

# Package ‘MeSHSim’

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**Type** Package

**Author** Jing Zhou, Yuxuan Shui

**Maintainer** Jing Zhou <12210240050@fudan.edu.cn>

**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

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docInfo	<i>Get details of documents</i>
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**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 title, abstract, MeSH headings

**Note**

Network connection is required for using this function.

**Examples**

```
docInfo("1111111")
```

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docSim	<i>Similarity between articles</i>
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**Description**

Calculate the similarity between two articles.

**Usage**

```
docSim(pmId1, pmId2, method="SP", frame="node", major=FALSE, env=NULL)
```

**Arguments**

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

**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")
```

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headingSetSim

*Similarity between two MeSH heading sets*

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**Description**

Calculate similarity between two MeSH heading sets.

**Usage**

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

**Arguments**

<code>headingSet1, headingSet2</code>	two lists of headings
<code>method</code>	similarity measurement method, see Details for available methods.
<code>frame</code>	available options are node and heading, decide whether using node-based or heading-based methods.
<code>env</code>	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**

*Similarity between headings*

## Description

Calculate similarity between two headings.

## Usage

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

## Arguments

heading1, heading2	two headings or two lists of headings
method	similarity measurement 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")
```

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**MeshSimData*****MeSH Dataset***

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**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.

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**mheadingSim*****Similarity between heading lists***

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**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 measurement 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)
```

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mnodeSim	<i>Similarity between node lists</i>
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## Description

Calculate similarity matrix between two MeSH node lists.

## Usage

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

## Arguments

nodeList1, nodeList2	two nodes or two lists of nodes
method	similarity measurement 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.

## 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)
```

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nodeInfo	<i>Details of nodes</i>
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**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)
```

---

nodeSim	<i>Similarity between nodes</i>
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**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 measurement 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 retrieve breif tree information of MeSH term
env	the dataset to use.

## Value

Hierarchy information of a given term

## Examples

```
termInfo("Body Regions")
```

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