Package: scoredec (via r-universe)

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Type Package
Title S-Core Graph Decomposition
Version 0.1.1
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Description S-Core Graph Decomposition algorithm for graphs. This is a method for decomposition of a weighted graph, as proposed by Eidsaa and Almaas (2013) <doi:10.1103 physreve.88.062819="">. The high speed and the low memory usage make it suitable for large graphs.</doi:10.1103>
License GPL-3
LinkingTo Rcpp
Encoding UTF-8
<pre>URL https://github.com/cadam00/scoredec, https://cadam00.github.io/scoredec/</pre>
<pre>BugReports https://github.com/cadam00/scoredec/issues</pre>
Imports Rcpp (>= 1.0.12), Rfast, igraph
Suggests knitr, rmarkdown, testthat (>= 3.0.0)
Config/testthat/edition 3
RoxygenNote 7.3.2
VignetteBuilder knitr, rmarkdown
NeedsCompilation yes
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Repository CRAN
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s-core community decomposition

Description

s-core community decomposition

Usage

```
s_coreness(g = NULL, W = NULL, mode = "all")
```

Arguments

g	igraph object. It is a weighted graph. If it is not weighted, then the igraph::coreness function will be used. It can be used as an alternative to W.
W	matrix object. It is an adjacency matrix. It can be used as an alternative to g.
mode	character object. It can be one of "all", "in" or "out".

Details

s-core community decomposition implementation. Only one of g or W must be provided.

While the source code is not as clear as the one at brainGraph::s_core, it is very speed and memory efficient. In case that the adjacency matrix W is provided instead of the graph g is provided, then this function is very speed and memory efficient.

Note that in cases that the adjacency matrix W is known to be symmetric (checked, for example, with base::isSymmetric orRfast::is.symmetric), then mode = "in" and mode = "out" will produce the same result more efficiently. For efficiency reasons not checking it is chosen, but user should do it.

Value

Integer vector with s-coreness attribute to each vertex.

References

Eidsaa, M. and Almaas, E. (2013) 's-core network decomposition: A generalization of k-core analysis to weighted networks', *Phys. Rev. E.*, American Physical Society, **88**, 062819. doi:10.1103/PhysRevE.88.062819.

Watson, C.G. (2024). brainGraph: Graph Theory Analysis of Brain MRI Data. R package version 3.1.0. doi:10.32614/CRAN.package.brainGraph.

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Examples

```
set.seed(42)
## Create a dummy symmetric adjacency matrix
W <- matrix(runif(n^2),n)</pre>
W[lower.tri(W)] = t(W)[lower.tri(W)]
diag(W) <- 0
print(scoredec::s_coreness(g = NULL, W = W, mode = "all"))
#> [1] 3 1 2 4 4
base::isSymmetric(W)
#> [1] TRUE
all.equal(scoredec::s_coreness(g = NULL, W = W, mode = "all"),
scoredec::s_coreness(g = NULL, W = W, mode = "in"))
#> [1] TRUE
# Create a dummy undirected graph
g <- igraph::graph_from_adjacency_matrix(adjmatrix = W,</pre>
                                         mode = "undirected",
                                         weighted = TRUE)
print(scoredec::s_coreness(g = g, W = NULL, mode = "all"))
#> [1] 3 1 2 4 4
```

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