

# Taxonomic Data

## with taxize

### Cheat Sheet



#### Use Cases

##### Get taxonomic IDs

**taxize::get\_ids/get\_colid/etc.**

*Puma concolor* → 2435099  
*Ursus americanus* → 2433407  
*Quercus robur* → 2878688

##### Get correct spellings.

**taxize::tnrs/taxize::gnr\_resolve**

*Pumma concolor* → *Puma concolor*

##### Retrieve higher taxonomic classification.

**taxize::classification**

Classification for *Chironomus riparius*

	name	rank	id
1	Animalia	Kingdom	22254411
2	Arthropoda	Phylum	22254412
3	Insecta	Class	22254618
4	Diptera	Order	22254663
5	Chironomidae	Family	22254676
6	Chironomus	Genus	22350527
7	Chironomus riparius	Species	8663146

##### Fetch all taxonomic names downstream from a target taxonomic group.

**taxize::downstream**

Species downstream from the genus *Apis*

	childtaxa_id	childtaxa_name	childtaxa_rank
1	6971712	Apis andreniformis	Species
2	6971713	Apis cerana	Species
3	6971714	Apis dorsata	Species
4	6971715	Apis florea	Species
5	6971716	Apis koschevnikovi	Species
6	6845885	Apis mellifera	Species
7	6971717	Apis nigrocincta	Species

#### Data Sources

[Encyclopedia of Life](#)  
[Taxonomic Name Resolution Service](#)  
[Integrated Taxonomic Information Service](#)  
[Phylomatic](#)  
[uBio](#)  
[Global Names Resolver](#)  
[Global Names Index](#)  
[IUCN Red List](#)  
[Tropicos](#)  
[Plantminer](#)  
[Theplantlist](#)  
[Catalogue of Life](#)  
[Global Invasive Species Database](#)  
[National Center for Biotechnology Information](#)  
[CANADENSYS Vscan name search API](#)  
[International Plant Names Index \(IPNI\)](#)  
[Barcode of Life Data Systems \(BOLD\)](#)  
[National Biodiversity Network \(UK\)](#)

#### Get taxonomic IDs

##### Options

- ☒ Interactively select names.
- ☒ Or, get back all names to process later.
- ☒ Or, get back certain records.

**taxize::get\_colid(<id>)**

If > 1 result, interactively select by row

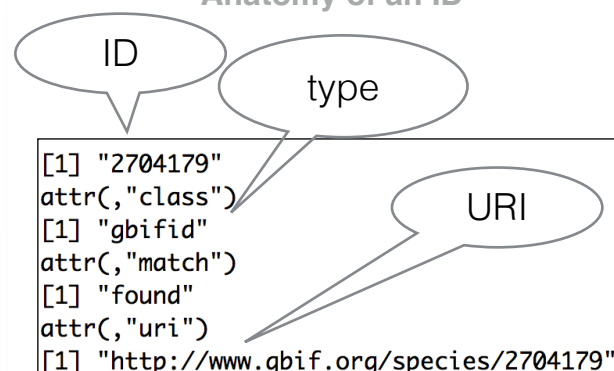
**taxize::get\_colid(<id>, rows = 1:4)**

Get rows 1 through 4

**taxize::get\_colid\_(<id>)**

Get all data

##### Anatomy of an ID



#### Correct Names

##### Out of the box frameworks

##### Global Names Resolver

**taxize::gnr\_resolve**(c("Poa anua", "Quercus rober"), data\_source\_ids = 11)

	submitted_name	matched_name
1	Poa anua	Poa annua L.
2	Quercus rober	Quercus robur L.

Notice that both submitted names are spelled incorrectly, and are matched to their correct spellings.

##### Others

**taxize::tnrs(<names>)**

Queries NCBI, Mammals dataset, and iPlant, only fuzzy matching on plants

**taxize::iplant\_resolve(<names>)**

iPlant has a nice name resolver - of course, it only works for plants

**Bonus:** **taxize::resolve(<names>)** is an interface to all three of the resolution functions

##### DIY correction

There are a lot of functions to search various taxonomic databases - they don't do name resolution per se. But after getting an id, you can do many other tasks. Some examples:

**taxize::synonyms(<id>)**

Get name synonyms

**taxize::ncbi\_search(<id>)**

Search NCBI

**taxize::tp\_search(<id>)**

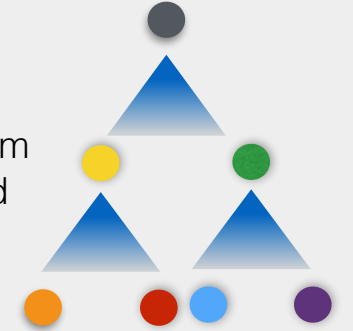
Search Tropicos

#### Down-/up-stream, children, and more

● Target Taxon

**taxize::downstream(<name>, "species")**

Downstream to desired rank



**taxize::children(<name>)**

Immediate children only



**taxize::upstream(<name>)**

Upstream to desired rank



**taxize::classification(<name>)**

