# Package 'leidenbase'

January 14, 2022

Type Package	
Title R and C wrappers to run the Leiden find_partition function	
Version 0.1.4	
Description An R to C interface that runs the Leiden community detection algorithm to find a basic partition. It runs the equivalent of the find_partition() function, which is given in the Leidenalg distribution file 'leiden/src/functions.py'. This package includes the required source code files from the official Leidenalg distribution and several functions from the R igraph package. The Leidenalg distribution is available from https://github.com/vtraag/leidenalg and the R igraph package is available from https://igraph.org/r/.  The Leiden algorithm is described in the article 'From Louvain to Leiden: guaranteeing well-connected communities', V. A. Traag and L. Waltman and N. J. van Eck, Scientific Reports (2019), DOI: 10.1038/s41598-019-41695-z.	
<b>Requires</b> R (>= 3.0.0)	
Imports igraph ( $>= 0.8.2$ )	
<b>License</b> GPL (>=2) + file LICENSE	
Encoding UTF-8	
LazyData true	
RoxygenNote 7.1.2	
Suggests testthat	
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leiden\_find\_partition Leiden find partition community detection function

## **Description**

R to C wrapper that runs the basic Leiden community detection algorithm, which is similar to the find partition() function in the python Leidenalg distribution.

#### Usage

```
leiden_find_partition(
  igraph,
  partition_type = c("CPMVertexPartition", "ModularityVertexPartition",
    "RBConfigurationVertexPartition", "RBERVertexPartition",
    "SignificanceVertexPartition", "SurpriseVertexPartition"),
  initial_membership = NULL,
  edge_weights = NULL,
  node_sizes = NULL,
  seed = NULL,
  resolution_parameter = 0.1,
 num_iter = 2,
  verbose = FALSE
)
```

# **Arguments**

igraph R igraph graph. partition\_type String partition type name. Default is CPMVertexParition.

initial\_membership

Numeric vector of initial membership assignments of nodes. These are 1-based indices. Default is one community per node.

edge\_weights Numeric vector of edge weights. Default is 1.0 for all edges.

node\_sizes Numeric vector of node sizes. Default is 1 for all nodes.

seed Numeric random number generator seed. The seed value must be either NULL

for random seed values or greater than 0 for a fixed seed value. Default is NULL.

resolution\_parameter

Numeric resolution parameter. The value must be greater than 0.0. Default is 0.1. The resolution\_parameter is ignored for the partition\_types ModularityVertexPartition, Significance VertexPartition, and Surprise VertexPartition.

num\_iter Numeric number of iterations. Default is 2.

verbose A logic flag to determine whether or not we should print run diagnostics. leiden\_find\_partition 3

#### **Details**

The Leiden algorithm is described in From Louvain to Leiden: guaranteeing well-connected communities. V. A. Traag and L. Waltman and N. J. van Eck Scientific Reports, 9(1) (2019) DOI: 10.1038/s41598-019-41695-z.

Significance is described in Significant Scales in Community Structure V. A. Traag, G. Krings, and P. Van Dooren Scientific Reports, 3(1) (2013) DOI: 10.1038/srep02930

Notes excerpted from leidenalg/src/VertexPartition.py

- *CPMVertexPartition* Implements Constant Potts Model. This quality function uses a linear resolution parameter and is well-defined for both positive and negative edge weights.
- *ModularityVertexPartition* Implements modularity. This quality function is well-defined only for positive edge weights.
- RBConfigurationVertexPartition Implements Reichardt and Bornholdt's Potts model with a configuration null model. This quality function uses a linear resolution parameter and is well-defined only for positive edge weights.
- RBERVertexPartition Implements Reichardt and Bornholdt's Potts model with an Erdos-Renyi null model. This quality function uses a linear resolution parameter and is well-defined only for positive edge weights.
- Significance Vertex Partition Implements Significance. This quality function is well-defined only for unweighted graphs.
- SurpriseVertexPartition Implements (asymptotic) Surprise. This quality function is well-defined only for positive edge weights.

#### Value

A named list consisting of a numeric vector of the node memberships (1-based indices), a numeric quality value, a numeric modularity, a numeric significance, a numeric vector of edge weights within each community, a numeric vector of edge weights from each community, a numeric vector of edge weights to each community, and total edge weight.

### References

V. A. Traag, L. Waltman, N. J. van Eck (2019). From Louvain to Leiden: guaranteeing well-connected communities. Scientific Reports, 9(1). DOI: 10.1038/s41598-019-41695-z

Significant Scales in Community Structure V. A. Traag, G. Krings, and P. Van Dooren Scientific Reports, 3(1) (2013) DOI: 10.1038/srep02930

# Examples

```
library(igraph)
fpath <- system.file( 'testdata', 'igraph_n1500_edgelist.txt.gz', package = 'leidenbase' )
zfp <- gzfile(fpath)
igraph <- read_graph( file = zfp, format='edgelist', n=1500 )
res <- leiden_find_partition(igraph=igraph, partition_type='CPMVertexPartition', resolution_parameter=1e-5)</pre>
```

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