

# Package: rtaxref (via r-universe)

September 9, 2024

**Type** Package

**Title** An R client for TaxRef the French Taxonomical Database

**Version** 0.1.0

**Description** Provides an R client to the TaxRef API

<<https://taxref.mnhn.fr/taxref-web/api/doc>>, the French Taxonomical Reference Database which indexes names of species with unique identifiers as well as conservation statuses, biological interactions and taxonomic relationships.

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**Imports** httr, tibble

**Suggests** testthat, knitr, vcr, rmarkdown

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.2.0

**Language** en-US

**VignetteBuilder** knitr

**URL** <https://github.com/Rekyt/rtaxref>

**BugReports** <https://github.com/Rekyt/rtaxref/issues>

**Repository** <https://rekyt.r-universe.dev>

**RemoteUrl** <https://github.com/Rekyt/rtaxref>

**RemoteRef** HEAD

**RemoteSha** bf897635dd5eb011d483551c480aff5ace80e435

## Contents

rt_biogeo_status . . . . .	2
rt_external_db . . . . .	3
rt_habitats . . . . .	3

rt_interactions_find_type . . . . .	4
rt_interactions_opgroup . . . . .	5
rt_interactions_search . . . . .	5
rt_interactions_type . . . . .	6
rt_languages . . . . .	7
rt_operational_groups . . . . .	7
rt_sources_find_term . . . . .	8
rt_sources_id . . . . .	8
rt_sources_match . . . . .	9
rt_source_uses . . . . .	9
rt_status_find_type . . . . .	10
rt_status_opgroup . . . . .	11
rt_status_search . . . . .	11
rt_status_type . . . . .	13
rt_taxa_autocomplete . . . . .	13
rt_taxa_children . . . . .	15
rt_taxa_externalids . . . . .	15
rt_taxa_factsheet . . . . .	16
rt_taxa_from_externalids . . . . .	16
rt_taxa_fuzzymatch . . . . .	17
rt_taxa_id . . . . .	17
rt_taxa_interactions . . . . .	18
rt_taxa_media . . . . .	18
rt_taxa_parent . . . . .	19
rt_taxa_search . . . . .	19
rt_taxa_sources . . . . .	22
rt_taxa_status . . . . .	23
rt_taxa_synonyms . . . . .	23
rt_taxa_taxrefhistory . . . . .	24
rt_taxa_vernacular . . . . .	24
rt_taxo_ranks . . . . .	25
rt_taxref_versions . . . . .	25
rt_vernacular_groups . . . . .	26

**Index****27**


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<b>rt_biogeo_status</b>	<i>Retrieve biogeographic statuses used in TAXREF</i>
-------------------------	---

---

**Description**

If the function is used without arguments returns the list of biogeographic statuses used in TAXREF

**Usage**

```
rt_biogeo_status(biogeo_id = NULL)
```

**Arguments**

biogeo\_id      NULL or character(1) [default = NULL]  
A biogeographic status id used in TAXREF

**Examples**

```
## Not run: rt_biogeo_status(biogeo_id = "P")
```

---

**rt\_external\_db**      *Retrieve external DBs referenced in TAXREF*

---

**Description**

If the function is used without arguments returns the list of external databases used in TAXREF.

**Usage**

```
rt_external_db(db_id = NULL)
```

**Arguments**

db\_id      NULL or character(1) [default = NULL]  
The id of an external db referenced in TAXREF

**Examples**

```
## Not run: rt_external_db(db_id = "worms")
```

---

**rt\_habitats**      *Retrieve habitats used in TAXREF*

---

**Description**

If the function is used without arguments returns the list of habitats used in TAXREF.

**Usage**

```
rt_habitats(habitat_id = NULL)
```

**Arguments**

habitat\_id      NULL or integer(1) [default = NULL]  
The id of a habitat referenced in TAXREF

## Habitats

- 1 = Marine
- 2 = Freshwater
- 3 = Terrestrial
- 4 = Marine and freshwater
- 5 = Marine and Terrestrial
- 6 = Brackish water
- 7 = Continental (terrestrial and/or freshwater)
- 8 = Continental (terrestrial and freshwater)

## Examples

```
## Not run: rt_habitats(habitat_id = 8)
```

## *rt\_interactions\_find\_type*

*Retrieve the status of all Taxa according to a Status Type*

## Description

Retrieve the status of all Taxa according to a Status Type

## Usage

```
rt_interactions_find_type(interaction_id = NULL, page = 1, size = 2000)
```

## Arguments

<code>interaction_id</code>	<code>character(1) [required]</code>
	The identifier of status (see <a href="#">rt_interactions_type()</a> for the list of status types)
<code>page</code>	<code>integer(1) [default = 1]</code>
	The page number returned
<code>size</code>	<code>integer(1) [default = 2000]</code>
	The number of lines returned per page (maximum = 2000)

## See Also

the list of status types [rt\\_interactions\\_type\(\)](#)

## Examples

```
## Not run: rt_interactions_find_type(interaction_id = "GALLES")
```

---

**rt\_interactions\_opgroup**

*Retrieve the interactions of all Taxa in an Operational Group*

---

**Description**

Retrieve the interactions of all Taxa in an Operational Group

**Usage**

```
rt_interactions_opgroup(opgroup_id = NULL, page = 1, size = 2000)
```

**Arguments**

opgroup_id	integer(1) [required]
	identifier of the operational group (see the list of operational groups <a href="#">rt_operational_groups()</a> )
page	integer(1) [default = 1]
	The page number returned
size	integer(1) [default = 2000]
	The number of lines returned per page (maximum = 2000)

**See Also**

the list of operational groups [rt\\_operational\\_groups\(\)](#)

**Examples**

```
## Not run: rt_interactions_opgroup(opgroup_id = 706)
```

---

**rt\_interactions\_search**

*Search Interactions based on a list of criteria*

---

**Description**

Search Interactions based on a list of criteria

**Usage**

```
rt_interactions_search(  
  id = NULL,  
  interaction_id = NULL,  
  opgroup_id = NULL,  
  page = 1,  
  size = 2000  
)
```

## Arguments

<code>id</code>	<code>NULL or integer(1+)</code> [default = <code>NULL</code> ] One or more integer giving the id(s) of taxa in TAXREF (also called cdNom)
<code>interaction_id</code>	<code>NULL or character(1)</code> [default = <code>NULL</code> ] A character identifying the type of interactions
<code>opgroup_id</code>	<code>NULL or numeric(1)</code> [default = <code>NULL</code> ] the id of an operational group referenced in TAXREF
<code>page</code>	<code>integer(1)</code> [default = 1] The page number returned
<code>size</code>	<code>integer(1)</code> [default = 2000] The number of lines returned per page (maximum = 2000)

## See Also

the list of interaction type `rt_interactions_type()` and the list of operational groups `rt_operational_groups()`

## Examples

```
## Not run:
# Interactors with 'Ophrys apifera' (id = 110335)
rt_interactions_search(id = 110335)

## End(Not run)
```

`rt_interactions_type`    *Retrieve Interactions Types used in TAXREF*

## Description

If used without argument returns the list of all interactions used in TAXREF

## Usage

```
rt_interactions_type(interaction_id = NULL)
```

## Arguments

<code>interaction_id</code>	<code>NULL or character(1)</code> [default = <code>NULL</code> ] A character identifying the type of interactions
-----------------------------	--

## Examples

```
## Not run: rt_interactions_type(interaction_id = "PREDATEUR")
```

---

rt_languages	<i>Retrieve the languages used in TAXREF</i>
--------------	--

---

### Description

If used without argument returns the list of languages used in TAXREF otherwise the specific

### Usage

```
rt_languages(languages_id = NULL)
```

### Arguments

languages_id	NULL or character(1) [default = NULL] three-letter lowercase code that is the id of a language referenced in TAXREF
--------------	--

### Examples

```
## Not run:  
# Retrieve the TAXREF definition in the Wayana Language  
rt_languages(languages_id = "way")  
  
## End(Not run)
```

---

rt_operational_groups	<i>Retrieve operational groups used in TAXREF</i>
-----------------------	---

---

### Description

Operational Groups are names of groups either taxonomic or vernacular used across TAXREF in French. If used without argument returns the list of all operational groups used in TAXREF.

### Usage

```
rt_operational_groups(opgroup_id = NULL)
```

### Arguments

opgroup_id	NULL or numeric(1) [default = NULL] the id of an operational group referenced in TAXREF
------------	--

### Examples

```
## Not run: rt_operational_groups(opgroup_id = 143)
```

**rt\_sources\_find\_term** *Search a source from a term*

### Description

Search a source from a term

### Usage

```
rt_sources_find_term(term = NULL)
```

### Arguments

term	character(1) <b>[required]</b>
	A term contained in the source

### Examples

```
## Not run:
rt_sources_find_term(term = "Ophrys")

## End(Not run)
```

**rt\_sources\_id** *Retrieve a source of taxonomic information using its id*

### Description

Retrieve a source of taxonomic information using its id

### Usage

```
rt_sources_id(source_id, bibtex = FALSE)
```

### Arguments

source_id	integer(1) <b>[required]</b>
	an integer giving the id of the source in TAXREF
bibtex	logical(1) [default = FALSE]
	should citation be returned as BibTeX?

## Examples

```
## Not run:  
rt_sources_id(source_id = 160688, bibtex = FALSE)  
rt_sources_id(source_id = 160688, bibtex = TRUE)  
  
## End(Not run)
```

---

rt\_sources\_match      *Search a source from a citation by fuzzy matching*

---

## Description

Search a source from a citation by fuzzy matching

## Usage

```
rt_sources_match(citation = NULL)
```

## Arguments

citation	character(1) [required]
	A string that could partially match the citation

## Examples

```
## Not run:  
rt_sources_match(  
  paste0("Chapman, T. A. 1909. Callophrys avis a new butterfly on ",  
        "southern France. The Entomologist's record and journal of ",  
        "variation, 21: 130-131.")  
)  
  
## End(Not run)
```

---

rt\_source\_uses      *Retrieve the list of source uses in TAXREF*

---

## Description

TAXREF references many publications across its documentation. The different sources are used for different reasons. This function returns the list of usages for a source when no arguments are provided, otherwise returns the specific type of use.

## Usage

```
rt_source_uses(usage_id = NULL)
```

**Arguments**

`usage_id`      NULL or numeric(1) [default = NULL]  
                   the id of a source use in TAXREF

**Examples**

```
## Not run: rt_source_uses(usage_id = 276)
```

`rt_status_find_type`      *Retrieve the status of all Taxa according to a Status Type*

**Description**

Retrieve the status of all Taxa according to a Status Type

**Usage**

```
rt_status_find_type(status_id = NULL, page = 1, size = 2000)
```

**Arguments**

<code>status_id</code>	character(1) <b>[required]</b>
	The identifier of status (see <a href="#">rt_status_type()</a> for the list of status types)
<code>page</code>	integer(1) [default = 1]
	The page number returned
<code>size</code>	integer(1) [default = 2000]
	The number of lines returned per page (maximum = 2000)

**See Also**

the list of status types [rt\\_status\\_type\(\)](#)

**Examples**

```
## Not run:  

# Find all taxa concerned by Habitat Directive referenced in TAXREF  

rt_status_find_type("DH")  
  

## End(Not run)
```

---

rt\_status\_opgroup      *Retrieve the status of all Taxa in an Operational Group*

---

## Description

Retrieve the status of all Taxa in an Operational Group

## Usage

```
rt_status_opgroup(opgroup_id, page = 1, size = 2000)
```

## Arguments

opgroup_id	integer(1) [required]
	identifier of the operational group (see the list of operational groups <a href="#">rt_operational_groups()</a> )
page	integer(1) [default = 1]
	The page number returned
size	integer(1) [default = 2000]
	The number of lines returned per page (maximum = 2000)

## See Also

the list of operational groups [rt\\_operational\\_groups\(\)](#)

## Examples

```
## Not run:  
  rt_status_opgroup(opgroup_id = 706)  
  
## End(Not run)
```

---

rt\_status\_search      *Search statuses based on a list of criteria*

---

## Description

Search statuses based on a list of criteria

**Usage**

```
rt_status_search(
  id = NULL,
  opgroup_id = NULL,
  location_id = NULL,
  latitude = NULL,
  longitude = NULL,
  page = 1,
  size = 2000,
  detail = FALSE
)
```

**Arguments**

<code>id</code>	<code>integer(1) [required]</code>
	an integer giving the id of the taxon in TAXREF (also called cdNom)
<code>opgroup_id</code>	<code>integer(1) [required]</code>
	identifier of the operational group (see the list of operational groups <a href="#">rt_operational_groups()</a> )
<code>location_id</code>	<code>NULL or character(1) [default = NULL]</code>
	a string identifying a location
<code>latitude</code>	<code>NULL or numeric(1) [default = NULL]</code>
	a latitude value
<code>longitude</code>	<code>NULL or numeric(1) [default = NULL]</code>
	a longitude value
<code>page</code>	<code>integer(1) [default = 1]</code>
	The page number returned
<code>size</code>	<code>integer(1) [default = 2000]</code>
	The number of lines returned per page (maximum = 2000)
<code>detail</code>	<code>logical(1) [default = FALSE]</code>
	Should the table be a long table detailing all the statuses or a compact table?

**See Also**

list of operational groups [rt\\_operational\\_groups\(\)](#)

**Examples**

```
## Not run:
rt_status_search(id = 443800, detail = FALSE)
rt_status_search(id = 443800, detail = TRUE)

## End(Not run)
```

---

**rt\_status\_type**      *Retrieve Status Types Used in TAXREF*

---

**Description**

If the function is provided without arguments returns the entire list of status types used in TAXREF.

**Usage**

```
rt_status_type(status_id = NULL)
```

**Arguments**

status_id	NULL or character(1) [default = NULL] A character identifying the type of status
-----------	---

---

**rt\_taxa\_autocomplete**      *Retrieve Taxon ID from partially matching its Name*

---

**Description**

Retrieve Taxon ID from partially matching its Name

**Usage**

```
rt_taxa_autocomplete(  
  term = NULL,  
  territories = NULL,  
  rank = NULL,  
  domain = NULL,  
  page = 1,  
  size = 200  
)
```

**Arguments**

term	character(1) <b>[required]</b> the beginning of the scientific name of the searched taxa
territories	NULL or character(1+) [default = NULL] One or more territories where species is present: <ul style="list-style-type: none"><li>• "fr" = Mainland France</li><li>• "gf" = French Guiana</li><li>• "gua" = Guadeloupe</li><li>• "mar" = Martinique</li></ul>

	<ul style="list-style-type: none"> <li>• "sm" = Saint Martin</li> <li>• "sb" = Saint Barthélemy</li> <li>• "spm" = Saint Pierre and Miquelon</li> <li>• "epa" = Scattered Islands in the Indian Ocean</li> <li>• "may" = Mayotte</li> <li>• "reu" = Réunion</li> <li>• "sa" = French Southern and Antarctic Lands</li> <li>• "ta" = Adélie Land</li> <li>• "nc" = New Caledonia</li> <li>• "wf" = Wallis and Futuna</li> <li>• "pf" = French Polynesia</li> <li>• "cli" = Clipperton Island</li> </ul>
rank	<p>NULL or character(1+) [default = NULL]</p> <p>One of more rank of the searched taxa (see <a href="#">rt_taxo_ranks()</a> for the list of all ranks available):</p> <ul style="list-style-type: none"> <li>• Dumm= Domain</li> <li>• KD= Kingdom</li> <li>• PH= Phylum</li> <li>• CL= Class</li> <li>• OR= Order</li> <li>• FM= Family</li> <li>• SBFM= Subfamily</li> <li>• TR= Tribe</li> <li>• GN= Genus</li> <li>• AGES= Aggregate</li> <li>• ES= Species</li> <li>• SSES= Subspecies</li> <li>• NAT= Natio</li> <li>• VAR= Variety</li> <li>• SVAR= Sub-variety</li> <li>• FO= Form</li> <li>• SSFO= Sub-form</li> <li>• RACE= Race</li> <li>• CAR= Cultivar</li> <li>• AB= Abberatio</li> </ul>
domain	<p>NULL or character(1) [default = NULL]</p> <p>The domain where the species is found:</p> <ul style="list-style-type: none"> <li>• "marin" = Marine</li> <li>• "continental" = Continental</li> </ul>
page	<p>integer(1) [default = 1]</p> <p>The page number returned</p>
size	<p>integer(1) [default = 200]</p> <p>The number of lines returned per page (maximum = 200)</p>

**See Also**

the list of taxonomic ranks available through [rt\\_taxo\\_ranks\(\)](#)

**Examples**

```
## Not run:  
rt_taxa_autocomplete(term = "Bradypus")  
  
## End(Not run)
```

---

**rt\_taxa\_children**      *Retrieve a taxon children*

---

**Description**

Retrieve a taxon children

**Usage**

```
rt_taxa_children(id)
```

**Arguments**

<b>id</b>	<b>integer(1) [required]</b>
	an integer giving the id of the taxon in TAXREF (also called cdNom)

**Examples**

```
## Not run: rt_taxa_children(id = 442585)
```

---

**rt\_taxa\_externalids**      *Retrieve IDs of the taxon in external databases*

---

**Description**

Retrieve IDs of the taxon in external databases

**Usage**

```
rt_taxa_externalids(id = NULL)
```

**Arguments**

<b>id</b>	<b>integer(1) [required]</b>
	an integer giving the id of the taxon in TAXREF (also called cdNom)

## Examples

```
## Not run: rt_taxa_externalids(id = 443800)
```

**rt\_taxa\_factsheet**      *Get factsheet for taxa*

## Description

For some taxa, a factsheet has been written. This text, often in French, describes the general ecology and some diagnosis of the submitted taxon.

## Usage

```
rt_taxa_factsheet(id = NULL)
```

## Arguments

id	integer(1) [required] an integer giving the id of the taxon in TAXREF (also called cdNom)
----	--

## Details

Beware, many taxa have no corresponding factsheet.

## Examples

```
## Not run: rt_taxa_factsheet(4540)
```

**rt\_taxa\_from\_externalids**  
*Retrieve Taxon using the ids from external databases*

## Description

Retrieve Taxon using the ids from external databases

## Usage

```
rt_taxa_from_externalids(external_db = NULL, external_id = NULL)
```

### Arguments

external_db	character(1) [required]
	The name of an external database see <a href="#">rt_external_db()</a> for the list of external database names
external_id	character(1) [required]
	The identifier of the taxon in the specified external database

### See Also

[rt\\_external\\_db\(\)](#) for the list of external databases linked to TAXREF; [rt\\_taxa\\_externalids\(\)](#) to get the external ids associated with a taxon in TAXREF

### Examples

```
## Not run: rt_taxa_from_externalids("TROPICOS", 26600197)
```

---

**rt\_taxa\_fuzzymatch**      *Find taxa by fuzzy matching their scientific Name*

---

### Description

Find taxa by fuzzy matching their scientific Name

### Usage

```
rt_taxa_fuzzymatch(sciname = NULL)
```

### Arguments

sciname	character(1) [required]
	a string giving the scientific name of the taxon with or without the author name

---

**rt\_taxa\_id**      *Retrieve a taxon information using its id*

---

### Description

Retrieve a taxon information using its id

### Usage

```
rt_taxa_id(id = NULL)
```

**Arguments**

id	<b>integer(1) [required]</b>
	an integer giving the id of the taxon in TAXREF (also called cdNom)

**Examples**

```
## Not run:
rt_taxa_id(id = 443800)
rt_taxa_id(id = 110335)

## End(Not run)
```

**rt\_taxa\_interactions**    *Retrieve Interactions of taxon with other taxa*

**Description**

Retrieve Interactions of taxon with other taxa

**Usage**

```
rt_taxa_interactions(id = NULL)
```

**Arguments**

id	<b>integer(1) [required]</b>
	an integer giving the id of the taxon in TAXREF (also called cdNom)

**Examples**

```
## Not run: rt_taxa_interactions(id = 110335)
```

**rt\_taxa\_media**    *Retrieve Media (photos & others) Associated with a Taxon*

**Description**

Retrieve Media (photos & others) Associated with a Taxon

**Usage**

```
rt_taxa_media(id = NULL)
```

**Arguments**

id	integer(1) [required]
	an integer giving the id of the taxon in TAXREF (also called cdNom)

**Examples**

```
## Not run: rt_taxa_media(id = 443800)
```

---

rt\_taxa\_parent      *Retrieve classification above the taxon*

---

**Description**

Retrieve classification above the taxon

**Usage**

```
rt_taxa_parent(id = NULL)
```

**Arguments**

id	integer(1) [required]
	an integer giving the id of the taxon in TAXREF (also called cdNom)

**Examples**

```
## Not run: rt_taxa_parent(id = 443800)
```

---

rt\_taxa\_search      *Search for a taxon information*

---

**Description**

Search for a taxon information

**Usage**

```
rt_taxa_search(
  id = NULL,
  sciname = NULL,
  fr_name = NULL,
  en_name = NULL,
  rank = NULL,
  territories = NULL,
  domain = NULL,
  habitats = NULL,
  vernacular = NULL,
  version = NULL,
  page = 1,
  size = 5000
)
```

**Arguments**

<b>id</b>	NULL or integer(1+) [default = NULL] One or more integer giving the id(s) of taxa in TAXREF (also called cdNom)
<b>sciname</b>	NULL or character(1+) [default = NULL] One or more scientific name of searched taxa
<b>fr_name</b>	NULL or character(1+) [default = NULL] One or more common name(s) in French of searched taxa
<b>en_name</b>	NULL or character(1+) [default = NULL] One or more common name(s) in English of searched taxa
<b>rank</b>	NULL or character(1+) [default = NULL] One of more rank of the searched taxa (see <a href="#">rt_taxo_ranks()</a> for the list of all ranks available): <ul style="list-style-type: none"> <li>• Dumm= Domain</li> <li>• KD= Kingdom</li> <li>• PH= Phylum</li> <li>• CL= Class</li> <li>• OR= Order</li> <li>• FM= Family</li> <li>• SBFM= Subfamily</li> <li>• TR= Tribe</li> <li>• GN= Genus</li> <li>• AGES= Aggregate</li> <li>• ES= Species</li> <li>• SSES= Subspecies</li> <li>• NAT= Natio</li> <li>• VAR= Variety</li> <li>• SVAR= Sub-variety</li> </ul>

	<ul style="list-style-type: none"> <li>• F0= Form</li> <li>• SSF0= Sub-form</li> <li>• RACE= Race</li> <li>• CAR= Cultivar</li> <li>• AB= Abberatio</li> </ul>
territories	<p>NULL or character(1+) [default = NULL]  One or more territories where species is present:</p> <ul style="list-style-type: none"> <li>• "fr" = Mainland France</li> <li>• "gf" = French Guiana</li> <li>• "gua" = Guadeloupe</li> <li>• "mar" = Martinique</li> <li>• "sm" = Saint Martin</li> <li>• "sb" = Saint Barthélemy</li> <li>• "spm" = Saint Pierre and Miquelon</li> <li>• "epa" = Scattered Islands in the Indian Ocean</li> <li>• "may" = Mayotte</li> <li>• "reu" = Réunion</li> <li>• "sa" = French Southern and Antarctic Lands</li> <li>• "ta" = Adélie Land</li> <li>• "nc" = New Caledonia</li> <li>• "wf" = Wallis and Futuna</li> <li>• "pf" = French Polynesia</li> <li>• "cli" = Clipperton Island</li> </ul>
domain	<p>NULL or character(1) [default = NULL]  The domain where the species is found:</p> <ul style="list-style-type: none"> <li>• "marin" = Marine</li> <li>• "continental" = Continental</li> </ul>
habitats	<p>NULL or integer(1+) [default = NULL]  One or more habitats where the species is found (see <a href="#">rt_habitats()</a> for a list of all habitats):</p> <ul style="list-style-type: none"> <li>• 1 = Marine</li> <li>• 2 = Freshwater</li> <li>• 3 = Terrestrial</li> <li>• 4 = Marine and freshwater</li> <li>• 5 = Marine and Terrestrial</li> <li>• 6 = Brackish water</li> <li>• 7 = Continental (terrestrial and/or freshwater)</li> <li>• 8 = Continental (terrestrial and freshwater)</li> </ul>
vernacular	<p>NULL or character(1+) [default = NULL]  Name of a vernacular group (see the list using <a href="#">rt_vernacular_groups()</a>)</p>

<code>version</code>	<code>NULL</code> or <code>character(1)</code> [default = <code>NULL</code> ] A string indicating which version of TAXREF should be queried. Should be of the form "2.0" up to "15.0". You can check the available TAXREF versions with <a href="#">rt_taxref_versions()</a> . If <code>NULL</code> uses the latest version of TAXREF available.
<code>page</code>	<code>integer(1)</code> [default = 1] The page number returned
<code>size</code>	<code>integer(1)</code> [default = 5000] The number of lines returned per page (maximum = 5000)

## See Also

the list of available taxonomic ranks [rt\\_taxo\\_ranks\(\)](#), the list of available habitats [rt\\_habitats\(\)](#), the list of vernacular groups [rt\\_taxref\\_versions\(\)](#), and the list of TAXREF versions [rt\\_taxref\\_versions\(\)](#).

## Examples

```
## Not run: rt_taxa_search(sciname = "Bradypus")
```

**rt\_taxa\_sources**      *Retrieve Sources Associated with a Taxon*

## Description

Retrieve Sources Associated with a Taxon

## Usage

```
rt_taxa_sources(id = NULL)
```

## Arguments

<code>id</code>	<code>integer(1)</code> [ <b>required</b> ] an integer giving the id of the taxon in TAXREF (also called cdNom)
-----------------	--

## See Also

[rt\\_source\\_uses\(\)](#) for types of source use

## Examples

```
## Not run: rt_taxa_sources(id = 443800)
```

---

**rt\_taxa\_status***Retrieve Taxon Conservation Status*

---

**Description**

Retrieve Taxon Conservation Status

**Usage**

```
rt_taxa_status(id = NULL, detail = FALSE)
```

**Arguments**

<b>id</b>	<b>integer(1) [required]</b>
	an integer giving the id of the taxon in TAXREF (also called cdNom)
<b>detail</b>	<b>logical(1) [default = FALSE]</b>
	Should the table be a long table detailing all the statuses or a compact table?

**Examples**

```
## Not run:  
rt_taxa_status(id = 443800, detail = FALSE)  
rt_taxa_status(id = 443800, detail = TRUE)  
  
## End(Not run)
```

---

**rt\_taxa\_synonyms***Retrieve Synonyms associated with a Taxon*

---

**Description**

Retrieve Synonyms associated with a Taxon

**Usage**

```
rt_taxa_synonyms(id = NULL)
```

**Arguments**

<b>id</b>	<b>integer(1) [required]</b>
	an integer giving the id of the taxon in TAXREF (also called cdNom)

**Examples**

```
## Not run: rt_taxa_synonyms(id = 107003)
```

---

`rt_taxa_taxrefhistory` *Retrieve history of a taxon through TAXREF versions*

---

### Description

Retrieve history of a taxon through TAXREF versions

### Usage

```
rt_taxa_taxrefhistory(id = NULL)
```

### Arguments

<code>id</code>	<code>integer(1) [required]</code>
	an integer giving the id of the taxon in TAXREF (also called cdNom)

### Examples

```
## Not run: rt_taxa_taxrefhistory(id = 443800)
```

---

`rt_taxa_vernacular` *Retrieve VernacularNames associated with a Taxon*

---

### Description

Retrieve VernacularNames associated with a Taxon

### Usage

```
rt_taxa_vernacular(id = NULL)
```

### Arguments

<code>id</code>	<code>integer(1) [required]</code>
	an integer giving the id of the taxon in TAXREF (also called cdNom)

### See Also

[rt\\_languages\(\)](#) for a list of languages referenced in TAXREF

### Examples

```
## Not run: rt_taxa_vernacular(id = 443800)
```

---

rt_taxo_ranks	<i>Retrieve taxonomic ranks in TAXREF</i>
---------------	---

---

### Description

If the function is used without arguments returns the entire list of taxonomic ranks used in TAXREF.

### Usage

```
rt_taxo_ranks(taxo_id = NULL)
```

### Arguments

taxo_id	NULL or character(1) [default = NULL] the id of a taxonomic rank in TAXREF.
---------	--

### Examples

```
## Not run: rt_taxo_ranks(taxo_id = "VAR")
```

---

rt_taxref_versions	<i>Retrieve the list of existing TAXREF versions</i>
--------------------	--

---

### Description

Retrieve the list of existing TAXREF versions

### Usage

```
rt_taxref_versions(version_id = NULL)
```

### Arguments

version_id	NULL or integer(1) [default = NULL] the id of a TAXREF version
------------	---

### Examples

```
## Not run: rt_taxref_versions(version_id = 12)
```

---

`rt_vernacular_groups`    *Retrieve the list of vernacular groups in TAXREF*

---

### Description

This function takes no argument.

### Usage

`rt_vernacular_groups()`

# Index

rt\_biogeo\_status, 2  
rt\_external\_db, 3  
rt\_external\_db(), 17  
rt\_habitats, 3  
rt\_habitats(), 21, 22  
rt\_interactions\_find\_type, 4  
rt\_interactions\_opgroup, 5  
rt\_interactions\_search, 5  
rt\_interactions\_type, 6  
rt\_interactions\_type(), 4, 6  
rt\_languages, 7  
rt\_languages(), 24  
rt\_operational\_groups, 7  
rt\_operational\_groups(), 5, 6, 11, 12  
rt\_source\_uses, 9  
rt\_source\_uses(), 22  
rt\_sources\_find\_term, 8  
rt\_sources\_id, 8  
rt\_sources\_match, 9  
rt\_status\_find\_type, 10  
rt\_status\_opgroup, 11  
rt\_status\_search, 11  
rt\_status\_type, 13  
rt\_status\_type(), 10  
rt\_taxa\_autocomplete, 13  
rt\_taxa\_children, 15  
rt\_taxa\_externalids, 15  
rt\_taxa\_externalids(), 17  
rt\_taxa\_factsheet, 16  
rt\_taxa\_from\_externalids, 16  
rt\_taxa\_fuzzymatch, 17  
rt\_taxa\_id, 17  
rt\_taxa\_interactions, 18  
rt\_taxa\_media, 18  
rt\_taxa\_parent, 19  
rt\_taxa\_search, 19  
rt\_taxa\_sources, 22  
rt\_taxa\_status, 23  
rt\_taxa\_synonyms, 23  
rt\_taxa\_taxrefhistory, 24  
rt\_taxa\_vernacular, 24  
rt\_taxo\_ranks, 25  
rt\_taxo\_ranks(), 14, 15, 20, 22  
rt\_taxref\_versions, 25  
rt\_taxref\_versions(), 22  
rt\_vernacular\_groups, 26  
rt\_vernacular\_groups(), 21