

# Package ‘ontologyPlot’

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**Type** Package

**Title** Functions for Visualising Sets of Ontological Terms

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**Description** Functions for visualising sets of ontological terms using the 'graphviz' layout system.

**License** GPL (>= 2)

**Depends** R (>= 3.0.0)

**Imports** methods, ontologyIndex, paintmap, Rgraphviz

**Suggests** knitr

**VignetteBuilder** knitr

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annotation_grid	<i>Get logical matrix of term annotation for group of cases</i>
-----------------	---

---

### Description

Get logical matrix of term annotation for group of cases

### Usage

```
annotation_grid(ontology, term_sets, all_terms = grid_terms(ontology,
  term_sets), remove_unanimous = FALSE, cluster_rows = TRUE,
  cluster_cols = TRUE)
```

### Arguments

ontology	ontology_index object
term_sets	List of character vectors of ontological term IDs
all_terms	Character vector giving terms to use in annotation.
remove_unanimous	Logical value determining whether to remove terms present in all term_sets.
cluster_rows	Logical value rows determining whether to use hclust to cluster term_sets.
cluster_cols	Logical value rows determining whether to use hclust to cluster terms (based on correlation of inclusion in term_sets).

**Value**

Logical matrix.

---

calibrate\_sizes      *Function to scale values between two given limits*

---

**Description**

Could be useful to modify a vector of sizes to between, say 1 and 3, before passing to ‘onto\_plot’.

**Usage**

```
calibrate_sizes(x, high, low)
```

**Arguments**

x	Numeric vector
high	Numeric value of largest size
low	Numeric value of smallest size

**Value**

Numeric vector

**Examples**

```
calibrate_sizes(c("HP:0000001"]=10, "HP:0000006"]=5), high=3, low=1)
```

---

colour\_by\_frequency      *Function to assign colours to terms based on frequency with which terms appear in term\_sets*

---

**Description**

Function to assign colours to terms based on frequency with which terms appear in term\_sets

**Usage**

```
colour_by_frequency(ontology, terms, term_sets,  
  colour_func = colorRampPalette(c("Yellow", "Green", "#0099FF")))
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms
term_sets	List of character vectors of ontological term IDs
colour_func	Function capable of returning a set of colours, given the number of colours it needs to return

**Value**

Character vector of colours, named by term

**See Also**

[colour\\_by\\_term\\_set](#), [colour\\_by\\_population\\_frequency](#)

---

colour\_by\_population\_frequency

*Function to assign colours to terms based on population frequency of terms*

---

**Description**

Function to assign colours to terms based on population frequency of terms

**Usage**

```
colour_by_population_frequency(ontology, terms, frequencies,
  colour_palette = colorRampPalette(c("Yellow", "Green", "#0099FF"))(10),
  max_colour_freq = max(terms_freq), min_colour_freq = min(terms_freq))
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms
frequencies	Numeric vector of term frequencies named by term IDs
colour_palette	Character vector of colours for the different information contents of the terms to be plotted, going from rare to common
max_colour_freq	Numeric value in [0, 1] giving the maximum frequency (to which the dullest color will be assigned)
min_colour_freq	Numeric value in [0, 1] giving the minimum frequency (to which the brightest color will be assigned)

**Value**

Character vector of colours, named by term

**See Also**

[colour\\_by\\_term\\_set](#), [colour\\_by\\_frequency](#)

---

colour_by_term_set	<i>Function to set colours of nodes in plot to distinguish terms belonging to different term sets</i>
--------------------	---

---

**Description**

Function to set colours of nodes in plot to distinguish terms belonging to different term sets

**Usage**

```
colour_by_term_set(ontology, terms, term_sets, colour_generator = rainbow,  
alpha = 0.5)
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms
term_sets	List of character vectors of ontological term IDs
colour_generator	Function which returns a vector of colours, e.g. rainbow or heat.colors.
alpha	alpha parameter to pass to colour_generator.

**Value**

Character vector of colours, named by term.

**See Also**

[colour\\_by\\_frequency](#), [colour\\_by\\_population\\_frequency](#)

---

dot_string	ontology_plot <i>object to dot string</i>
------------	---

---

**Description**

ontology\_plot object to dot string

**Usage**

```
dot_string(ontology_plot)
```

**Arguments**

ontology\_plot    Object of class 'ontology\_plot' to export.

**Value**

String

**See Also**

[onto\\_plot](#)

---

get_adjacency_matrix	<i>Get an adjacency matrix for a set of ontological terms</i>
----------------------	---

---

**Description**

Get an adjacency matrix for a set of ontological terms

**Usage**

```
get_adjacency_matrix(ontology, terms)
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms

**Value**

A logical matrix representing the adjacency matrix of terms based on the directed acyclic graph of ontology. A TRUE entry means the term corresponding to the column is a parent of the row term in ontology.

### See Also

[get\\_pseudo\\_adjacency\\_matrix](#)

### Examples

```
library(ontologyIndex)
data(hpo)
get_adjacency_matrix(hpo, c("HP:0000118", "HP:0001873", "HP:0011877"))
```

---

`get_node_friendly_long_names`

*Split up node labels across lines so they fit in nodes better*

---

### Description

Split up node labels across lines so they fit in nodes better

### Usage

```
get_node_friendly_long_names(ontology, terms, official_names = FALSE)
```

### Arguments

<code>ontology</code>	ontology_index object
<code>terms</code>	Character vector of ontological terms
<code>official_names</code>	Logical value indicating whether to use the exact names from the ontology. Otherwise, shortened, capitalised names are used.

### Value

Character vector.

### Examples

```
library(ontologyIndex)
data(hpo)
get_node_friendly_long_names(hpo, c("HP:0001873", "HP:0011877"))
```

---

get\_ontology\_plot      *Get ontology\_plot object*

---

### Description

Function to create ontology\_plot objects where all graphical parameters to be used must be specified.

### Usage

```
get_ontology_plot(ontology, terms, edge_attributes = list(color = "#000000",
  lty = "solid"), ...)
```

### Arguments

ontology	ontology_index object
terms	Character vector of ontological terms
edge_attributes	List of properties to set for arrows (note, these properties will be used for all arrow).
...	Named graphical parameters. These must either be vectors of values the same length as terms, or of length 1 if they should be used for all terms.

### Value

ontology\_plot object.

---

get\_pseudo\_adjacency\_matrix  
*Get an adjacency matrix for a set of ontological terms*

---

### Description

Get an adjacency matrix for a set of ontological terms

### Usage

```
get_pseudo_adjacency_matrix(ontology, terms)
```

### Arguments

ontology	ontology_index object
terms	Character vector of ontological terms



**Value**

A logical matrix representing the adjacency matrix of terms based on the directed acyclic graph of ontology. A TRUE entry means the term corresponding to the column is a parent of the row term within terms.

**See Also**

[get\\_adjacency\\_matrix](#)

**Examples**

```
library(ontologyIndex)
data(hpo)
get_pseudo_adjacency_matrix(hpo, c("HP:0000118", "HP:0001873", "HP:0011877"))
```

---

get_shortened_names	<i>Get human readable, shortened (where possible) ontological term names</i>
---------------------	--

---

**Description**

Get human readable, shortened (where possible) ontological term names

**Usage**

```
get_shortened_names(ontology, terms)
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms

**Value**

Character vector

**Examples**

```
library(ontologyIndex)
data(hpo)
get_shortened_names(hpo, c("HP:0001873", "HP:0011877"))
```

---

grid_terms	<i>Get set of HPO terms appropriate for showing in a grid</i>
------------	---

---

**Description**

Get set of HPO terms appropriate for showing in a grid

**Usage**

```
grid_terms(ontology, term_sets)
```

**Arguments**

ontology	ontology_index object
term_sets	List of character vectors of ontological term IDs

**Value**

Character vector of terme IDs.

---

label_by_frequency	<i>Function to get plot labels for terms based on frequency in term_sets</i>
--------------------	--

---

**Description**

Function to get plot labels for terms based on frequency in term\_sets

**Usage**

```
label_by_frequency(ontology, terms, term_sets)
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms
term_sets	List of character vectors of ontological term IDs

**Value**

Character vector of labels, named by term.

**See Also**

[simple\\_labels](#), [long\\_labels](#)

---

label_by_term_set	<i>Function to label nodes by term_set</i>
-------------------	--

---

**Description**

Function to label nodes by term\_set

**Usage**

```
label_by_term_set(ontology, terms, term_sets)
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms
term_sets	List of character vectors of ontological term IDs

**Value**

Character vector of colours, named by term.

**See Also**

[simple\\_labels](#), [label\\_by\\_frequency](#), [long\\_labels](#)

---

long_labels	<i>Function to assign detailed node labels to terms</i>
-------------	---

---

**Description**

Label includes term ID, term name, number of instances of term amongst term\_sets and percentage frequency in population.

**Usage**

```
long_labels(ontology, terms, term_sets, frequencies)
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms
term_sets	List of character vectors of ontological term IDs
frequencies	Numeric vector of term frequencies named by term IDs

**Value**

Character vector of labels, named by term.

**See Also**

[simple\\_labels](#), [label\\_by\\_frequency](#), [label\\_by\\_term\\_set](#)

---

n\_most\_frequent\_terms *Select n most prevalent terms in term\_sets*

---

**Description**

Selects n most prevalent terms in set of term sets/annotations including implicit terms. If more than one term are tied at the nth position, all terms are included in the result.

**Usage**

```
n_most_frequent_terms(ontology, term_sets, n,  
  terms = unique(unlist(term_sets)))
```

**Arguments**

ontology	ontology_index object
term_sets	List of character vectors of ontological term IDs
n	Integer
terms	Character vector of ontological terms

**Value**

Character vector of length at most n

**See Also**

[remove\\_terms\\_with\\_less\\_than\\_n\\_occurrences](#)

**Examples**

```
library(ontologyIndex)  
data(hpo)  
n_most_frequent_terms(hpo, c("HP:0001873"),  
  list(term_sets=list("HP:0001873", "HP:0001902")), n=2)
```

---

official_labels	<i>Get official names for terms</i>
-----------------	-------------------------------------

---

**Description**

Get official names for terms

**Usage**

```
official_labels(ontology, terms)
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms

**Value**

Character vector of labels, named by term.

**See Also**

[simple\\_labels](#)

---

ontologyPlot	<i>Functions for Visualising Sets of Ontological Terms</i>
--------------	--

---

**Description**

Functions for visualising sets of ontological terms using the ‘graphviz’ layout system.

**Details**

Package:	ontologyPlot
Type:	Package
Version:	1.0
Date:	2016-01-11
License:	GPL (>= 2)

This package succeeds the package hpoPlot with an improved interface and focusing on general ontologies. The key function is [onto\\_plot](#), which creates an object of class ontology\_plot which can be displayed as a graph or exported to dot format.

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**References**

‘The Human Phenotype Ontology project: linking molecular biology and disease through phenotype data’, Nucl. Acids Res. (1 January 2014) 42 (D1): D966-D974 doi:10.1093/nar/gkt1026 Westbury, S. K. et al. (2015). Human Phenotype Ontology annotation and cluster analysis to unravel genetic defects in 707 cases with unexplained bleeding and platelet disorders. Genome Medicine. 7 (2015)

---

onto\_plot

*Get ontology\_plot object*

---

**Description**

A convenience wrapper for the [get\\_ontology\\_plot](#) function, enabling functions to be passed to generate graphical parameters for terms automatically.

**Usage**

```
onto_plot(ontology, term_sets = NULL, frequencies = NULL,
  terms = remove_uninformative_terms(ontology, term_sets),
  edge_attributes = list(color = "#000000", lty = "solid"),
  fillcolor = "powderblue", label = simple_labels, color = "transparent",
  width = 0.75, fontsize = 30, style = "filled", fixedsize = "true",
  shape = "circle", ...)
```

**Arguments**

ontology	ontology_index object
term_sets	List of character vectors of ontological term IDs
frequencies	Numeric vector of term frequencies named by term IDs
terms	Character vector of ontological terms
edge_attributes	List of properties to set for arrows (note, these properties will be used for all arrow).
fillcolor	Character vector of colours to fill nodes corresponding to terms with. Alternatively a function to set the colours of the nodes in the graph based on term_sets.
label	Character vector of labels (or function to set them).
color	Character vector of colours for borders of nodes representing terms (or function to set them).
width	Numeric vector of widths for nodes (of function to set them).

fontsize	Numeric vector of font sizes for the text to be placed in the nodes (or function to set them).
style	Display style for nodes, defaults to "filled".
fixedsize	Character indicating whether nodes should be fixed size, "true", or adjusted to fit around the contained text, "false".
shape	Character vector of shape names for nodes (or function to set them). Defaults to "circle".
...	Other node attributes for dot format.

**Value**

ontology\_plot object.

**See Also**

[get\\_ontology\\_plot](#), [write\\_dot](#)

**Examples**

```
library(ontologyIndex)
data(hpo)
hpo_phenotypes <- c(
  A=c("HP:0001382", "HP:0004272", "HP:0007917", "HP:0004912", "HP:0001596"),
  B=c("HP:0001382", "HP:0004272", "HP:0002165", "HP:0004800", "HP:0004912"),
  C=c("HP:0004800", "HP:0001382", "HP:0004912", "HP:0007917", "HP:0008743"),
  D=c("HP:0001257", "HP:0001382", "HP:0007917", "HP:0012623", "HP:0002165"),
  E=c("HP:0007917", "HP:0004800", "HP:0004272", "HP:0001596", "HP:0002165")
)

onto_plot(
  ontology=hpo,
  term_sets=hpo_phenotypes
)
```

---

plot.ontology\_plot      *Plotting function for ontology\_plot object*

---

**Description**

Plotting function for ontology\_plot object

**Usage**

```
## S3 method for class 'ontology_plot'
plot(x, ...)
```

**Arguments**

x                    Object of class ontologicalPlot.  
 ...                  Other options passed to plot().

**Value**

Nothing, side-effect: plots a graph.

---

plot\_annotation\_grid    *Plot a logical matrix of term annotation*

---

**Description**

Plot a logical matrix of term annotation

**Usage**

```
plot_annotation_grid(..., on_colour = "#FF0000FF", off_colour = "#FFFFBFFF")
```

**Arguments**

...                    Arguments to be passed to [annotation\\_grid](#).  
 on\_colour            Colour to use to show presence of term.  
 off\_colour           Colour to use to show absence of term.

**Value**

Plots heatmap.

---

print.ontology\_plot    *Print function for ontology\_plot object*

---

**Description**

Print function for ontology\_plot object

**Usage**

```
## S3 method for class 'ontology_plot'  
print(x, ...)
```

**Arguments**

x                    Object of class ontologicalPlot.  
 ...                  Other options passed to be passed to plot().



**Value**

Nothing. Side-effect: plots graphs.

---

*p\_values\_for\_occurrence\_of\_term\_in\_group*

*Get p-values for observing at least as many of each term as occur in term\_sets given the population frequencies of the terms*

---

**Description**

Get p-values for observing at least as many of each term as occur in term\_sets given the population frequencies of the terms

**Usage**

`p_values_for_occurrence_of_term_in_group(ontology, term_sets, terms_freq)`

**Arguments**

- ontology            ontology\_index object
- term\_sets           List of character vectors of ontological term IDs
- terms\_freq         Numeric vector of population frequencies of terms.

**Value**

Numeric vector of log p-values named by corresponding term.

**See Also**

[width\\_by\\_significance](#)

---

*remove\_links*

*Remove terms which just link two other terms together in a subontology*

---

**Description**

Remove terms which just link two other terms together in a subontology

**Usage**

`remove_links(ontology, terms, hard = FALSE)`

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms
hard	Logical value determining whether to multiple edges to leaf terms are kept - <code>`hard=FALSE`</code> , or removed - <code>`hard=TRUE`</code> .

**Value**

Character vector.

**See Also**

[remove\\_uninformative\\_terms](#)

**Examples**

```
library(ontologyIndex)
data(hpo)
remove_links(hpo, c("HP:0001873", "HP:0001872", "HP:0011873", "HP:0011877"))
```

---

remove\_terms\_with\_less\_than\_n\_occurrences

*Remove terms with less than certain number of occurrences*

---

**Description**

Remove terms with less than certain number of occurrences

**Usage**

```
remove_terms_with_less_than_n_occurrences(ontology, term_sets, n,
  terms = unique(unlist(term_sets)))
```

**Arguments**

ontology	ontology_index object
term_sets	List of character vectors of ontological term IDs
n	Integer
terms	Character vector of ontological terms

**Value**

Character vector

**See Also**

[n\\_most\\_frequent\\_terms](#)

## Examples

```
library(ontologyIndex)
data(hpo)
remove_terms_with_less_than_n_occurrences(hpo,
term_sets=list("HP:0001873", "HP:0001902"), n=2)
```

---

remove\_uninformative\_terms

*Remove uninformative terms from union of all terms in set of annotations*

---

## Description

For a set of ontological annotation sets, remove terms annotated to the same objects as all their children. Useful for selecting terms for summarising a set of annotation sets, as it can lead to a significant reduction in the number of terms.

## Usage

```
remove_uninformative_terms(ontology, term_sets)
```

## Arguments

ontology	ontology_index object
term_sets	List of character vectors of ontological term IDs

## Value

Character vector of terms

## Examples

```
library(ontologyIndex)
data(hpo)
remove_uninformative_terms(hpo, list(Patient1=c("HP:0001873", "HP:0000118")))
```

simple\_cap                      *Capitalise words in character vector*

---

**Description**

Capitalise words in character vector

**Usage**

```
simple_cap(x)
```

**Arguments**

x                      Character vector

**Value**

Character vector

**Examples**

```
simple_cap(c("a simple test", "Another-test"))
```

---

simple\_labels                      *Get simplified labels for terms*

---

**Description**

Get simplified labels for terms

**Usage**

```
simple_labels(ontology, terms)
```

**Arguments**

ontology                      ontology\_index object  
terms                      Character vector of ontological terms

**Value**

Character vector of labels, named by term.

**See Also**

[official\\_labels](#)

---

to_svg_string	<i>Convert ontology_plot to SVG string</i>
---------------	--

---

**Description**

Note that by setting "id" and "class" attributes it enables nodes to be selected for manipulation using Javascript if interactivity is desired.

**Usage**

```
to_svg_string(op)
```

**Arguments**

op	Object of class ontology_plot.
----	--------------------------------

**Value**

Character vector of length 1 containing SVG representation of node.

**See Also**

[onto\\_plot](#), [get\\_ontology\\_plot](#)

---

width_by_frequency	<i>Function to get node sizes for terms based on frequency in term_sets</i>
--------------------	---

---

**Description**

Function to get node sizes for terms based on frequency in term\_sets

**Usage**

```
width_by_frequency(ontology, terms, term_sets)
```

**Arguments**

ontology	ontology_index object
terms	Character vector of ontological terms
term_sets	List of character vectors of ontological term IDs

**Value**

Character vector of sizes, named by term

**See Also**

[width\\_by\\_significance](#)

---

`width_by_significance` *Function to get node sizes for terms based on statistical significance of seeing at least this number of each term in term\_sets*

---

### Description

Function to get node sizes for terms based on statistical significance of seeing at least this number of each term in term\_sets

### Usage

```
width_by_significance(ontology, terms, term_sets, frequencies)
```

### Arguments

<code>ontology</code>	ontology_index object
<code>terms</code>	Character vector of ontological terms
<code>term_sets</code>	List of character vectors of ontological term IDs
<code>frequencies</code>	Numeric vector of term frequencies named by term IDs

### Value

Character vector of sizes, named by term

### See Also

[width\\_by\\_frequency](#)

---

`write_dot` *Export ontology\_plot object as dot file*

---

### Description

Export ontology\_plot object as dot file

### Usage

```
write_dot(ontology_plot, file)
```

### Arguments

<code>ontology_plot</code>	Object of class 'ontology_plot' to export.
<code>file</code>	Character value of target file path.

**Value**

Nothing, side effect - writes to file.

**See Also**

[dot\\_string](#)

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