

Package ‘screenmedR’

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Type Package

Title What the Package Does (Title Case)

Version 0.1.0

Author Who wrote it

Maintainer The package maintainer <yourself@somewhere.net>

Description More about what it does (maybe more than one line)

Use four spaces when indenting paragraphs within the Description.

License What license is it under?

Encoding UTF-8

LazyData true

RoxygenNote 7.1.1

Imports rentrez,

XML,

dplyr,

purrr,

tm,

stringr,

RISmed,

tidyr,

proxy,

lsa

Suggests knitr,

rmarkdown

VignetteBuilder knitr

R topics documented:

abstractsofgroup	2
diagnosis_clean	2
mesh_by_name	2
mesh_by_name_bq	3
mesh_clean	3
mesh_clean_bq	3
screenmed	4

Index	5
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abstractsofgroup	<i>Screens Pubmed PMID numbers in terms of those of 5 initial ones</i>
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Description

This package predicts which abstracts are closer in terms of words with an initial group of n abstracts that the user provides.

Usage

```
abstractsofgroup(clustering, group_number, initial_search)
```

Arguments

```
clustering, group_number, initial_search
```

diagnosis_clean	<i>Screens Pubmed PMID numbers in terms of those of 5 initial ones</i>
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Description

This package predicts which abstracts are closer in terms of words with an initial group of n abstracts that the user provides.

Usage

```
diagnosis_clean(x)
```

Arguments

```
x
```

mesh_by_name	<i>Screens Pubmed PMID numbers in terms of the names of Descriptors & Qualifiers</i>
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Usage

```
mesh_by_name(x, Descriptor, Qualifier)
```

Arguments

```
x, Descriptor, Qualifier
```

mesh_by_name_bq	<i>Screens Pubmed PMID numbers in terms of the names of Descriptors & Qualifiers</i>
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Usage

```
mesh_by_name_bq(x, d, q)
```

Arguments

x, Descriptor, Qualifier

mesh_clean	<i>Screens Pubmed PMID numbers in terms of those of 5 initial ones</i>
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Description

This package predicts which abstracts are closer in terms of words with an initial group of n abstracts that the user provides.

Usage

```
mesh_clean(x, y, d, q)
```

Arguments

x, y, d, q

mesh_clean_bq	<i>Screens Pubmed PMID numbers in terms of those of 5 initial ones</i>
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Description

This package predicts which abstracts are closer in terms of words with an initial group of n abstracts that the user provides.

Usage

```
mesh_clean_bq(x, y, d, q)
```

Arguments

x, y, d, q

screenmed*Screens Pubmed PMID numbers in terms of those of 5 initial ones*

Description

This package predicts which abstracts are closer in terms of words with an initial group of n abstracts that the user provides.

Usage

```
screenmed(initial_search, filtered, sparsity, group_number)
```

Arguments

```
initial_search, filtered, sparsity, group_number
```

Index

`abstractsofgroup`, [2](#)

`diagnosis_clean`, [2](#)

`mesh_by_name`, [2](#)

`mesh_by_name_bq`, [3](#)

`mesh_clean`, [3](#)

`mesh_clean_bq`, [3](#)

`screenmed`, [4](#)