

# Package ‘SteinerNet’

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**Title** Steiner approach for Biological Pathway Graph Analysis

**License** GPL-3

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**Depends** R (>= 2.0), igraph0 , RBGL , limma

**Imports** igraph0, RBGL, limma

**Description** A set of graph functions written in R to perform steiner tree network data analysis on biological pathway data

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generate\_st\_samples  
*generate\_st\_samples*

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## Description

This function generates simulation data. It creates random graphs with randomly selected terminals.

**Usage**

```
generate_st_samples(test, graph, folder= NULL, listofterminaltest, repetition)
```

**Arguments**

<code>test</code>	<code>test</code> selects the test type to make random data for it. the random walk for exact algorithm makes subgraphs that include random terminals, but for heuristics it selects random terminals on the base graph and returns the terminal set only
<code>graph</code>	<code>graph</code> is the base graph for generating random subgraphs and random terminal set.
<code>folder</code>	<code>folder</code> specifies a folder name to store the simulated data inside it.
<code>listofterminaltest</code>	<code>listofterminaltest</code> is an input list. Elements of the list are number of terminals to select for a simulation.
<code>repetition</code>	<code>repetition</code> is a list of probabilities. Its length declares the number of elements if random data set that is created for terminals. Each element of the list is the probability of selecting a node as terminal while the random walk is traversing the base graph.

**Details**

This function generates random data for two type of simulations. For experiments that include exact algorithms, it generates random subgraphs with randomly selected terminals. Otherwise it returns only a set of random terminals to be used with the base graph.

Test specifies the type of simulation. `test` can be `exact` or `appr`,

`exact` refers to generation of data for an experiment that includes exact Steiner tree algorithm.

`appr` refers to generation of data for a experiment that involves only approximate Steiner tree algorithms.

If `folder` is `NULL`, it will use default value "steinerdata2" for folder when type is `exact` and "steinerdata" when type is `appr`.

`listofterminaltest` in our study was made of 5, 8, 20, 50, 70 for comparing approximate algorithms and it was 5, 8 for experiments that included exact Steiner tree algorithm. [1]

In our study, we repeated the tests 50 times, and we made the random walk to select a node to be terminal with 0.5 probability. Therefore `repetition` in our comparison was a list of fifty 0.5 values.

**Value**

The function stores the random data in address that is stated in `folder`. When `test` is `exact` the output includes random subgraphs and random set of terminals. When `test` is `appr` the function returns random sets of terminals.

**Author(s)**

Afshin Sadeghi, Holger Froehlich

## References

1. Please refer to the paper that is published with this package.

## Examples

```
library(SteinerNet)
g <- graph.ring(10)
#generate_st_samples("exact", g, "testfolder", c(2,3), c(.8,.8))
```

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steinertree

*steinertree*


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## Description

A function which involves a set of steiner tree algorithms on networks. This set involves an exact algorithm and five heuristic algorithms.

## Usage

```
steinertree(type, coloring = FALSE, ter_list = NULL, enumerate = FALSE, graph)
```

## Arguments

<code>type</code>	<code>type</code> specifies which steiner algorithm to use.
<code>coloring</code>	<code>coloring</code> is a boolean value. When it is TRUE, the function will return a list that consists of the result stienner tree and a copy of the input graph while including the random terminals within the input graph are specified by coloring of nodes. In this graph, green nodes would represent steiner nodes and red nodes represent terminals.
<code>ter_list</code>	<code>ter_list</code> is an input list of terminals. Steiner tree algorithm finds the result according to terminals.
<code>enumerate</code>	<code>enumerate</code> a boolean value to force the steiner tree function to return merged enumerated solutions.
<code>graph</code>	<code>graph</code> is an input igraph object which is delivered to one of the steiner tree algorithms of the package.

## Details

This function withholds six steiner tree algorithms for networks. `type` specifes the steiner algorithm to deploy to the input graph and terminal set. `type` can be SP, KB, RSP, EXA or SPM. The explanation of the abbreviations is listed below.

SP refers to the shortest path heuristic algorithm. [1,2]

KB exerts to Kruskal-Based Heuristic algorithm. [3]

RSP exerts a random approximation algorithm developed by the package developers. [4]

EXA in single mode uses an exact algorithm to return one of the optimal solutions of the problem. In enumerate mode, returns the merged enumerated solution. [4,5]

SPM in single mode returns one of heuristic enumeration algorithm solutions for the problem. In enumerate mode, returns the merged enumerated solution.[4]

EXA and SPM algorithms can return a single solution or run in enumerating mode. The boolean value of enumerate specifies one of the two cases. If this value is FALSE they return one of their enumerated steiner solutions without merging it to other solutions. If it is TRUE they return the merged enumerated solutions of the steiner tree problem.

According to our knowledge RSP, EXA Enumeration, SPM are represented for the first time in this package and are new algorithms. [4]

ter\_list value can be NULL. In this case, the function will observe graph vertex colors to find terminals. To call the function in this approach, the terminal nodes should be colored in red and the non-terminal nodes should be yellow.

This function handles input igraph objects which their vertices have labels and names. To apply the function on graphs with no label and name, steinertree function automatically recognizes non-labeled graph vertices and creates names and labels for them. The new labels and names for vertices are created according to the vertex ID of each vertex.

## Value

When coloring is FALSE returns a Steiner tree in form of a new igraph object. When coloring is TRUE returns a list that consists of two objects. The first is a steiner tree and the second object is a colored version of the input graph with distinguished steiner nodes and terminals.

## Author(s)

Afshin Sadeghi, Holger Froehlich

## References

1. Path heuristic and Original path heuristic ,Section 4.1.3 of the book "The Steiner tree Problem", Petter,L,Hammer
2. "An approximate solution for the Steiner problem in graphs" , H Takahashi, A Matsuyama
3. F K. Hwang, D S. Richards and P Winter,"The steiner tree Problem", Kruskal-Based Heuristic Section 4.1.4,ISBN: 978-0-444-89098-6
4. Please refer to the paper that is pulished with this package.
5. F K. Hwang, D S. Richards and P Winter,"The steiner tree Problem", Kruskal-Based Heuristic Section 4.1.4, The Optimal solution for stiner trees on networks,ISBN: 978-0-444-89098-6.

## Examples

```
#example 1
library(SteinerNet)
g <- graph.ring(10)
ter_list= c(1,2,9)
SPM=steinertree("SPM", FALSE, ter_list, TRUE, g)
```

```

#example 2
g2 <- graph( c(0,2,1,2,2,3,3,4,5,6,3,6,2,7,2,5,2,6,5,8), directed=FALSE)
V(g2)$color="yellow"
V(g2)$color[c(1,4)]= "red"
OP=steinertree("SP", TRUE, NULL, FALSE, g2)

#example 3: A case study with a sample graph and a given gene list

g <- graph( c(0,2,1,2,2,3,3,4,5,6,3,6,2,7,2,5,2,6,5,8), directed=FALSE)
V(g)$name=c(1058, 51203, 6515, 83879, 160897, 10531, 8659, 2947, 643008)
geneid_list= c(1058,83879, 160897,643008)
list =as.character(geneid_list)
#we include into the test those geneIDes who exist within the base graph.
r =1:(length(list))
t =sapply (r ,function(r) !is.na(match(list[r],c(g)[[9]][[3]]$name ) ))
IDlist1=list[t==TRUE]
ST1=steinertree( "SP", FALSE,IDlist1, FALSE ,g)
#result1= graphEnt_Sym(ST1[[1]]) # converts the Gene entrez ID taken from the graph node "
#tkplot(result1) # tkplot function displays labels instead of names

```

---

```

steiner_comparison_plots
      steiner_comparison_plots

```

---

## Description

This function plots the comparison results of Steiner tree algorithms excutions on simulated data.

## Usage

```
steiner_comparison_plots (test_name, test_folder =NULL, outputname = NULL, listoft
```

## Arguments

test_name	test_name selects the plot type to creat. 14 type of comparison are available to perform.
test_folder	testfolder specifies a folder name to read the result of steiner tree simulations.
outputname	outputname is name of a pdf file to store the result.
listofterminaltest	listofterminaltest is an input list. Elements of the list are number of terminals that are selected for a simulation.
repetition	repetition is a list of probabilities. Its length declares the number of elements if random data set that is created for each terminal number.

## Details

This function creates 12 different comparison types and depicts them by plots. `test_name` specifies the type of comparison.

`test_name` can be one of the following character values.

`exact` refers to time and edge number comparison of Steiner tree algorithms including the exact algorithm.

`appr` refers to time and edge number comparison of Steiner tree algorithms without the exact algorithm.

`Enum` refers to time and edge number comparison of Steiner tree enumeration algorithms.

`Enum-median-venn-node-edge` refers to edge and node number comparison of subgraphs made by Steiner tree enumeration algorithms via Venn diagram.

`org` refers to edge number comparison of random subgraphs that are made by simulations.

`org-dens-e` refers to edge density comparison of random graphs that are made by random graph generator.

`appr-vfreq` refers to vertex frequency comparison of Steiner tree algorithms without the exact algorithm.

`exact-vfreq` refers to vertex frequency comparison of Steiner tree algorithms including the exact algorithm.

`Enum-vfreq` refers to vertex frequency comparison of Steiner tree enumeration algorithms.

`appr-density-e` refers to edge density comparison of steiner tree algorithms excluding the exact algorithm.

`exact-density-e` refers to edge density comparison of steiner tree algorithms including the exact algorithm.

`Enum-density-e` refers to edge density comparison of steiner tree enumeration algorithms.

If `testfolder` is `NULL`, it will use default value "steinerdatae" for folder when `test` is `exact` and "steinerdataenum" when `test` is `enum`.

When `outputname` is `NULL`, a default value would be used for output pdf file name with consideration of selected type.

`listofterminaltest` in our study was made of 5, 8, 20, 50, 70 for comparing approximate algorithms and it was 5, 8 for experiments that included exact Steiner tree algorithm. [1]

In our study, we repeated the tests 50 times, and we made the random walk to select a node to be terminal with 0.5 probability while it traverses the base graph. Therefore `repetition` in our comparison was a list of fifty 0.5 values. If `repetition` is `NULL`, the function regards the `repetition` and `listofterminaltest` values that were used in our study.

## Value

The function stores a resulted plot in a PDF file.

## Author(s)

Afshin Sadeghi, Holger Froehlich

## References

1. Please refer to the paper that is published with this package.

## Examples

```
library(SteinerNet)
g <- graph.ring(10)
#generate_st_samples("exact", g, "testfolder", c(2,3), c(.8,.8))
#steiner_simulation("exact", "testfolder", c(2,3), c(.8,.8))
#steiner_comparison_plots ("exact", "testfolder", c(2,3), c(.8,.8))
```

---

steiner\_simulation *steiner\_simulation*

---

## Description

This function executes Steiner algorithms on simulated data and stores their results into files.

## Usage

```
steiner_simulation(test, listofterminaltest, repetition, testfolder = NULL)
```

## Arguments

test	test selects the test type to apply the simulation. It can be exact, appr, or enum.
listofterminaltest	listofterminaltest is an input list. Elements of the list are number of terminals that are selected for a simulation.
repetition	repetition is a list of probabilities. Its length declares the number of elements if random data set that is created for each terminal number.
testfolder	testfolder specifies a folder name to read the simulated data from it and to store Steiner tree algorithms results inside it.

## Details

This function performs three type of experiments. Test specifies the type of comparison to perform.

test can be exact or appr or enum,

exact refers to executing the set of steiner tree algorithms including the exact algorithm.

appr forces to executing the set of steiner tree algorithms without the exact algorithm.

enum refers to to executing the set of steiner tree enumeration algorithms.

listofterminaltest in our study was made of 5, 8, 20, 50, 70 for comparing approximate algorithms and it was 5, 8 for experiments that included exact Steiner tree algorithm. [1]

If testfolder is NULL, it will use default value "steinerdata2" for folder when test is exact and "steinerdataenum" when test is enum.

In our study, we repeated the tests 50 times, and we made the random walk to select a node to be terminal with 0.5 probability. Therefore repetition in our comparison was a list of fifty 0.5 values.

**Value**

The function stores the result of execution of Steiner trees and the time of their executions in an address that is stated in `testfolder`.

**Author(s)**

Afshin Sadeghi, Holger Froehlich

**References**

1. Please refer to the paper that is published with this package.

**Examples**

```
library(SteinerNet)
g <- graph.ring(10)
#generate_st_samples("exact", g, "testfolder", c(2,3), c(.8,.8))
#steiner_simulation("exact", "testfolder", c(2,3), c(.8,.8))
```



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\*Topic **graphs, protein interaction,  
network, pathway data  
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