

Package: SymbiotaR2 (via r-universe)

October 28, 2024

Type Package

Title Downloading Data from Symbiota2 Portals into R

Version 0.0-1

Maintainer Austin Koontz <austinkoontz11@gmail.com>

Description Download data from Symbiota2 portals using Symbiota's API. Covers the Checklists, Collections, Crowdsourcing, Exsiccata, Glossary, ImageProcessor, Key, Media, Occurrence, Reference, Taxa, Traits, and UserRoles API families. Each Symbiota2 portal owner can load their own plugins (and modified code), and so this package may not cover every possible API endpoint from a given Symbiota2 instance.

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URL <https://github.com/pearselab/SymbiotaR2>

BugReports <https://github.com/pearselab/SymbiotaR2/issues>

VignetteBuilder knitr

LazyLoad yes

Suggests knitr (>= 1.23), testthat (>= 2.0.0), vcr

Imports httr (>= 1.0.0), rjson (>= 0.2.20), fs (>= 1.2.6), curl (>= 3.3)

Encoding UTF-8

RoxygenNote 7.1.0

Repository <https://ropensci.r-universe.dev>

RemoteUrl <https://github.com/ropensci/SymbiotaR2>

RemoteRef master

RemoteSha f0231b51066d15e7d6a64baf5c63eb37236a23bd

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Checklists	<i>Retrieves Checklists resources from the Symbiota2 server</i>
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Description

Functions that retrieve Checklist resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

ChecklistProjects(id, page, url = NULL)

Coordinates(id, page, url = NULL)

TaxaLink(id, page, url = NULL)

Checklists(id, page, url = NULL)

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using `id`, the specific resource specified; if using `page`, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:
# Pulling a Coordinates resource (id = 1), from a (nonexistent) dummy portal
object <- Coordinates(id = 1, url = "http://dummy-portal.com/api/")

## End(Not run)
```

Collection

Retrieves Collection resources from the Symbiota2 server

Description

Functions that retrieve Collection resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
Categories(id, page, url = NULL)
```

```
Institutions(id, page, url = NULL)
```

```
Stats(id, page, url = NULL)
```

```
Collections(id, page, url = NULL)
```

Arguments

<code>id</code>	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
<code>page</code>	numeric value referring to the page of resources to pull. If neither an <code>id</code> or a <code>page</code> parameter is provided, function will pull the first page of resources (i.e. <code>page=1</code>)
<code>url</code>	URL string of the Symbiota2 portal to be connected to. A trailing <code>/</code> will be appended, if it is not given.

Value

If using `id`, the specific resource specified; if using `page`, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:
# Pulling in a page of Institutions, from a (nonexistent) dummy portal
ints <- Institutions(page = 3, url = "http://dummy-portal.com/api/")

## End(Not run)
```

Crowdsourcing

Retrieves Crowdsourcing resources from the Symbiota2 server

Description

Functions that retrieve Crowdsourcing resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
Central(id, page, url = NULL)
```

```
Queue(id, page, url = NULL)
```

Arguments

<code>id</code>	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
<code>page</code>	numeric value referring to the page of resources to pull. If neither an <code>id</code> or a <code>page</code> parameter is provided, function will pull the first page of resources (i.e. <code>page=1</code>)
<code>url</code>	URL string of the Symbiota2 portal to be connected to. A trailing <code>/</code> will be appended, if it is not given.

Value

If using `id`, the specific resource specified; if using `page`, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:  
# Acquiring a Queue resource (id = 2), from a (nonexistent) dummy portal  
object <- Queue(id = 2, url = "http://dummy-portal.com/api/")  
  
## End(Not run)
```

Exsiccati

Retrieves Exsiccati resources from the Symbiota2 server

Description

Functions that retrieve Exsiccati resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
Numbers(id, page, url = NULL)
```

```
Titles(id, page, url = NULL)
```

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:
# Acquiring a Titles resource (id = 3), from a (nonexistent) dummy portal
object <- Titles(id = 3, url = "http://dummy-portal.com/api/")

## End(Not run)
```

Glossary

Retrieves Glossary resources from the Symbiota2 server

Description

Functions that retrieve Glossary resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
Glossary(id, page, url = NULL)
```

```
TermLink(id, page, url = NULL)
```

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:  
# Acquiring a page of Glossary resources, from a (nonexistent) dummy portal  
glossPage <- Glossary(page = 1, url = "http://dummy-portal.com/api/")  
  
## End(Not run)
```

ImageProcessor

Retrieves ImageProcessor resources from the Symbiota2 server

Description

Functions that retrieve ImageProcessor resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
Projects(id, page, url = NULL)
```

```
RawLabels(id, page, url = NULL)
```

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:
# Acquiring a RawLabels resource (id = 1), from a (nonexistent) dummy portal
object <- RawLabels(id = 1, url = "http://dummy-portal.com/api/")

## End(Not run)
```

Key	<i>Retrieves Key resources from the Symbiota2 server</i>
-----	--

Description

Functions that retrieve Key resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
CharacterHeading(id, page, url = NULL)

Characters(id, page, url = NULL)

CharacterStateImages(id, page, url = NULL)

CharacterStates(id, page, url = NULL)

DescriptionDeletions(id, page, url = NULL)
```

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:
# Pulling a Characters resource (id = 3), from a (nonexistent) dummy portal
object <- Characters(id = 3, url = "http://dummy-portal.com/api/")

## End(Not run)
```

Media

Retrieves Media resources from the Symbiota2 server

Description

Functions that retrieve Media resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
TagKey(id, page, url = NULL)
```

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:
# Pulling a page of TagKey resources, from a (nonexistent) dummy portal
object <- TagKey(page = 1, url = "http://dummy-portal.com/api/")

## End(Not run)
```

Miscellaneous

Retrieves Miscellaneous resources from the Symbiota2 server

Description

Functions that retrieve Miscellaneous resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
Configurations(id, page, url = NULL)
```

```
LookupLanguages(id, page, url = NULL)
```

```
SchemaVersion(id, page, url = NULL)
```

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:  
# Pulling a Configurations resource (id = 4), from a (nonexistent) dummy portal  
object <- Configurations(id = 4, url = "http://dummy-portal.com/api/")  
  
## End(Not run)
```

Occurrences

Retrieves Occurrence resources from the Symbiota2 server

Description

Functions that retrieve Occurrence resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

AccessStats(id, page, url = NULL)
Determinations(id, page, url = NULL)
Duplicates(id, page, url = NULL)
EditLocks(id, page, url = NULL)
Edits(id, page, url = NULL)
FullText(id, page, url = NULL)
GuidDeterminations(id, page, url = NULL)
GuidOccurrences(id, page, url = NULL)
LookupChronostratigraphy(id, page, url = NULL)
LookupCounties(id, page, url = NULL)
LookupCountries(id, page, url = NULL)
LookupStateProvinces(id, page, url = NULL)
UploadMappings(id, page, url = NULL)
UploadParameters(id, page, url = NULL)
Verification(id, page, url = NULL)
Associations(id, page, url = NULL)
Comments(id, page, url = NULL)
DatasetLink(id, page, url = NULL)
Datasets(id, page, url = NULL)

```
Exchange(id, page, url = NULL)
```

```
Loans(id, page, url = NULL)
```

```
Occurrences(id, page, url = NULL)
```

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:
# Pulling a page of Occurrences, from a (nonexistent) dummy portal
entries <- Occurrence(page = 6, url = "http://dummy-portal.com/api/")

## End(Not run)
```

Reference

Retrieves Reference resources from the Symbiota2 server

Description

Functions that retrieve Reference resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
LookupReferenceTypes(id, page, url = NULL)
```

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:  
# Pulling a LookupReferenceType resource (id = 1),  
from a (nonexistent) dummy portal  
object <- LookupReferenceType(id = 1, url = "http://dummy-portal.com/api/")  
  
## End(Not run)
```

SymbiotaR2

Downloading data from Symbiota2 portals into R

Description

This package allows users to access and download from Symbiota2, a content management system for biodiversity data.

About

Symbiota2 is the improved, refactored version of Symbiota, an open source content management system for biological specimen data. SymbiotaR2 allows users to access the data available at Symbiota2 portals. By specifying the URL of the relevant portal, and the resource to be downloaded, users can use SymbiotaR2 to deliver biological specimen-data in an R format.

Code Structure

Package functions are organized by API family, which generally group the functions by the type of resource they pull from the portal. Each function can either return an individual resources (through specifying the ‘id’ argument) or a collection of resources (through specifying the ‘page’ argument). After providing either the ‘id’ or the ‘page’ of resources, and the URL of the relevant portal, SymbiotaR2 will return an R object (for ‘id’, usually a list; for ‘page’, usually a data.frame).

Portal Specification

All SymbiotaR2 commands require a URL that directs to the Symbiota2 portal to download data from. Users need to make sure they are granted access to a Symbiota2 portal before trying to download data from it.

The address of a Symbiota2 portal is provided as the ‘url’ string argument to each function. To specify a default URL, use the ‘SymbiotaR2_setup’ function, which will be the default url to your .Rprofile.

This package only allows users to access data from existing Symbiota2 portals; to create a new Symbiota2 portal, see the documentation at <https://symbiota2.github.io/Symbiota2/setup/installation.html>

References

<https://symbiota.org/docs/>

Gries, C., Gilbert, E. E., & Franz, N. M. (2014). Symbiota - A virtual platform for creating voucher-based biodiversity information communities. *Biodiversity Data Journal*, 2, e1114.

Examples

```
## Not run:
myURL <- "http://ImaginarySymbiota2Portal.com/api"
myTaxa <- Taxa(id = 12, url = myURL)
str(myTaxa)

myOccurrences <- Occurrence(page = 2, url = myURL)
length(myOccurrences)

## End(Not run)
```

SymbiotaR2_setup

Set default URL for Symbiota2 portal download

Description

Sets the SymbiotaR2_url option for you, optionally, by appending it to your .Rprofile. Checks whether you’ve specified a valid URL that can be reached, and attempts to pull a resource from the API, to confirm that the URL does specify a Symbiota2 portal.

Usage

```
SymbiotaR2_setup(url, append = FALSE, verbose = TRUE)
```

Arguments

url	URL of Symbiota2 portal (a trailing / will be appended, if it is not given)
append	logical of whether to attempt to append this to your .Rprofile file, making this your default every time you start up R
verbose	logical of whether or not to display output

Value

Invisibly, the URL that has been stored

Author(s)

Will Pearse

Examples

```
## Not run:
# An example (that doesn't work because it's not a real portal)
SymbiotaR2_setup("http://nonexistent-portal.com/api/")
# Trying to save a non-existence portal
SymbiotaR2_setup("http://nonexistent-portal.com/api/", TRUE)

## End(Not run)
```

Taxa

Retrieves Taxa resources from the Symbiota2 server

Description

Functions that retrieve Taxa resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

Taxa(id, page, url = NULL)

Authorities(id, page, url = NULL)

DescriptionBlock(id, page, url = NULL)

Synonymy(id, page, url = NULL)

Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:  
# Pulling a page of Taxa resources, from a (nonexistent) dummy portal  
object <- Taxa(page = 2, url = "http://dummy-portal.com/api/")  
  
## End(Not run)
```

Traits

Retrieves Traits resources from the Symbiota2 server

Description

Functions that retrieve Traits resources from the server previously connected to. Each function either retrieves an individual resource or a page of resources, depending on the arguments provided.

Usage

```
Traits(id, page, url = NULL)
```


Arguments

id	id value (usually numeric, but not always) used to refer to the specific resource to pull from the database
page	numeric value referring to the page of resources to pull. If neither an id or a page parameter is provided, function will pull the first page of resources (i.e. page=1)
url	URL string of the Symbiota2 portal to be connected to. A trailing / will be appended, if it is not given.

Value

If using id, the specific resource specified; if using page, the page specified of resources

Note

To specify a default URL to refer to, see [SymbiotaR2_setup\(\)](#)

Author(s)

Austin Koontz

Examples

```
## Not run:  
# Pulling a Traits resource (id = 4), from a (nonexistent) dummy portal  
object <- Traits(id = 4, url = "http://dummy-portal.com/api/")  
  
## End(Not run)
```

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