# Package 'MeSHSim'

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Type Package

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Title MeSH(Medical Subject Headings) Semantic Similarity Measures

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Imports XML, RCurl

**Depends** R(>= 3.0.0)

**Description** Provide for measuring semantic similarity over MeSH headings and MEDLINE documents

License GPL-2

biocViews Clustering, Software

NeedsCompilation no

# **R** topics documented:

docInfo	
docSim	
headingSetSim	3
headingSim	4
MeshSimData	
mheadingSim	5
mnodeSim	5
nodeInfo	
nodeSim	
termInfo	7

9

Index

docInfo

## Description

Fetch information of a given article from PubMed.

## Usage

docInfo(pmid, verbose=FALSE, major=FALSE)

## Arguments

pmid	pmid of the desired article.
verbose	whether the title and abstract of the article should be print out.
major	whether only major terms should be returned.

## Value

Document information of given PMID including titile, abstract, MeSH headings

## Note

Network connection is required for using this function.

## Examples

docInfo("1111111")

docSim

Similarity between articles

## Description

Calculate the similarity between two articles.

## Usage

```
docSim(pmid1, pmid2, method="SP", frame="node", major=FALSE, env=NULL)
```

## Arguments

pmid1, pmid2	pmids of two articles whose similarity is needed to be calculated.
method	similarity measurment method, see Details for available methods.
frame	available options are node and heading, decide whether using node-based or heading-based methods.
env	the dataset to use.
major	whether the calculation should only be based on major terms

#### headingSetSim

#### Details

Available methods: SP: Shortest Path method, WL: Weighted Link method, WP: Wu and Palmer's method, LC: Leacock and Chodorow's method, Li: Li's method, Lord: Lord's method, Resnik: Resnik's method, Lin: Lin's method, JC: Jiang and Conrath's method.

## Value

Semantic similarity between two MEDLINE documents, whose value is between 0 and 1.

#### Note

Network connection is required for using this function.

#### Examples

docSim("1111113","1111111")

headingSetSim Similarity between two MeSH heading sets

#### Description

Calculate similarity between two MeSH heading sets.

#### Usage

```
headingSetSim(headingSet1, headingSet2, method="SP", frame="node", env=NULL)
```

#### Arguments

headingSet1,	headingSet2
	two lists of headings
method	similarity measurment method, see Details for available methods.
frame	available options are node and heading, decide whether using node-based or heading-based methods.
env	the dataset to use.

#### Details

Available methods: SP: Shortest Path method, WL: Weighted Link method, WP: Wu and Palmer's method, LC: Leacock and Chodorow's method, Li: Li's method, Lord: Lord's method, Resnik: Resnik's method, Lin: Lin's method, JC: Jiang and Conrath's method.

#### Value

Semantic similarity between two MeSH heading sets, whose value is between 0 and 1.

#### Examples

```
headingSet1<-c("Lumbosacral Region", "Body Regions")
headingSet2<-c("Body Regions", "Abdomen", "Abdominal Cavity")
headingSetSim(headingSet1,headingSet2,'SP','node')
```

headingSim

#### Description

Calculate similarity between two headings.

#### Usage

```
headingSim(heading1, heading2, method="SP", frame="node", env=NULL)
```

#### Arguments

heading1, head	ing2
	two headings or two lists of headings
method	similarity measurment method, see Details for available methods.
frame	available options are node and heading, decide whether using node-based or heading-based methods.
env	the dataset to use.

## Details

Available methods: SP: Shortest Path method, WL: Weighted Link method, WP: Wu and Palmer's method, LC: Leacock and Chodorow's method, Li: Li's method, Lord: Lord's method, Resnik: Resnik's method, Lin: Lin's method, JC: Jiang and Conrath's method.

#### Value

Semantic similarity between two MeSH headings, whose value is between 0 and 1.

#### Examples

headingSim("Lumbosacral Region", "Body Regions")

|--|

## Description

These contents data of the whole MeSH tree, as well as information contents for every node and term.

This dataset will be auto loaded by the first invoked function of this package, if no other dataset is specified.

mheadingSim

#### Description

Calculate similarity matrix between two heading lists.

#### Usage

```
mheadingSim(headingList1, headingList2, method="SP", frame="node", env=NULL)
```

#### Arguments

headingList1,	headingList2
	two headings or two lists of headings
method	similarity measurment method, see Details for available methods.
frame	available options are node and heading, decide whether using node-based or heading-based methods.
env	the dataset to use.

#### Details

Available methods: SP: Shortest Path method, WL: Weighted Link method, WP: Wu and Palmer's method, LC: Leacock and Chodorow's method, Li: Li's method, Lord: Lord's method, Resnik: Resnik's method, Lin: Lin's method, JC: Jiang and Conrath's method.

#### Value

Semantic similarity matrix between two MeSH heading lists.

#### Examples

```
headingList1<-c("Body Regions", "Abdomen", "Abdominal Cavity")
headingList2<-c("Lumbosacral Region", "Body Regions")
mheadingSim(headingList1,headingList2)</pre>
```

mnodeSim

Similarity between node lists

#### Description

Calculate similarity matrix between two MeSH node lists.

## Usage

```
mnodeSim(nodeList1, nodeList2, method="SP", frame="node", env=NULL)
```

nodeInfo

## Arguments

nodeList1, node	eList2
	two nodes or two lists of nodes
method	similarity measurment method, see Details for available methods.
frame	available options are node and heading, decide whether using node-based or heading-based methods.
env	the dataset to use.

#### Details

Available methods: SP: Shortest Path method, WL: Weighted Link method, WP: Wu and Palmer's method, LC: Leacock and Chodorow's method, Li: Li's method, Lord: Lord's method, Resnik: Resnik's method, Lin: Lin's method, JC: Jiang and Conrath's method.

## Value

Semantic similarity matrix between two MeSH node lists.

Details of nodes

## Examples

```
nodeList1<-c("B03.440.400.425.340.590", "B03.440.400.425.117.800.200")
nodeList2<-c("B03.440.400.425.340.590", "B03.440.400.425.117.800.200", "B03.440.400.425.127.100")
mnodeSim(nodeList1, nodeList2)</pre>
```

nodeInfo

## Description

This function returns a tree contains the given node.

## Usage

```
nodeInfo(node, brief, env=NULL)
```

#### Arguments

node	a node name.
brief	brief model for nodeInfo
env	the dataset to use.

## Value

Hierarchy information of node a

## Examples

```
nodeInfo("B03.440.400.425.127")
nodeInfo("B03.440.400", brief=TRUE)
```

6

nodeSim

#### Description

Calculate similarity between two MeSH nodes.

## Usage

```
nodeSim(node1, node2, method="SP", frame="node", env=NULL)
```

#### Arguments

node1, node2	two nodes or two lists of nodes
method	similarity measurment method, see Details for available methods.
frame	available options are node and heading, decide whether using node-based or heading-based methods.
env	the dataset to use.

## Details

Available methods: SP: Shortest Path method, WL: Weighted Link method, WP: Wu and Palmer's method, LC: Leacock and Chodorow's method, Li: Li's method, Lord: Lord's method, Resnik: Resnik's method, Lin: Lin's method, JC: Jiang and Conrath's method.

### Value

Semantic similarity between two MeSH nodes, whose value is between 0 and 1.

## Examples

nodeSim("B03.440.400.425.340.590", "B03.440.400.425.117.800.200")

termInfo

Details of MeSH terms

## Description

This function returns a tree contains the given term.

## Usage

termInfo(term, brief,env=NULL)

#### Arguments

term	a term name.
brief	whether to retrive breif tree information of MeSH term
env	the dataset to use.

termInfo

## Value

Hierarchy information of a given term

## Examples

termInfo("Body Regions")

# Index

mheadingSim, 5 mnodeSim, 5

nodeInfo, 6
nodeSim, 7

termInfo,7