

# Package ‘gglyph’

May 8, 2026

**Title** Network-Style Visualization of Directed Pairwise Relationships

**Version** 0.2.0

**Description** Create network-style visualizations of pairwise relationships using custom edge glyphs built on top of 'ggplot2'. The package supports both statistical and non-statistical data and allows users to represent directed relationships. This enables clear, publication-ready graphics for exploring and communicating relational structures in a wide range of domains. The method was first used in Abu-Akel et al. (2021) <doi:10.1371/journal.pone.0245100>. Code is released under the MIT License; included datasets are licensed under the Creative Commons Attribution 4.0 International (CC BY 4.0).

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**URL** <https://valentinsvelev.github.io/gglyph/>

**BugReports** <https://github.com/valentinsvelev/gglyph/issues/>

**Depends** R (>= 4.1.0)

**Imports** dplyr, ggplot2, ggtext, grid, magrittr, rlang, stats, tibble, tidy, utils

**Suggests** ggthemes, haven, kableExtra, knitr, patchwork, psych, purrr, readr, rmarkdown, rstatix, spelling, svglite, testthat (>= 3.0.0), tidyverse, viridis, viridisLite

**VignetteBuilder** knitr

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**Repository** CRAN

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generate_mock_data	<i>Generate mock data for gglyph::geom_glyph()</i>
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### Description

Generates custom mock data to be passed to gglyph::geom\_glyph().

### Usage

```
generate_mock_data(
  n_nodes = 5,
  n_edges = 7,
  n_groups = 1,
  statistical = FALSE,
  p_threshold = 0.05
)
```

### Arguments

n_nodes	Number of nodes in the graph. Default is 5.
n_edges	Number of edges to generate. Default is 7.
n_groups	Number of groups (for faceting). Default is 1 (ungrouped).
statistical	If TRUE, generates mock p-values for edges. Default is FALSE.
p_threshold	The significance threshold for filtering edges. Default is 0.05.

### Value

A data frame with mock data for nodes and edges.

### Examples

```
# For non-grouped data
mock_data <- generate_mock_data(
  n_nodes = 5,
  n_edges = 7,
  n_groups = 1,
  statistical = FALSE,
```

```
    p_threshold = 0.05
  )

# For grouped data
mock_data <- generate_mock_data(
  n_nodes = 5,
  n_edges = 7,
  n_groups = 3,
  statistical = TRUE,
  p_threshold = 0.05
)
```

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**geom\_glyph***Create a directed network-style graph*

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## Description

Create a network-style graph that illustrates directed pairwise relationships using custom edges.

## Usage

```
geom_glyph(
  mapping = NULL,
  data = NULL,
  edge_size = 1,
  edge_colour = "grey",
  edge_fill = NULL,
  edge_alpha = 1,
  node_size = 1,
  node_colour = "black",
  node_fill = NULL,
  node_alpha = 1,
  node_shape = 21,
  node_spacing = 1,
  label_size = 12,
  group_label_size = 13,
  legend_title = NULL,
  legend_subtitle = NULL,
  ...,
  stat = "identity",
  position = "identity",
  na.rm = FALSE,
  show.legend = TRUE,
  inherit.aes = TRUE
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>aes()</code> . You must supply mapping if there is no plot mapping.
data	A <code>DataFrame</code> with preprocessed data from either <code>gglyph::preprocess_data_general()</code> or <code>gglyph::preprocess_data_statistical()</code> . To be passed to <code>ggplot2::ggplot()</code> .
edge_size	A numeric scaling factor indicating the size/width of the edges. Default is 1.
edge_colour	Color(s) of the edge outlines. Can be a single string (for non-grouped data) or a vector of strings or a function (for grouped data). Default is "grey".
edge_fill	Color(s) for the edge fill. Can be a single string, a vector of strings, or a color function. If <code>NULL</code> , defaults to <code>edge_colour</code> .
edge_alpha	A numeric value indicating the transparency of the edges. Default is 1.
node_size	A numeric value indicating the size of the nodes. Default is 8.
node_colour	Color(s) of the node outlines. Can be a single string (for non-grouped data) or a vector of strings or a function (for grouped data). Default is "black".
node_fill	Color for the node fill. If <code>NULL</code> , defaults to <code>node_colour</code> .
node_alpha	A numeric value indicating the transparency of the nodes. Default is 1.
node_shape	A numeric value specifying the shape of the nodes, following <code>ggplot2</code> 's shape specifications. Default is 21 (a circle with a border).
node_spacing	A numeric scaling factor for the distance between nodes. Values $> 1$ will push nodes further apart, while values $< 1$ will bring them closer. Default is 1.
label_size	A numeric value indicating the size of the node labels. Default is 12.
group_label_size	A numeric value indicating the size of group label. Default is 13.
legend_title	Title for the legend as a string.
legend_subtitle	Subtitle for the legend as a string.
...	Additional arguments passed to <code>ggplot2</code> layer.
stat	The statistical transformation to use on the data for this layer.
position	A position adjustment to use on the data for this layer.
na.rm	If <code>FALSE</code> , the default, missing values are removed with a warning. If <code>TRUE</code> , missing values are silently removed.
show.legend	Should this layer be included in the legends? Default is <code>TRUE</code> .
inherit.aes	If <code>FALSE</code> , overrides the default aesthetics, rather than combining with them. Default is <code>FALSE</code> .

**Value**

A `ggplot2` layer with custom network-based graph.

**See Also**

[ggplot2::ggsave\(\)](#)

## Examples

```
# For non-grouped/-facetted plot
data <- gglyph::generate_mock_data(n_groups = 1)

ggplot2::ggplot(data = data) +
  gglyph::geom_glyph()

ggplot2::ggplot(data = data) +
  gglyph::geom_glyph(edge_colour = "purple", node_colour = "blue")

ggplot2::ggplot(data = data) +
  gglyph::geom_glyph(edge_colour = "purple", node_colour = "blue") +
  ggplot2::labs(title = "A beautiful glyph")

# For grouped/facetted plot
data <- gglyph::generate_mock_data(n_groups = 3)

ggplot2::ggplot(data = data) +
  gglyph::geom_glyph() +
  ggplot2::facet_wrap(~ group)

ggplot2::ggplot(data = data) +
  gglyph::geom_glyph(edge_colour = viridis::viridis, node_colour = viridis::viridis) +
  ggplot2::facet_wrap(~ group)

ggplot2::ggplot(data = data) +
  gglyph::geom_glyph(edge_colour = viridis::viridis, node_colour = viridis::viridis) +
  ggplot2::facet_wrap(~ group) +
  ggplot2::labs(title = "Beautiful glyphs")
```

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pisa\_2022

*PISA 2022 data*

---

## Description

Results of pairwise t-tests (with Bonferroni correction) performed on a subset from the PISA 2022 data.

## Usage

```
data(pisa_2022)
```

## Format

A data frame with 492 rows and 3 variables:

from Category A of educational level (ISCED) attained by the parents of the respondent (character).

to Category B of educational level (ISCED) attained by the parents of the respondent (character).

group Country of the respondent (character).  
sig p-value of the pairwise t-test (numeric).

### Source

Data obtained from:

- OECD (2023). *PISA 2022 Database* [Data Set]. Zenodo. doi:10.5281/zenodo.13382904

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### References

Additional reference(s) for further reading:

- OECD (2024). *PISA 2022 Technical Report*. OECD Publishing, Paris. doi:10.1787/01820d6d-en

### Examples

```
data(pisa_2022)
head(pisa_2022)
```

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process\_data\_general    *Process general/non-statistical data*

---

### Description

Prepare general/non-statistical data for plotting with `gglyph::geom_glyph()`.

### Usage

```
process_data_general(data, from = "from", to = "to", group = NULL)
```

### Arguments

data	A DataFrame or tibble containing the input data to be processed.
from	A string indicating the column name for the start nodes.
to	A string indicating the column name for the end nodes.
group	A string indicating the column name for the grouping variable.

### Value

A DataFrame with the preprocessed data that is to be passed to `gglyph::geom_glyph()`.

## Examples

```
data(sipri_milex_1995_2023)

# For non-grouped data
processed_data <- process_data_general(
  data = sipri_milex_1995_2023,
  from = "from",
  to = "to"
)

# For grouped data
processed_data <- process_data_general(
  data = sipri_milex_1995_2023,
  from = "from",
  to = "to",
  group = "group"
)
```

---

process\_data\_statistical

*Process statistical data*

---

## Description

Prepare statistical data for plotting with `gglyph::geom_glyph()`.

## Usage

```
process_data_statistical(
  data,
  from = "from",
  to = "to",
  group = NULL,
  sig = "sig",
  thresh = 0.05
)
```

## Arguments

<code>data</code>	A <code>DataFrame</code> or <code>tibble</code> containing the input data to be processed.
<code>from</code>	A string indicating the column name for the start nodes.
<code>to</code>	A string indicating the column name for the end nodes.
<code>group</code>	A string indicating the column name for the grouping variable.
<code>sig</code>	A string indicating the column name for the significance level.
<code>thresh</code>	A single number indicating the significance threshold. Default is 0.05.

**Value**

A DataFrame with the preprocessed data that is to be passed to `gglyph::geom_glyph()`.

**Examples**

```
data(pisa_2022)

# For non-grouped data
processed_data <- process_data_statistical(
  data = pisa_2022,
  from = "from",
  to = "to",
  sig = "sig",
  thresh = 0.05
)

# For grouped data
processed_data <- process_data_statistical(
  data = pisa_2022,
  from = "from",
  to = "to",
  sig = "sig",
  group = "group",
  thresh = 0.05
)
```

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sipri\_milex\_1995\_2023 *SIPRI Military Expenditure 1995-2023 data*

---

**Description**

A subset of the SIPRI Military Expenditure 1949-2023 data.

**Usage**

```
data(sipri_milex_1995_2023)
```

**Format**

A data frame with 77 rows and 3 variables:

from Name of country A (character).

to Name of country B (character).

group Year (numeric).

### **Source**

Data obtained from:

- SIPRI (2025). *SIPRI Military Expenditure Database* [Data Set]. doi:10.55163/CQGC9685

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### **Examples**

```
data(sipri_milex_1995_2023)
head(sipri_milex_1995_2023)
```

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