

taxize: taxonomic search and retrieval in R

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Summary

Taxize is a R package that provides an interface to various taxonomic data sources around the web. We hope that taxize will be

Data Sources



Taxize currently provides simple and programmatic access to taxonomic data from 14 data sources around the web.

Features

Resolve taxonomic names

- We often have a list of species names and we want to know
- a) if we have the most up-to-date names,
 - b) if our names are spelled correctly,
 - c) and the scientific name for a common name.
- Taxize provides an interface to the EOL Global Names Resolver and Taxonomic Name Resolution Service, e.g.

```
gnr_resolve('Baetis roodani')
```

```
##      submitted_name  matched_name
## 1 Baetis roodani Baetis rhodani
```

Retrieve higher taxonomic names

Another common

```
classification('Baetis rhodani', db='col')
```

```
##      name      rank
## 1 Animalia Kingdom
## 2 Arthropoda Phylum
## 3 Insecta    Class
## 4 Ephemeroptera Order
## 5 Baetoidea Superfamily
## 6 Baetidae   Family
## 7 Baetis     Genus
## 8 Baetis rhodani Species
```

Retrieve children taxa

```
downstream('Baetis', db = 'col', downto = 'Species')
```

```
##      childtaxa_name childtaxa_rank
## 1 Baetis acceptus Species
## 2 Baetis aculeatus Species
## 3 Baetis acuminatus Species
## 4 Baetis adonis Species
## 5 Baetis aeneus Species
```

Aggregate data to a specific taxonomic rank

test

```
1+1
## [1] 2
```

Contribution

taxize is currently developed collaboratively in github. Feature requests, bug reports and contributions are strongly encouraged!

 <https://github.com/ropensci/taxize>

