Package 'specieshindex'

November 12, 2020

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Type Package	
Title How (scientifically) popular is a given species?	
Version 0.0.1	
Date 2020-07-10	
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Description Finds the h-index of a species.	
Depends R (>= 3.1.0)	
LazyData true	
Imports rscopus, taxize, dplyr, httr, XML, rlang	
Suggests testthat,	
knitr, roxygen2, devtools, rmarkdown, ggplot2	
License MIT + file LICENSE	
<pre>URL https://github.com/jessicatytam/Honours</pre>	
RoxygenNote 7.1.1	
VignetteBuilder knitr	
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Allindices

Index summary

Description

This function returns a dataframe of the summary of all of the indices.

Usage

Index

```
Allindices(data, genus, species)
```

Arguments

data The dataframe generated from FetchSpT or FetchSpTAK.

genus Genus classification from the binomial name.
species Species classification from the binomial name.

Value

A datarame of all of the indices in the package.

```
data(Woylie)
Allindices(Woylie, genus = "genus_name", species = "species_name")
```

ARRatio 3

ARRatio Article: Review ratio

Description

This function calculates the percentage ratio of article:rerview.

Usage

```
ARRatio(data)
```

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

A character value of the percentage ratio of the number of articles and reviews.

Examples

```
data(Woylie)
ARRatio(Woylie)
```

CountSpT

Search count - title only

Description

This function counts the total number of search results. It counts only the publications with the binomial name in their title. A check will be conducted via gnr_resolve to validify the genus and species names.

Usage

```
CountSpT(genus, species, APIkey, datatype = "application/xml")
```

Arguments

genus	Genus classification from the binomial name.
species	Species classification from the binomial name.
APIkey	Scopus API key needed to access and download data from their database.

datatype Formats the URL to be sent to the API. The default is "application/xml".

Value

Search count of the species with the given genus and species.

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References

Chamberlain, S. & Szocs, E. (2013). taxize - taxonomic search and retrieval in R. *F1000Research*, 2, 191.

CountSpTAK

Search count - title, abstract and keywords

Description

This function counts the total number of search results. It counts the publications with the binomial name in the title, abstract and keywords. A check will be conducted via gnr_resolve to validify the genus and species names.

Usage

```
CountSpTAK(genus, species, APIkey, datatype = "application/xml")
```

Arguments

genus Genus classification from the binomial name.

species Species classification from the binomial name.

APIkey Scopus API key needed to access and download data from their database.

datatype Formats the URL to be sent to the API. The default is "application/xml".

Value

Search count of the species with the given genus and species.

References

Chamberlain, S. & Szocs, E. (2013). taxize - taxonomic search and retrieval in R. *F1000Research*, 2, 191.

```
## Not run:
CountSpTAK("Bettongia", "penicillata", "myAPI")

#lower case letter in genus is also accepted and will return identical results
CountSpTAK("bettongia", "penicillata", "myAPI")

## End(Not run)
```

FetchSpT 5

Description

This function fetches citation information from Scopus using genus and species name found in the title of the publications. Duplicates are removed after fetching the data.

Usage

```
FetchSpT(genus, species, APIkey)
```

Arguments

genus Genus classification from the binomial name.
species Species classification from the binomial name.

APIkey Scopus API key needed to access and download data from their database.

Value

A dataframe of the species' citation records with the given genus and species.

Examples

```
## Not run:
FetchSpT("Bettongia", "penicillata", "myAPI")

#lower case letter in genus is also accepted and will return identical results
FetchSpT("bettongia", "penicillata", "myAPI")
## End(Not run)
```

FetchSpTAK

Fetch data - title, abstract and keywords

Description

This function fetches citation information from Scopus using genus and species name found in the title, abstract and keywords of the publications. Duplicates are removed after fetching the data.

Usage

```
FetchSpTAK(genus, species, APIkey)
```

Arguments

genus Genus classification from the binomial name.
species Species classification from the binomial name.

APIkey Scopus API key needed to access and download data from their database.

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Value

A dataframe of the species' citation records with the given genus and species.

Examples

```
## Not run:
FetchSpT("Bettongia", "penicillata", "myAPI")

#lower case letter in genus is also accepted and will return identical results
FetchSpT("bettongia", "penicillata", "myAPI")
## End(Not run)
```

Koala

Koala dataset

Description

Citation records of koala (Phascolarctos cinereus) from Scopus. Data was retrieved on 10 July 2020.

Usage

Koala

Format

A data frame with 773 rows and 20 variables

Source

http://api.elsevier.com/content/search/scopus

Platypus

Platypus dataset

Description

Citation records of platypus (Ornithorhynchus anatinus) from Scopus. Data was retrieved on 10 July 2020.

Usage

Platypus

Format

A data frame with 321 rows and 20 variables

Source

http://api.elsevier.com/content/search/scopus

Quokka 7

Quokka

Quokka dataset

Description

Citation records of quokka (Setonix brachyurus) from Scopus. Data was retrieved on 10 July 2020.

Usage

Quokka

Format

A data frame with 242 rows and 20 variables

Source

```
http://api.elsevier.com/content/search/scopus
```

SpH5

Species h5 index

Description

This function calculates the h-index of a species in the past 5 years.

Usage

SpH5(data)

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

H5 index.

References

Suzuki, H. (2012). *Google Scholar Metrics for Publications*. Retrieved from https://scholar.googleblog.com/2012/04/google-scholar-metrics-for-publications.html.

```
data(Woylie)
SpH5(Woylie)
```

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SpHAfterdate

Species h-index with a given time frame

Description

This function calculates the h-index using a given date up till the newest record.

Usage

```
SpHAfterdate(data, date)
```

Arguments

data The dataframe generated from FetchSpT or FetchSpTAK.

date The lower limit of the timeframe.

Value

H-index of the given time period.

Examples

```
data(Woylie)
SpHAfterdate(Woylie, "2000-01-01")
```

SpHindex

Species h-index

Description

This function calculates the h-index of a species.

Usage

```
SpHindex(data)
```

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

H-index.

References

Bertoli-Barsotti, L. & Lando, T. (2015). On a formula for the h-index. *Journal of Informetrics*, 9(4), 762-776.

Hirsch, J. (2005). An index to quantify an individual's scientific research output. *Proceedings of the National Academy of Sciences of the United States of America*, 102(46), 16569-16572.

Spi10 9

Examples

```
data(Woylie)
SpHindex(Woylie)
```

Spi10

Species i10 index

Description

This function calculates the i10 index of a species. i10 index counts all of the publications with 10 or more citations.

Usage

```
Spi10(data)
```

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

i10 index.

References

Cornell University (2019). *i10-index*. Retrieved from https://guides.library.cornell.edu/c.php?g=32272&p=203393.

Examples

```
data(Woylie)
Spi10(Woylie)
```

SpMindex

Species m-index

Description

This function calculates the m-index of species. M-index uses the h-index and divides it by the number of years of activity.

Usage

```
SpMindex(data)
```

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

TotalArt

Value

M-index.

References

University of Pittsburgh (2019). *Research Impact and Metrics: Author metrics*. Retrieved from https://pitt.libguides.com/bibliometricIndicators/AuthorMetrics.

Examples

```
data(Woylie)
SpMindex(Woylie)
```

TotalArt

Total Article

Description

This function calculates the total number of articles.

Usage

TotalArt(data)

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

An integer of the total number of articles.

```
data(Woylie)
TotalArt(Woylie)
```

TotalCite 11

TotalCite

Total citations

Description

This function calculates the total number of citations.

Usage

```
TotalCite(data)
```

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

A numerical value of the total number of citations.

Examples

```
data(Woylie)
TotalCite(Woylie)
```

TotalJournals

Total journals

Description

This function calculates the total number of journals.

Usage

```
TotalJournals(data)
```

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

An integer of the total number of journals.

```
data(Woylie)
TotalJournals(Woylie)
```

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TotalPub

Total publications

Description

This function calculates the total number of publications.

Usage

```
TotalPub(data)
```

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

An integer of the total number of publications.

Examples

```
data(Woylie)
TotalPub(Woylie)
```

TotalRev

Total reviews

Description

This function calculates the total number of reviews.

Usage

```
TotalRev(data)
```

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

An integer of the total number of reviews.

```
data(Woylie)
TotalRev(Woylie)
```

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Woylie

Woylie dataset

Description

Citation records of woylie (Bettongia penicillata) from Scopus. Data was retrieved on 10 July 2020.

Usage

Woylie

Format

A data frame with 113 rows and 20 variables

Source

http://api.elsevier.com/content/search/scopus

 ${\it YearsPublishing}$

Years since first publication

Description

The number of years since the first publication in relation to the species.

Usage

YearsPublishing(data)

Arguments

data

The dataframe generated from FetchSpT or FetchSpTAK.

Value

Number of years.

```
data(Woylie)
YearsPublishing(Woylie)
```

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