
width	The width of the bounding rectangle

**Details**

Different approaches to dividing the rectangles in a treemap exists; all with their strengths and weaknesses. Currently only the split algorithm is implemented which strikes a good balance between aspect ratio and order preservation, but other, more well-known, algorithms such as squarify and slice-and-dice will eventually be implemented.

**Algorithms***Split* (default)

The Split algorithm was developed by Bjorn Engdahl in order to address the downsides of both the original slice-and-dice algorithm (poor aspect ratio) and the popular squarify algorithm (no ordering of nodes). It works by finding the best cut in the ordered list of children in terms of making sure that the two rectangles associated with the split will have optimal aspect ratio.

**Value**

A data.frame with the columns `x`, `y`, `width`, `height`, `leaf`, `depth`, `circular` as well as any information stored as node variables in the `tbl_graph` object.

**Note**

Treemap is a layout intended for trees, that is, graphs where nodes only have one parent and zero or more children. If the provided graph does not fit this format an attempt to convert it to such a format will be made.

**References**

Engdahl, B. (2005). *Ordered and unordered treemap algorithms and their applications on handheld devices*. Master's Degree Project.

Johnson, B., & Ben Shneiderman. (1991). *Tree maps: A Space-Filling Approach to the Visualization of Hierarchical Information Structures*. IEEE Visualization, 284-291. doi:10.1109/VISUAL.1991.175815

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustree\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_unrooted\(\)](#)

---

layout\_tbl\_graph\_unrooted

*Create an unrooted layout using equal-angle or equal-daylight*

---

**Description**

When drawing unrooted trees the standard dendrogram layout is a bad fit as it implicitly creates a visual root node. Instead it is possible to spread the leafs out on the plane without putting any special emphasis on a particular node using an unrooted layout. The standard algorithm is the equal angle algorithm, but it can struggle with optimising the leaf distribution for large trees with very uneven branch length. The equal daylight algorithm modifies the output of the equal angle algorithm to better disperse the leaves, at the cost of higher computational cost and the possibility of edge crossings for very large unbalanced trees. For standard sized trees the daylight algorithm is far superior and not too heavy so it is the default.

**Usage**

```
layout_tbl_graph_unrooted(
  graph,
  daylight = TRUE,
  length = NULL,
  tolerance = 0.05,
  rotation_mod = 1,
  maxiter = 100,
  circular = FALSE
)
```

**Arguments**

graph	A tbl_graph object
daylight	Should equal-daylight adjustments be made
length	An expression evaluating to the branch length of each edge
tolerance	The threshold for mean angular adjustment before terminating the daylight adjustment
rotation_mod	A modifier for the angular adjustment of each branch. Set it below 1 to let the daylight adjustment progress more slowly
maxiter	The maximum number of iterations in the the daylight adjustment
circular	ignored

**Value**

A data.frame with the columns x, y, circular, leaf as well as any information stored as node variables in the tbl\_graph object.

**Note**

Unrooted is a layout intended for undirected trees, that is, graphs with no cycles. If the provided graph does not fit this format an attempt to convert it to such a format will be made.

**References**

Felsenstein, J. (2004) *Drawing Trees*, in *Inferring Phylogenies*. Sinauer Assoc., pp 573-584

**See Also**

Other layout\_tbl\_graph\_\*: [layout\\_tbl\\_graph\\_auto\(\)](#), [layout\\_tbl\\_graph\\_backbone\(\)](#), [layout\\_tbl\\_graph\\_cactustree\(\)](#), [layout\\_tbl\\_graph\\_centrality\(\)](#), [layout\\_tbl\\_graph\\_circlepack\(\)](#), [layout\\_tbl\\_graph\\_dendrogram\(\)](#), [layout\\_tbl\\_graph\\_eigen\(\)](#), [layout\\_tbl\\_graph\\_fabric\(\)](#), [layout\\_tbl\\_graph\\_focus\(\)](#), [layout\\_tbl\\_graph\\_hive\(\)](#), [layout\\_tbl\\_graph\\_htree\(\)](#), [layout\\_tbl\\_graph\\_igraph\(\)](#), [layout\\_tbl\\_graph\\_linear\(\)](#), [layout\\_tbl\\_graph\\_manual\(\)](#), [layout\\_tbl\\_graph\\_matrix\(\)](#), [layout\\_tbl\\_graph\\_metro\(\)](#), [layout\\_tbl\\_graph\\_partition\(\)](#), [layout\\_tbl\\_graph\\_pmds\(\)](#), [layout\\_tbl\\_graph\\_sf\(\)](#), [layout\\_tbl\\_graph\\_stress\(\)](#), [layout\\_tbl\\_graph\\_treemap\(\)](#)

---

node\_angle

*Get the angle of nodes and edges*


---

**Description**

These helper functions makes it easy to calculate the angle associated with nodes and edges. For nodes the angle is defined as the angle of the vector pointing towards the node position, and is thus mainly suited for circular layouts where it can be used to calculate the angle of labels. For edges it is simply the angle of the vector describing the edge.

**Usage**

```
node_angle(x, y, degrees = TRUE, avoid_flip = TRUE)

edge_angle(x, y, xend, yend, degrees = TRUE, avoid_flip = TRUE)
```

**Arguments**

<code>x, y</code>	A vector of positions
<code>degrees</code>	Logical. Should the angle be returned in degree (TRUE) or radians (FALSE). Defaults to TRUE.
<code>avoid_flip</code>	Logical. Should the angle be adjusted so that text is always upside-down
<code>xend, yend</code>	The end position of the edge

**Value**

A vector with the angle of each node/edge

**Examples**

```
require(tidygraph)
flareGraph <- tbl_graph(flare$vertices, flare$edges)

ggraph(flareGraph, 'dendrogram', circular = TRUE) +
  geom_edge_diagonal0() +
  geom_node_text(aes(filter = leaf, angle = node_angle(x, y), label = shortName),
    hjust = 'outward', size = 2
  ) +
  expand_limits(x = c(-1.3, 1.3), y = c(-1.3, 1.3))
```

---

pack_circles	<i>Pack circles together</i>
--------------	------------------------------

---

**Description**

This function is a direct interface to the circle packing algorithm used by [layout\\_tbl\\_graph\\_circlepack](#). It takes a vector of sizes and returns the x and y position of each circle as a two-column matrix.

**Usage**

```
pack_circles(areas)
```

**Arguments**

<code>areas</code>	A vector of circle areas
--------------------	--------------------------

**Value**

A matrix with two columns and the same number of rows as the length of the "areas" vector. The matrix has the following attributes added: "enclosing\_radius" giving the radius of the smallest enclosing circle, and "front\_chain" giving the terminating members of the front chain (see Wang *et al.* 2006).

**References**

Wang, W., Wang, H. H., Dai, G., & Wang, H. (2006). *Visualization of large hierarchical data by circle packing*. *Chi*, 517-520.

**Examples**

```
library(ggforce)
sizes <- sample(10, 100, TRUE)

position <- pack_circles(sizes)
data <- data.frame(x = position[,1], y = position[,2], r = sqrt(sizes/pi))

ggplot() +
  geom_circle(aes(x0 = x, y0 = y, r = r), data = data, fill = 'steelblue') +
  geom_circle(aes(x0 = 0, y0 = 0, r = attr(position, 'enclosing_radius'))) +
  geom_polygon(aes(x = x, y = y),
              data = data[attr(position, 'front_chain'), ],
              fill = NA,
              colour = 'black')
```

---

scale\_edge\_alpha      *Edge alpha scales*

---

**Description**

This set of scales defines new alpha scales for edge geoms equivalent to the ones already defined by ggplot2. See [ggplot2::scale\\_alpha\(\)](#) for more information. The different geoms will know whether to use edge scales or the standard scales so it is not necessary to write edge\_alpha in the call to the geom - just use alpha.

**Usage**

```
scale_edge_alpha(..., range = c(0.1, 1))

scale_edge_alpha_continuous(..., range = c(0.1, 1))

scale_edge_alpha_discrete(..., range = c(0.1, 1))

scale_edge_alpha_binned(..., range = c(0.1, 1))
```

```
scale_edge_alpha_manual(..., values, breaks = waiver(), na.value = NA)
```

```
scale_edge_alpha_identity(..., guide = "none")
```

### Arguments

...	Other arguments passed on to <a href="#">continuous_scale()</a> , <a href="#">binned_scale()</a> , or <a href="#">discrete_scale()</a> as appropriate, to control name, limits, breaks, labels and so forth.
range	Output range of alpha values. Must lie between 0 and 1.
values	a set of aesthetic values to map data values to. The values will be matched in order (usually alphabetical) with the limits of the scale, or with breaks if provided. If this is a named vector, then the values will be matched based on the names instead. Data values that don't match will be given <code>na.value</code> .
breaks	One of: <ul style="list-style-type: none"> <li>• <code>NULL</code> for no breaks</li> <li>• <code>waiver()</code> for the default breaks (the scale limits)</li> <li>• A character vector of breaks</li> <li>• A function that takes the limits as input and returns breaks as output</li> </ul>
<code>na.value</code>	The aesthetic value to use for missing (NA) values
guide	Guide to use for this scale. Defaults to "none".

### Value

A ggproto object inheriting from `Scale`

### See Also

Other `scale_edge_*`: [scale\\_edge\\_colour](#), [scale\\_edge\\_fill](#), [scale\\_edge\\_linetype\(\)](#), [scale\\_edge\\_shape\(\)](#), [scale\\_edge\\_size\(\)](#), [scale\\_edge\\_width\(\)](#), [scale\\_label\\_size\(\)](#)

---

scale\_edge\_colour      *Edge colour scales*

---

### Description

This set of scales defines new colour scales for edge geoms equivalent to the ones already defined by `ggplot2`. The parameters are equivalent to the ones from `ggplot2` so there is nothing new under the sun. The different geoms will know whether to use edge scales or the standard scales so it is not necessary to write `edge_colour` in the call to the geom - just use `colour`.



**Usage**

```
scale_edge_colour_hue(  
  ...,  
  h = c(0, 360) + 15,  
  c = 100,  
  l = 65,  
  h.start = 0,  
  direction = 1,  
  na.value = "grey50",  
  aesthetics = "edge_colour"  
)  
  
scale_edge_colour_brewer(  
  ...,  
  type = "seq",  
  palette = 1,  
  direction = 1,  
  aesthetics = "edge_colour"  
)  
  
scale_edge_colour_distiller(  
  ...,  
  type = "seq",  
  palette = 1,  
  direction = -1,  
  values = NULL,  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_colour"  
)  
  
scale_edge_colour_gradient(  
  ...,  
  low = "#132B43",  
  high = "#56B1F7",  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_colour"  
)  
  
scale_edge_colour_gradient2(  
  ...,  
  low = muted("red"),  
  mid = "white",  
  high = muted("blue"),  
  midpoint = 0,
```

```
space = "Lab",
na.value = "grey50",
guide = "edge_colourbar",
aesthetics = "edge_colour"
)

scale_edge_colour_gradientn(
  ...,
  colours,
  values = NULL,
  space = "Lab",
  na.value = "grey50",
  guide = "edge_colourbar",
  aesthetics = "edge_colour",
  colors
)

scale_edge_colour_grey(
  ...,
  start = 0.2,
  end = 0.8,
  na.value = "red",
  aesthetics = "edge_colour"
)

scale_edge_colour_identity(..., guide = "none", aesthetics = "edge_colour")

scale_edge_colour_manual(
  ...,
  values,
  aesthetics = "edge_colour",
  breaks = waiver(),
  na.value = "grey50"
)

scale_edge_colour_viridis(
  ...,
  alpha = 1,
  begin = 0,
  end = 1,
  direction = 1,
  discrete = FALSE,
  option = "D",
  aesthetics = "edge_colour"
)

scale_edge_colour_steps(
  ...,
```

```
    low = "#132B43",
    high = "#56B1F7",
    space = "Lab",
    na.value = "grey50",
    guide = "edge_coloursteps",
    aesthetics = "edge_colour"
)

scale_edge_colour_steps2(
  ...,
  low = muted("red"),
  mid = "white",
  high = muted("blue"),
  midpoint = 0,
  space = "Lab",
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_colour"
)

scale_edge_colour_stepsn(
  ...,
  colours,
  values = NULL,
  space = "Lab",
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_colour",
  colors
)

scale_edge_colour_fermenter(
  ...,
  type = "seq",
  palette = 1,
  direction = -1,
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_colour"
)

scale_edge_colour_continuous(
  ...,
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
  guide = "edge_colourbar",
```

```
    aesthetics = "edge_colour"
  )

scale_edge_colour_discrete(
  ...,
  h = c(0, 360) + 15,
  c = 100,
  l = 65,
  h.start = 0,
  direction = 1,
  na.value = "grey50",
  aesthetics = "edge_colour"
)

scale_edge_colour_binned(
  ...,
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_colour"
)

scale_edge_color_hue(
  ...,
  h = c(0, 360) + 15,
  c = 100,
  l = 65,
  h.start = 0,
  direction = 1,
  na.value = "grey50",
  aesthetics = "edge_colour"
)

scale_edge_color_brewer(
  ...,
  type = "seq",
  palette = 1,
  direction = 1,
  aesthetics = "edge_colour"
)

scale_edge_color_distiller(
  ...,
  type = "seq",
  palette = 1,
  direction = -1,
```

```
    values = NULL,  
    space = "Lab",  
    na.value = "grey50",  
    guide = "edge_colourbar",  
    aesthetics = "edge_colour"  
  )  
  
scale_edge_color_gradient(  
  ...,  
  low = "#132B43",  
  high = "#56B1F7",  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_colour"  
)  
  
scale_edge_color_gradient2(  
  ...,  
  low = muted("red"),  
  mid = "white",  
  high = muted("blue"),  
  midpoint = 0,  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_colour"  
)  
  
scale_edge_color_gradientn(  
  ...,  
  colours,  
  values = NULL,  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_colour",  
  colors  
)  
  
scale_edge_color_grey(  
  ...,  
  start = 0.2,  
  end = 0.8,  
  na.value = "red",  
  aesthetics = "edge_colour"  
)
```

```
scale_edge_color_identity(..., guide = "none", aesthetics = "edge_colour")

scale_edge_color_manual(
  ...,
  values,
  aesthetics = "edge_colour",
  breaks = waiver(),
  na.value = "grey50"
)

scale_edge_color_continuous(
  ...,
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
  guide = "edge_colourbar",
  aesthetics = "edge_colour"
)

scale_edge_color_discrete(
  ...,
  h = c(0, 360) + 15,
  c = 100,
  l = 65,
  h.start = 0,
  direction = 1,
  na.value = "grey50",
  aesthetics = "edge_colour"
)

scale_edge_color_viridis(
  ...,
  alpha = 1,
  begin = 0,
  end = 1,
  direction = 1,
  discrete = FALSE,
  option = "D",
  aesthetics = "edge_colour"
)

scale_edge_color_steps(
  ...,
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
```

```
    guide = "edge_coloursteps",
    aesthetics = "edge_colour"
  )

scale_edge_color_steps2(
  ...,
  low = muted("red"),
  mid = "white",
  high = muted("blue"),
  midpoint = 0,
  space = "Lab",
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_colour"
)

scale_edge_color_stepsn(
  ...,
  colours,
  values = NULL,
  space = "Lab",
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_colour",
  colors
)

scale_edge_color_fermenter(
  ...,
  type = "seq",
  palette = 1,
  direction = -1,
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_colour"
)

scale_edge_color_binned(
  ...,
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_colour"
)
```

**Arguments**

...

Arguments passed on to [discrete\\_scale](#)

**palette** A palette function that when called with a single integer argument (the number of levels in the scale) returns the values that they should take (e.g., [scales::pal\\_hue\(\)](#)).

**breaks** One of:

- NULL for no breaks
- [waiver\(\)](#) for the default breaks (the scale limits)
- A character vector of breaks
- A function that takes the limits as input and returns breaks as output. Also accepts rlang [lambda](#) function notation.

**limits** One of:

- NULL to use the default scale values
- A character vector that defines possible values of the scale and their order
- A function that accepts the existing (automatic) values and returns new ones. Also accepts rlang [lambda](#) function notation.

**drop** Should unused factor levels be omitted from the scale? The default, TRUE, uses the levels that appear in the data; FALSE uses all the levels in the factor.

**na.translate** Unlike continuous scales, discrete scales can easily show missing values, and do so by default. If you want to remove missing values from a discrete scale, specify `na.translate = FALSE`.

**scale\_name** **[Deprecated]** The name of the scale that should be used for error messages associated with this scale.

**labels** One of:

- NULL for no labels
- [waiver\(\)](#) for the default labels computed by the transformation object
- A character vector giving labels (must be same length as breaks)
- An expression vector (must be the same length as breaks). See [?plot-math](#) for details.
- A function that takes the breaks as input and returns labels as output. Also accepts rlang [lambda](#) function notation.

**guide** A function used to create a guide or its name. See [guides\(\)](#) for more information.

**expand** For position scales, a vector of range expansion constants used to add some padding around the data to ensure that they are placed some distance away from the axes. Use the convenience function [expansion\(\)](#) to generate the values for the expand argument. The defaults are to expand the scale by 5% on each side for continuous variables, and by 0.6 units on each side for discrete variables.

**position** For position scales, The position of the axis. `left` or `right` for y axes, `top` or `bottom` for x axes.

**call** The call used to construct the scale for reporting messages.

**super** The super class to use for the constructed scale



h	range of hues to use, in [0, 360]
c	chroma (intensity of colour), maximum value varies depending on combination of hue and luminance.
l	luminance (lightness), in [0, 100]
h.start	hue to start at
direction	direction to travel around the colour wheel, 1 = clockwise, -1 = counter-clockwise
na.value	Colour to use for missing values
aesthetics	Character string or vector of character strings listing the name(s) of the aesthetic(s) that this scale works with. This can be useful, for example, to apply colour settings to the colour and fill aesthetics at the same time, via <code>aesthetics = c("colour", "fill")</code> .
type	One of "seq" (sequential), "div" (diverging) or "qual" (qualitative)
palette	If a string, will use that named palette. If a number, will index into the list of palettes of appropriate type. The list of available palettes can be found in the Palettes section.
values	if colours should not be evenly positioned along the gradient this vector gives the position (between 0 and 1) for each colour in the colours vector. See <a href="#">rescale()</a> for a convenience function to map an arbitrary range to between 0 and 1.
space	colour space in which to calculate gradient. Must be "Lab" - other values are deprecated.
guide	Type of legend. Use "colourbar" for continuous colour bar, or "legend" for discrete colour legend.
low, high	Colours for low and high ends of the gradient.
mid	colour for mid point
midpoint	The midpoint (in data value) of the diverging scale. Defaults to 0.
colours, colors	Vector of colours to use for n-colour gradient.
start	grey value at low end of palette
end	grey value at high end of palette
breaks	One of: <ul style="list-style-type: none"> <li>• NULL for no breaks</li> <li>• <code>waiver()</code> for the default breaks (the scale limits)</li> <li>• A character vector of breaks</li> <li>• A function that takes the limits as input and returns breaks as output</li> </ul>
alpha	The alpha transparency, a number in [0,1], see argument alpha in <a href="#">hsv</a> .
begin	The (corrected) hue in [0,1] at which the color map begins.
discrete	Generate a discrete palette? (default: FALSE - generate continuous palette).
option	A character string indicating the color map option to use. Eight options are available: <ul style="list-style-type: none"> <li>• "magma" (or "A")</li> </ul>

- "inferno" (or "B")
- "plasma" (or "C")
- "viridis" (or "D")
- "cividis" (or "E")
- "rocket" (or "F")
- "mako" (or "G")
- "turbo" (or "H")

### Value

A ggproto object inheriting from Scale

### See Also

Other scale\_edge\_\*: [scale\\_edge\\_alpha\(\)](#), [scale\\_edge\\_fill](#), [scale\\_edge\\_linetype\(\)](#), [scale\\_edge\\_shape\(\)](#), [scale\\_edge\\_size\(\)](#), [scale\\_edge\\_width\(\)](#), [scale\\_label\\_size\(\)](#)

---

scale_edge_fill	<i>Edge fill scales</i>
-----------------	-------------------------

---

### Description

This set of scales defines new fill scales for edge geoms equivalent to the ones already defined by ggplot2. The parameters are equivalent to the ones from ggplot2 so there is nothing new under the sun. The different geoms will know whether to use edge scales or the standard scales so it is not necessary to write edge\_fill in the call to the geom - just use fill.

### Usage

```
scale_edge_fill_hue(
  ...,
  h = c(0, 360) + 15,
  c = 100,
  l = 65,
  h.start = 0,
  direction = 1,
  na.value = "grey50",
  aesthetics = "edge_fill"
)
```

```
scale_edge_fill_brewer(
  ...,
  type = "seq",
  palette = 1,
  direction = 1,
  aesthetics = "edge_fill"
```

```
)  
  
scale_edge_fill_distiller(  
  ...,  
  type = "seq",  
  palette = 1,  
  direction = -1,  
  values = NULL,  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_fill"  
)  
  
scale_edge_fill_gradient(  
  ...,  
  low = "#132B43",  
  high = "#56B1F7",  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_fill"  
)  
  
scale_edge_fill_gradient2(  
  ...,  
  low = muted("red"),  
  mid = "white",  
  high = muted("blue"),  
  midpoint = 0,  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_fill"  
)  
  
scale_edge_fill_gradientn(  
  ...,  
  colours,  
  values = NULL,  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_fill",  
  colors  
)  
  
scale_edge_fill_grey(  
  ...,  
  values = NULL,  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_fill"  
)
```

```
    ...,
    start = 0.2,
    end = 0.8,
    na.value = "red",
    aesthetics = "edge_fill"
  )

scale_edge_fill_identity(..., guide = "none", aesthetics = "edge_fill")

scale_edge_fill_manual(
  ...,
  values,
  aesthetics = "edge_fill",
  breaks = waiver(),
  na.value = "grey50"
)

scale_edge_fill_viridis(
  ...,
  alpha = 1,
  begin = 0,
  end = 1,
  direction = 1,
  discrete = FALSE,
  option = "D",
  aesthetics = "edge_fill"
)

scale_edge_fill_steps(
  ...,
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_fill"
)

scale_edge_fill_steps2(
  ...,
  low = muted("red"),
  mid = "white",
  high = muted("blue"),
  midpoint = 0,
  space = "Lab",
  na.value = "grey50",
  guide = "edge_coloursteps",
  aesthetics = "edge_fill"
)
```

```
)  
  
scale_edge_fill_stepsn(  
  ...,  
  colours,  
  values = NULL,  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_coloursteps",  
  aesthetics = "edge_fill",  
  colors  
)  
  
scale_edge_fill_fermenter(  
  ...,  
  type = "seq",  
  palette = 1,  
  direction = -1,  
  na.value = "grey50",  
  guide = "edge_coloursteps",  
  aesthetics = "edge_fill"  
)  
  
scale_edge_fill_continuous(  
  ...,  
  low = "#132B43",  
  high = "#56B1F7",  
  space = "Lab",  
  na.value = "grey50",  
  guide = "edge_colourbar",  
  aesthetics = "edge_fill"  
)  
  
scale_edge_fill_discrete(  
  ...,  
  h = c(0, 360) + 15,  
  c = 100,  
  l = 65,  
  h.start = 0,  
  direction = 1,  
  na.value = "grey50",  
  aesthetics = "edge_fill"  
)  
  
scale_edge_fill_binned(  
  ...,  
  low = "#132B43",  
  high = "#56B1F7",
```

```

space = "Lab",
na.value = "grey50",
guide = "edge_coloursteps",
aesthetics = "edge_fill"
)

```

## Arguments

- ... Arguments passed on to [discrete\\_scale](#)
- palette** A palette function that when called with a single integer argument (the number of levels in the scale) returns the values that they should take (e.g., [scales::pal\\_hue\(\)](#)).
- breaks** One of:
- NULL for no breaks
  - [waiver\(\)](#) for the default breaks (the scale limits)
  - A character vector of breaks
  - A function that takes the limits as input and returns breaks as output. Also accepts rlang [lambda](#) function notation.
- limits** One of:
- NULL to use the default scale values
  - A character vector that defines possible values of the scale and their order
  - A function that accepts the existing (automatic) values and returns new ones. Also accepts rlang [lambda](#) function notation.
- drop** Should unused factor levels be omitted from the scale? The default, TRUE, uses the levels that appear in the data; FALSE uses all the levels in the factor.
- na.translate** Unlike continuous scales, discrete scales can easily show missing values, and do so by default. If you want to remove missing values from a discrete scale, specify `na.translate = FALSE`.
- scale\_name** **[Deprecated]** The name of the scale that should be used for error messages associated with this scale.
- labels** One of:
- NULL for no labels
  - [waiver\(\)](#) for the default labels computed by the transformation object
  - A character vector giving labels (must be same length as breaks)
  - An expression vector (must be the same length as breaks). See [?plot-math](#) for details.
  - A function that takes the breaks as input and returns labels as output. Also accepts rlang [lambda](#) function notation.
- guide** A function used to create a guide or its name. See [guides\(\)](#) for more information.
- expand** For position scales, a vector of range expansion constants used to add some padding around the data to ensure that they are placed some distance away from the axes. Use the convenience function [expansion\(\)](#) to generate the values for the expand argument. The defaults are to expand the

	scale by 5% on each side for continuous variables, and by 0.6 units on each side for discrete variables.
position	For position scales, The position of the axis. left or right for y axes, top or bottom for x axes.
call	The call used to construct the scale for reporting messages.
super	The super class to use for the constructed scale
h	range of hues to use, in [0, 360]
c	chroma (intensity of colour), maximum value varies depending on combination of hue and luminance.
l	luminance (lightness), in [0, 100]
h.start	hue to start at
direction	direction to travel around the colour wheel, 1 = clockwise, -1 = counter-clockwise
na.value	Colour to use for missing values
aesthetics	Character string or vector of character strings listing the name(s) of the aesthetic(s) that this scale works with. This can be useful, for example, to apply colour settings to the colour and fill aesthetics at the same time, via aesthetics = c("colour", "fill").
type	One of "seq" (sequential), "div" (diverging) or "qual" (qualitative)
palette	If a string, will use that named palette. If a number, will index into the list of palettes of appropriate type. The list of available palettes can found in the Palettes section.
values	if colours should not be evenly positioned along the gradient this vector gives the position (between 0 and 1) for each colour in the colours vector. See <a href="#">rescale()</a> for a convenience function to map an arbitrary range to between 0 and 1.
space	colour space in which to calculate gradient. Must be "Lab" - other values are deprecated.
guide	Type of legend. Use "colourbar" for continuous colour bar, or "legend" for discrete colour legend.
low, high	Colours for low and high ends of the gradient.
mid	colour for mid point
midpoint	The midpoint (in data value) of the diverging scale. Defaults to 0.
colours, colors	Vector of colours to use for n-colour gradient.
start	grey value at low end of palette
end	grey value at high end of palette
breaks	One of: <ul style="list-style-type: none"> <li>• NULL for no breaks</li> <li>• waiver() for the default breaks (the scale limits)</li> <li>• A character vector of breaks</li> <li>• A function that takes the limits as input and returns breaks as output</li> </ul>

alpha	The alpha transparency, a number in [0,1], see argument alpha in <a href="#">hsv</a> .
begin	The (corrected) hue in [0,1] at which the color map begins.
discrete	Generate a discrete palette? (default: FALSE - generate continuous palette).
option	A character string indicating the color map option to use. Eight options are available: <ul style="list-style-type: none"> <li>• "magma" (or "A")</li> <li>• "inferno" (or "B")</li> <li>• "plasma" (or "C")</li> <li>• "viridis" (or "D")</li> <li>• "cividis" (or "E")</li> <li>• "rocket" (or "F")</li> <li>• "mako" (or "G")</li> <li>• "turbo" (or "H")</li> </ul>

**Value**

A ggproto object inheriting from Scale

**See Also**

Other scale\_edge\_\*: [scale\\_edge\\_alpha\(\)](#), [scale\\_edge\\_colour](#), [scale\\_edge\\_linetype\(\)](#), [scale\\_edge\\_shape\(\)](#), [scale\\_edge\\_size\(\)](#), [scale\\_edge\\_width\(\)](#), [scale\\_label\\_size\(\)](#)

---

scale\_edge\_linetype    *Edge linetype scales*

---

**Description**

This set of scales defines new linetype scales for edge geoms equivalent to the ones already defined by ggplot2. See [ggplot2::scale\\_linetype\(\)](#) for more information. The different geoms will know whether to use edge scales or the standard scales so it is not necessary to write edge\_linetype in the call to the geom - just use linetype.

**Usage**

```
scale_edge_linetype(..., na.value = "blank")
```

```
scale_edge_linetype_continuous(...)
```

```
scale_edge_linetype_discrete(..., na.value = "blank")
```

```
scale_edge_linetype_binned(..., na.value = "blank")
```

```
scale_edge_linetype_manual(..., values, breaks = waiver(), na.value = "blank")
```

```
scale_edge_linetype_identity(..., guide = "none")
```



**Arguments**

...	Arguments passed on to <code>discrete_scale</code>
<code>palette</code>	A palette function that when called with a single integer argument (the number of levels in the scale) returns the values that they should take (e.g., <code>scales::pal_hue()</code> ).
<code>breaks</code>	One of: <ul style="list-style-type: none"> <li>• NULL for no breaks</li> <li>• <code>waiver()</code> for the default breaks (the scale limits)</li> <li>• A character vector of breaks</li> <li>• A function that takes the limits as input and returns breaks as output. Also accepts rlang <code>lambda</code> function notation.</li> </ul>
<code>limits</code>	One of: <ul style="list-style-type: none"> <li>• NULL to use the default scale values</li> <li>• A character vector that defines possible values of the scale and their order</li> <li>• A function that accepts the existing (automatic) values and returns new ones. Also accepts rlang <code>lambda</code> function notation.</li> </ul>
<code>drop</code>	Should unused factor levels be omitted from the scale? The default, TRUE, uses the levels that appear in the data; FALSE uses all the levels in the factor.
<code>na.translate</code>	Unlike continuous scales, discrete scales can easily show missing values, and do so by default. If you want to remove missing values from a discrete scale, specify <code>na.translate = FALSE</code> .
<code>aesthetics</code>	The names of the aesthetics that this scale works with.
<code>scale_name</code>	<b>[Deprecated]</b> The name of the scale that should be used for error messages associated with this scale.
<code>labels</code>	One of: <ul style="list-style-type: none"> <li>• NULL for no labels</li> <li>• <code>waiver()</code> for the default labels computed by the transformation object</li> <li>• A character vector giving labels (must be same length as breaks)</li> <li>• An expression vector (must be the same length as breaks). See <code>?plot-math</code> for details.</li> <li>• A function that takes the breaks as input and returns labels as output. Also accepts rlang <code>lambda</code> function notation.</li> </ul>
<code>guide</code>	A function used to create a guide or its name. See <code>guides()</code> for more information.
<code>call</code>	The call used to construct the scale for reporting messages.
<code>super</code>	The super class to use for the constructed scale
<code>na.value</code>	The linetype to use for NA values.
<code>values</code>	a set of aesthetic values to map data values to. The values will be matched in order (usually alphabetical) with the limits of the scale, or with breaks if provided. If this is a named vector, then the values will be matched based on the names instead. Data values that don't match will be given <code>na.value</code> .
<code>breaks</code>	One of:

- NULL for no breaks
  - `waiver()` for the default breaks (the scale limits)
  - A character vector of breaks
  - A function that takes the limits as input and returns breaks as output
- guide Guide to use for this scale. Defaults to "none".

**Value**

A ggproto object inheriting from `Scale`

**See Also**

Other `scale_edge_*`: [scale\\_edge\\_alpha\(\)](#), [scale\\_edge\\_colour](#), [scale\\_edge\\_fill](#), [scale\\_edge\\_shape\(\)](#), [scale\\_edge\\_size\(\)](#), [scale\\_edge\\_width\(\)](#), [scale\\_label\\_size\(\)](#)

---

scale\_edge\_shape      *Edge shape scales*

---

**Description**

This set of scales defines new shape scales for edge geoms equivalent to the ones already defined by `ggplot2`. See [ggplot2::scale\\_shape\(\)](#) for more information. The different geoms will know whether to use edge scales or the standard scales so it is not necessary to write `edge_shape` in the call to the geom - just use `shape`.

**Usage**

```
scale_edge_shape(..., solid = TRUE)

scale_edge_shape_discrete(..., solid = TRUE)

scale_edge_shape_continuous(...)

scale_edge_shape_binned(..., solid = TRUE)

scale_edge_shape_manual(..., values, breaks = waiver(), na.value = NA)

scale_edge_shape_identity(..., guide = "none")
```

**Arguments**

... Arguments passed on to [discrete\\_scale](#)

palette A palette function that when called with a single integer argument (the number of levels in the scale) returns the values that they should take (e.g., [scales::pal\\_hue\(\)](#)).

breaks One of:

- NULL for no breaks
- `waiver()` for the default breaks (the scale limits)
- A character vector of breaks
- A function that takes the limits as input and returns breaks as output. Also accepts rlang `lambda` function notation.

`limits` One of:

- NULL to use the default scale values
- A character vector that defines possible values of the scale and their order
- A function that accepts the existing (automatic) values and returns new ones. Also accepts rlang `lambda` function notation.

`drop` Should unused factor levels be omitted from the scale? The default, TRUE, uses the levels that appear in the data; FALSE uses all the levels in the factor.

`na.translate` Unlike continuous scales, discrete scales can easily show missing values, and do so by default. If you want to remove missing values from a discrete scale, specify `na.translate = FALSE`.

`na.value` If `na.translate = TRUE`, what aesthetic value should the missing values be displayed as? Does not apply to position scales where NA is always placed at the far right.

`aesthetics` The names of the aesthetics that this scale works with.

`scale_name` **[Deprecated]** The name of the scale that should be used for error messages associated with this scale.

`labels` One of:

- NULL for no labels
- `waiver()` for the default labels computed by the transformation object
- A character vector giving labels (must be same length as breaks)
- An expression vector (must be the same length as breaks). See `?plot-math` for details.
- A function that takes the breaks as input and returns labels as output. Also accepts rlang `lambda` function notation.

`guide` A function used to create a guide or its name. See `guides()` for more information.

`call` The call used to construct the scale for reporting messages.

`super` The super class to use for the constructed scale

`solid` Should the shapes be solid, TRUE, or hollow, FALSE?

`values` a set of aesthetic values to map data values to. The values will be matched in order (usually alphabetical) with the limits of the scale, or with breaks if provided. If this is a named vector, then the values will be matched based on the names instead. Data values that don't match will be given `na.value`.

`breaks` One of:

- NULL for no breaks
- `waiver()` for the default breaks (the scale limits)
- A character vector of breaks
- A function that takes the limits as input and returns breaks as output

na.value      The aesthetic value to use for missing (NA) values  
 guide         Guide to use for this scale.

**Value**

A ggproto object inheriting from Scale

**See Also**

Other scale\_edge\_\*: [scale\\_edge\\_alpha\(\)](#), [scale\\_edge\\_colour](#), [scale\\_edge\\_fill](#), [scale\\_edge\\_linetype\(\)](#), [scale\\_edge\\_size\(\)](#), [scale\\_edge\\_width\(\)](#), [scale\\_label\\_size\(\)](#)

---

scale\_edge\_size      *Edge size scales*

---

**Description**

This set of scales defines new size scales for edge geoms equivalent to the ones already defined by ggplot2. See [ggplot2::scale\\_size\(\)](#) for more information. The different geoms will know whether to use edge scales or the standard scales so it is not necessary to write edge\_size in the call to the geom - just use size.

**Usage**

```
scale_edge_size_continuous(  
  name = waiver(),  
  breaks = waiver(),  
  labels = waiver(),  
  limits = NULL,  
  range = c(1, 6),  
  trans = "identity",  
  guide = "legend"  
)
```

```
scale_edge_radius(  
  name = waiver(),  
  breaks = waiver(),  
  labels = waiver(),  
  limits = NULL,  
  range = c(1, 6),  
  trans = "identity",  
  guide = "legend"  
)
```

```
scale_edge_size(  
  name = waiver(),  
  breaks = waiver(),
```

```

  labels = waiver(),
  limits = NULL,
  range = c(1, 6),
  trans = "identity",
  guide = "legend"
)

scale_edge_size_discrete(...)

scale_edge_size_binned(
  name = waiver(),
  breaks = waiver(),
  labels = waiver(),
  limits = NULL,
  range = c(1, 6),
  n.breaks = NULL,
  nice.breaks = TRUE,
  trans = "identity",
  guide = "bins"
)

scale_edge_size_area(..., max_size = 6)

scale_edge_size_binned_area(..., max_size = 6)

scale_edge_size_manual(..., values, breaks = waiver(), na.value = NA)

scale_edge_size_identity(..., guide = "none")

```

## Arguments

name	The name of the scale. Used as the axis or legend title. If <code>waiver()</code> , the default, the name of the scale is taken from the first mapping used for that aesthetic. If <code>NULL</code> , the legend title will be omitted.
breaks	One of: <ul style="list-style-type: none"> <li>• <code>NULL</code> for no breaks</li> <li>• <code>waiver()</code> for the default breaks computed by the <a href="#">transformation object</a></li> <li>• A numeric vector of positions</li> <li>• A function that takes the limits as input and returns breaks as output (e.g., a function returned by <code>scales::extended_breaks()</code>). Also accepts <code>rlang</code> <a href="#">lambda</a> function notation.</li> </ul>
labels	One of: <ul style="list-style-type: none"> <li>• <code>NULL</code> for no labels</li> <li>• <code>waiver()</code> for the default labels computed by the transformation object</li> <li>• A character vector giving labels (must be same length as breaks)</li> <li>• An expression vector (must be the same length as breaks). See <code>?plotmath</code> for details.</li> </ul>

	<ul style="list-style-type: none"> <li>• A function that takes the breaks as input and returns labels as output. Also accepts rlang <a href="#">lambda</a> function notation.</li> </ul>
limits	<p>One of:</p> <ul style="list-style-type: none"> <li>• NULL to use the default scale range</li> <li>• A numeric vector of length two providing limits of the scale. Use NA to refer to the existing minimum or maximum</li> <li>• A function that accepts the existing (automatic) limits and returns new limits. Also accepts rlang <a href="#">lambda</a> function notation. Note that setting limits on positional scales will <b>remove</b> data outside of the limits. If the purpose is to zoom, use the limit argument in the coordinate system (see <a href="#">coord_cartesian()</a>).</li> </ul>
range	a numeric vector of length 2 that specifies the minimum and maximum size of the plotting symbol after transformation.
trans	<b>[Deprecated]</b> Deprecated in favour of transform.
guide	A function used to create a guide or its name. See <a href="#">guides()</a> for more information.
...	Arguments passed on to <a href="#">continuous_scale</a>
minor_breaks	<p>One of:</p> <ul style="list-style-type: none"> <li>• NULL for no minor breaks</li> <li>• <a href="#">waiver()</a> for the default breaks (one minor break between each major break)</li> <li>• A numeric vector of positions</li> <li>• A function that given the limits returns a vector of minor breaks. Also accepts rlang <a href="#">lambda</a> function notation. When the function has two arguments, it will be given the limits and major breaks.</li> </ul>
oob	<p>One of:</p> <ul style="list-style-type: none"> <li>• Function that handles limits outside of the scale limits (out of bounds). Also accepts rlang <a href="#">lambda</a> function notation.</li> <li>• The default (<a href="#">scales:::censor()</a>) replaces out of bounds values with NA.</li> <li>• <a href="#">scales:::squish()</a> for squishing out of bounds values into range.</li> <li>• <a href="#">scales:::squish_infinite()</a> for squishing infinite values into range.</li> </ul>
na.value	Missing values will be replaced with this value.
expand	For position scales, a vector of range expansion constants used to add some padding around the data to ensure that they are placed some distance away from the axes. Use the convenience function <a href="#">expansion()</a> to generate the values for the expand argument. The defaults are to expand the scale by 5% on each side for continuous variables, and by 0.6 units on each side for discrete variables.
position	For position scales, The position of the axis. left or right for y axes, top or bottom for x axes.
call	The call used to construct the scale for reporting messages.
super	The super class to use for the constructed scale

n.breaks	An integer guiding the number of major breaks. The algorithm may choose a slightly different number to ensure nice break labels. Will only have an effect if breaks = waiver(). Use NULL to use the default number of breaks given by the transformation.
nice.breaks	Logical. Should breaks be attempted placed at nice values instead of exactly evenly spaced between the limits. If TRUE (default) the scale will ask the transformation object to create breaks, and this may result in a different number of breaks than requested. Ignored if breaks are given explicitly.
max_size	Size of largest points.
values	a set of aesthetic values to map data values to. The values will be matched in order (usually alphabetical) with the limits of the scale, or with breaks if provided. If this is a named vector, then the values will be matched based on the names instead. Data values that don't match will be given na.value.
na.value	The aesthetic value to use for missing (NA) values

**Value**

A ggproto object inheriting from Scale

**Note**

In ggplot2 size conflates both line width and point size into one scale. In ggraph there is also a width scale ([scale\\_edge\\_width\(\)](#)) that is used for linewidth. As edges are often represented by lines the width scale is the most common.

**See Also**

Other scale\_edge\_\*: [scale\\_edge\\_alpha\(\)](#), [scale\\_edge\\_colour](#), [scale\\_edge\\_fill](#), [scale\\_edge\\_linetype\(\)](#), [scale\\_edge\\_shape\(\)](#), [scale\\_edge\\_width\(\)](#), [scale\\_label\\_size\(\)](#)

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scale_edge_width	<i>Edge width scales</i>
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---

**Description**

This set of scales defines width scales for edge geoms. Of all the new edge scales defined in ggraph, this is the only one not having an equivalent in ggplot2. In essence it mimics the use of size in [ggplot2::geom\\_line\(\)](#) and related. As almost all edge representations are lines of some sort, edge\_width will be used much more often than edge\_size. It is not necessary to spell out that it is an edge scale as the geom knows if it is drawing an edge. Just write width and not edge\_width in the call to geoms.

**Usage**

```
scale_edge_width_continuous(  
  name = waiver(),  
  breaks = waiver(),  
  labels = waiver(),  
  limits = NULL,  
  range = c(1, 6),  
  trans = "identity",  
  guide = "legend"  
)  
  
scale_edge_width(  
  name = waiver(),  
  breaks = waiver(),  
  labels = waiver(),  
  limits = NULL,  
  range = c(1, 6),  
  trans = "identity",  
  guide = "legend"  
)  
  
scale_edge_width_discrete(...)  
  
scale_edge_width_binned(  
  name = waiver(),  
  breaks = waiver(),  
  labels = waiver(),  
  limits = NULL,  
  range = c(1, 6),  
  n.breaks = NULL,  
  nice.breaks = TRUE,  
  trans = "identity",  
  guide = "bins"  
)  
  
scale_edge_width_manual(..., values, breaks = waiver(), na.value = NA)  
  
scale_edge_width_identity(..., guide = "none")
```

**Arguments**

name	The name of the scale. Used as the axis or legend title. If <code>waiver()</code> , the default, the name of the scale is taken from the first mapping used for that aesthetic. If <code>NULL</code> , the legend title will be omitted.
breaks	One of: <ul style="list-style-type: none"><li>• <code>NULL</code> for no breaks</li><li>• <code>waiver()</code> for the default breaks computed by the <a href="#">transformation object</a></li></ul>



	<ul style="list-style-type: none"> <li>• A numeric vector of positions</li> <li>• A function that takes the limits as input and returns breaks as output (e.g., a function returned by <code>scales::extended_breaks()</code>). Also accepts rlang <code>lambda</code> function notation.</li> </ul>
labels	<p>One of:</p> <ul style="list-style-type: none"> <li>• NULL for no labels</li> <li>• <code>waiver()</code> for the default labels computed by the transformation object</li> <li>• A character vector giving labels (must be same length as breaks)</li> <li>• An expression vector (must be the same length as breaks). See <code>?plotmath</code> for details.</li> <li>• A function that takes the breaks as input and returns labels as output. Also accepts rlang <code>lambda</code> function notation.</li> </ul>
limits	<p>One of:</p> <ul style="list-style-type: none"> <li>• NULL to use the default scale range</li> <li>• A numeric vector of length two providing limits of the scale. Use NA to refer to the existing minimum or maximum</li> <li>• A function that accepts the existing (automatic) limits and returns new limits. Also accepts rlang <code>lambda</code> function notation. Note that setting limits on positional scales will <b>remove</b> data outside of the limits. If the purpose is to zoom, use the limit argument in the coordinate system (see <code>coord_cartesian()</code>).</li> </ul>
range	a numeric vector of length 2 that specifies the minimum and maximum size of the plotting symbol after transformation.
trans	<b>[Deprecated]</b> Deprecated in favour of <code>transform</code> .
guide	A function used to create a guide or its name. See <code>guides()</code> for more information.
...	Arguments passed on to <code>continuous_scale</code>
minor_breaks	<p>One of:</p> <ul style="list-style-type: none"> <li>• NULL for no minor breaks</li> <li>• <code>waiver()</code> for the default breaks (one minor break between each major break)</li> <li>• A numeric vector of positions</li> <li>• A function that given the limits returns a vector of minor breaks. Also accepts rlang <code>lambda</code> function notation. When the function has two arguments, it will be given the limits and major breaks.</li> </ul>
oob	<p>One of:</p> <ul style="list-style-type: none"> <li>• Function that handles limits outside of the scale limits (out of bounds). Also accepts rlang <code>lambda</code> function notation.</li> <li>• The default (<code>scales::censor()</code>) replaces out of bounds values with NA.</li> <li>• <code>scales::squish()</code> for squishing out of bounds values into range.</li> <li>• <code>scales::squish_infinite()</code> for squishing infinite values into range.</li> </ul>
na.value	Missing values will be replaced with this value.

expand	For position scales, a vector of range expansion constants used to add some padding around the data to ensure that they are placed some distance away from the axes. Use the convenience function <code>expansion()</code> to generate the values for the <code>expand</code> argument. The defaults are to expand the scale by 5% on each side for continuous variables, and by 0.6 units on each side for discrete variables.
position	For position scales, The position of the axis. <code>left</code> or <code>right</code> for y axes, <code>top</code> or <code>bottom</code> for x axes.
call	The call used to construct the scale for reporting messages.
super	The super class to use for the constructed scale
n.breaks	An integer guiding the number of major breaks. The algorithm may choose a slightly different number to ensure nice break labels. Will only have an effect if <code>breaks = waiver()</code> . Use <code>NULL</code> to use the default number of breaks given by the transformation.
nice.breaks	Logical. Should breaks be attempted placed at nice values instead of exactly evenly spaced between the limits. If <code>TRUE</code> (default) the scale will ask the transformation object to create breaks, and this may result in a different number of breaks than requested. Ignored if breaks are given explicitly.
values	a set of aesthetic values to map data values to. The values will be matched in order (usually alphabetical) with the limits of the scale, or with breaks if provided. If this is a named vector, then the values will be matched based on the names instead. Data values that don't match will be given <code>na.value</code> .
na.value	The aesthetic value to use for missing (NA) values

**Value**

A ggproto object inheriting from `Scale`

**See Also**

Other `scale_edge_*`: [scale\\_edge\\_alpha\(\)](#), [scale\\_edge\\_colour](#), [scale\\_edge\\_fill](#), [scale\\_edge\\_linetype\(\)](#), [scale\\_edge\\_shape\(\)](#), [scale\\_edge\\_size\(\)](#), [scale\\_label\\_size\(\)](#)

---

scale_label_size	<i>Edge label size scales</i>
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---

**Description**

This set of scales defines new size scales for edge labels in order to allow for separate sizing of edges and their labels.

**Usage**

```
scale_label_size_continuous(  
  name = waiver(),  
  breaks = waiver(),  
  labels = waiver(),  
  limits = NULL,  
  range = c(1, 6),  
  trans = "identity",  
  guide = "legend"  
)  
  
scale_label_size(  
  name = waiver(),  
  breaks = waiver(),  
  labels = waiver(),  
  limits = NULL,  
  range = c(1, 6),  
  trans = "identity",  
  guide = "legend"  
)  
  
scale_label_size_discrete(...)  
  
scale_label_size_binned(  
  name = waiver(),  
  breaks = waiver(),  
  labels = waiver(),  
  limits = NULL,  
  range = c(1, 6),  
  n.breaks = NULL,  
  nice.breaks = TRUE,  
  trans = "identity",  
  guide = "bins"  
)  
  
scale_label_size_manual(..., values, breaks = waiver(), na.value = NA)  
  
scale_label_size_identity(..., guide = "none")
```

**Arguments**

name	The name of the scale. Used as the axis or legend title. If <code>waiver()</code> , the default, the name of the scale is taken from the first mapping used for that aesthetic. If <code>NULL</code> , the legend title will be omitted.
breaks	One of: <ul style="list-style-type: none"><li>• <code>NULL</code> for no breaks</li><li>• <code>waiver()</code> for the default breaks computed by the <a href="#">transformation object</a></li></ul>

	<ul style="list-style-type: none"> <li>• A numeric vector of positions</li> <li>• A function that takes the limits as input and returns breaks as output (e.g., a function returned by <code>scales::extended_breaks()</code>). Also accepts rlang <code>lambda</code> function notation.</li> </ul>
labels	<p>One of:</p> <ul style="list-style-type: none"> <li>• NULL for no labels</li> <li>• <code>waiver()</code> for the default labels computed by the transformation object</li> <li>• A character vector giving labels (must be same length as breaks)</li> <li>• An expression vector (must be the same length as breaks). See <code>?plotmath</code> for details.</li> <li>• A function that takes the breaks as input and returns labels as output. Also accepts rlang <code>lambda</code> function notation.</li> </ul>
limits	<p>One of:</p> <ul style="list-style-type: none"> <li>• NULL to use the default scale range</li> <li>• A numeric vector of length two providing limits of the scale. Use NA to refer to the existing minimum or maximum</li> <li>• A function that accepts the existing (automatic) limits and returns new limits. Also accepts rlang <code>lambda</code> function notation. Note that setting limits on positional scales will <b>remove</b> data outside of the limits. If the purpose is to zoom, use the limit argument in the coordinate system (see <code>coord_cartesian()</code>).</li> </ul>
range	a numeric vector of length 2 that specifies the minimum and maximum size of the plotting symbol after transformation.
trans	<b>[Deprecated]</b> Deprecated in favour of <code>transform</code> .
guide	A function used to create a guide or its name. See <code>guides()</code> for more information.
...	Arguments passed on to <code>continuous_scale</code>
minor_breaks	<p>One of:</p> <ul style="list-style-type: none"> <li>• NULL for no minor breaks</li> <li>• <code>waiver()</code> for the default breaks (one minor break between each major break)</li> <li>• A numeric vector of positions</li> <li>• A function that given the limits returns a vector of minor breaks. Also accepts rlang <code>lambda</code> function notation. When the function has two arguments, it will be given the limits and major breaks.</li> </ul>
oob	<p>One of:</p> <ul style="list-style-type: none"> <li>• Function that handles limits outside of the scale limits (out of bounds). Also accepts rlang <code>lambda</code> function notation.</li> <li>• The default (<code>scales::censor()</code>) replaces out of bounds values with NA.</li> <li>• <code>scales::squish()</code> for squishing out of bounds values into range.</li> <li>• <code>scales::squish_infinite()</code> for squishing infinite values into range.</li> </ul>
na.value	Missing values will be replaced with this value.

	<p><code>expand</code> For position scales, a vector of range expansion constants used to add some padding around the data to ensure that they are placed some distance away from the axes. Use the convenience function <code>expansion()</code> to generate the values for the <code>expand</code> argument. The defaults are to expand the scale by 5% on each side for continuous variables, and by 0.6 units on each side for discrete variables.</p> <p><code>position</code> For position scales, The position of the axis. <code>left</code> or <code>right</code> for y axes, <code>top</code> or <code>bottom</code> for x axes.</p> <p><code>call</code> The call used to construct the scale for reporting messages.</p> <p><code>super</code> The super class to use for the constructed scale</p>
<code>n.breaks</code>	An integer guiding the number of major breaks. The algorithm may choose a slightly different number to ensure nice break labels. Will only have an effect if <code>breaks = waiver()</code> . Use <code>NULL</code> to use the default number of breaks given by the transformation.
<code>nice.breaks</code>	Logical. Should breaks be attempted placed at nice values instead of exactly evenly spaced between the limits. If <code>TRUE</code> (default) the scale will ask the transformation object to create breaks, and this may result in a different number of breaks than requested. Ignored if breaks are given explicitly.
<code>values</code>	a set of aesthetic values to map data values to. The values will be matched in order (usually alphabetical) with the limits of the scale, or with breaks if provided. If this is a named vector, then the values will be matched based on the names instead. Data values that don't match will be given <code>na.value</code> .
<code>na.value</code>	The aesthetic value to use for missing (NA) values

**Value**

A ggproto object inheriting from `Scale`

**See Also**

Other `scale_edge_*`: [scale\\_edge\\_alpha\(\)](#), [scale\\_edge\\_colour](#), [scale\\_edge\\_fill](#), [scale\\_edge\\_linetype\(\)](#), [scale\\_edge\\_shape\(\)](#), [scale\\_edge\\_size\(\)](#), [scale\\_edge\\_width\(\)](#)

---

theme\_graph

*A theme tuned for graph visualizations*

---

**Description**

When plotting graphs, networks, and trees the coordinate values are often of no importance and axes are thus a distraction. `ggraph` comes with a build-in theme that removes redundant elements in order to put focus on the data. Furthermore the default behaviour is to use a narrow font so text takes up less space. Theme colour is defined by a background and foreground colour where the background defines the colour of the whole graphics area and the foreground defines the colour of the strip and border. By default strip and border is turned off as it is an unnecessary element unless facetting is used. To add a foreground colour to a plot that is already using `theme_graph` the `th_foreground`

helper is provided. In order to use this appearance as default use the `set_graph_style` function. An added benefit of this is that it also changes the default text-related values in the different geoms for a completely coherent look. `unset_graph_style` can be used to revert the defaults back to their default settings (that is, they are not necessarily reverted back to what they were prior to calling `set_graph_style`). The `th_no_axes()` helper is provided to modify an existing theme so that grid and axes are removed.

## Usage

```
theme_graph(
  base_family = "Arial Narrow",
  base_size = 11,
  background = "white",
  foreground = NULL,
  border = TRUE,
  text_colour = "black",
  bg_text_colour = text_colour,
  fg_text_colour = text_colour,
  title_family = base_family,
  title_size = 18,
  title_face = "bold",
  title_margin = 10,
  title_colour = bg_text_colour,
  subtitle_family = base_family,
  subtitle_size = 12,
  subtitle_face = "plain",
  subtitle_margin = 15,
  subtitle_colour = bg_text_colour,
  strip_text_family = base_family,
  strip_text_size = 10,
  strip_text_face = "bold",
  strip_text_colour = fg_text_colour,
  caption_family = base_family,
  caption_size = 9,
  caption_face = "italic",
  caption_margin = 10,
  caption_colour = bg_text_colour,
  plot_margin = margin(30, 30, 30, 30)
)

th_foreground(foreground = "grey80", fg_text_colour = NULL, border = FALSE)

th_no_axes()

set_graph_style(
  family = "Arial Narrow",
  face = "plain",
  size = 11,
  text_size = 11,
```

```

    text_colour = "black",
    ...
  )

  unset_graph_style()

```

### Arguments

`base_size`, `size`, `text_size`, `title_size`, `subtitle_size`, `strip_text_size`, `caption_size`  
 The size to use for the various text elements. `text_size` will be used as geom defaults

`background`      The colour to use for the background. This theme sets all background elements except for `plot.background` to `element_blank` so this controls the background for all elements of the plot. Set to `NA` to remove the background (thus making the plot transparent)

`foreground`      The colour of foreground elements, specifically `strip` and `border`. Set to `NA` to remove.

`border`            Logical. Should border be drawn if a foreground colour is provided?

`text_colour`, `bg_text_colour`, `fg_text_colour`, `title_colour`, `subtitle_colour`, `strip_text_colour`, `caption_colour`  
 The colour of the text in the various text elements

`title_margin`, `subtitle_margin`, `caption_margin`  
 The margin to use between the text elements and the plot area

`plot_margin`      The plot margin

`family`, `base_family`, `title_family`, `subtitle_family`, `strip_text_family`, `caption_family`  
 The font to use for the different elements

`face`, `title_face`, `subtitle_face`, `strip_text_face`, `caption_face`  
 The fontface to use for the various text elements

...                Parameters passed on the `theme_graph`

### Examples

```

library(tidygraph)
graph <- as_tbl_graph(highschool)

ggraph(graph) + geom_edge_link() + geom_node_point() + theme_graph()

```

---

whigs

*Membership network of American Whigs*


---

### Description

This dataset shows the membership of 136 colonial Americans in 5 whig organization and is a bipartite graph. The data appeared in the appendix to David Hackett Fischer's *Paul Revere's Ride* (Oxford University Press, 1995) and compiled by Kieran Healy for the blog post [Using Metadata to Find Paul Revere](#).

**Usage**

whigs

**Format**

The data is stored as an incidence matrix with persons as rows and organizations as columns. A 0 means no membership while a one means membership.

**Source**

<https://github.com/kjhealy/revere/blob/master/data/PaulRevereAppD.csv> adapted from:  
Fischer, David H. (1995) *Paul Revere's Ride*. Oxford University Press



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