Package 'hypothesisr'

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Type Package

Title Wrapper for the 'Hypothes.is' Web Annotation Service				
Version 0.1.1				
Description Interact with the application programming interface for the web annotation service 'Hypothes.is' (See http://hypothes.is for more information.) Allows users to download data about public annotations, and create, retrieve, update, and delete their own annotations.				
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hs_create

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Description

Create annotations

Usage

```
hs_create(token, uri, user, permissions = list(read = "group:__world__",
   update = user, delete = user, admin = user), document = NULL,
   target = NULL, tags = NULL, text, custom = NULL)
```

Arguments

token	Character. Your account token, which you can generate at https://hypothes.is/register
uri	Character. The URI to be annotated.
user	Character. Your user account, normally in the format acct:username@hypothes.is
permissions	A named list with read, update, delete, and admin permissions. Defaults to setting global read permissions (group:world) and setting the user string for update, delete, and admin permissions.
document	A list describing the document. CURRENTLY IGNORED.
target	A list describing the highlight position of the annotation. CURRENTLY IGNORED
tags	Character. (optional) Tags to apply to the annotation.
text	Character. Text to put in the body of the annotation. This will be coerced into a character vector of length 1 using paste.
custom	Add arbitrary fields to the JSON object submitted to hypothes.is by means of a named list.

Value

Upon successful creation, returns a 22-character annotation ID. This ID may be retrieved using hs_read.

Source

https://h.readthedocs.io/en/latest/api/#create

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Examples

```
## Not run:
hs_create(token = user_token,
uri = "https://github.com/mdlincoln/hypothesisr",
user = "acct:mdlincoln@hypothes.is", tags = c("testing", "R"),
text = "R made me!")
## End(Not run)
```

hs_delete

Delete an annotation

Description

Delete an annotation

Usage

```
hs_delete(token, id)
```

Arguments

token Character. Your account token, which you can generate at https://hypothes.

is/register

id Character. A hypothes.is annotation id.

Value

TRUE on successful deletion.

Source

```
https://h.readthedocs.io/en/latest/api/#delete
```

hs_read

Retrieve a single annotation by ID

Description

Retrieve a single annotation by ID

Usage

```
hs_read(id)
```

hs_reply

Arguments

id

Character. A hypothes.is annotation id.

Value

A dataframe with the contents of that annotation.

Source

```
https://h.readthedocs.io/en/latest/api/#read
```

Examples

```
hs_read("WFMnSC3FEeaNvLeGkQeJbg")
```

hs_reply

Create a reply to a comment

Description

This is a utility wrapper around hs_create that takes an original comment ID and creates a reply to it by adding the custom references field when constructing the annotation. Normal fields like

Usage

```
hs_reply(token, user, id, text, ...)
```

Arguments

token	Character. Your account token, which you can generate at https://hypothes.is/register
user	$Character.\ Your\ user\ account,\ normally\ in\ the\ format\ acct: \verb"username@hypothes.is"$
id	Character. The annotation ID to reply to.
text	Character. Text to put in the body of the annotation. This will be coerced into a character vector of length 1 using paste.
	Other arguments to pass to hs_create.

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hs_search Search hypothes.is annotations
--

Description

Search hypothes.is annotations.

Usage

```
hs_search(limit = NULL, offset = NULL, sort = "updated", order = "asc",
  uri = NULL, user = NULL, text = NULL, any = NULL, custom = list())
```

Arguments

limit	Integer. The maximum number of annotations to return. (Default: 20)
offset	Integer The minimum number of initial annotations to skip. This is used for pagination. For example if there are 65 annotations matching our search query and we're retrieving up to 30 annotations at a time, then to retrieve the last 5, set offset = 5. (Default = 0)
sort	Character. Specify which field the annotations should be sorted by: consumer, created, id, text, updated, uri, user.
order	Character. Specify which order annotations should be sorted by: asc or desc.
uri	Character. Search for annotations of a particular URI, for example www.example.com. URI searches will also find annotations of equivalent URIs. For example if the HTML document at http://www.example.com/document.html? includes a link rel="canonical" href="http://www.example.com/canonical_document.html"> then annotations of http://www.example.com/canonical_document.html will also be included in the search results. Other forms of document equivalence that are supported include rel="alternate" links, DOIs, PDF file IDs, and more.
user	Character. Search for annotations by a particular user. For example, tim will find all annotations by users named tim at any provider, while tim@hypothes.is will only find annotations by tim on hypothes.is
text	Character. Search for annotations whose body text contains some text, for example: foobar.
any	Character. Search for annotations whose quote, tags, text, uri.parts or user fields match some query text.
custom	A named list of any field in the results returned by hypothes.is as a name, and

Value

A dataframe with annotation data.

Note

If any vectors are passed to these arguments, only the first values will be used.

hs_search_all

Source

https://h.readthedocs.io/en/latest/api/#search

Examples

```
# Search for no more than 5 annotations containing the text "ulysses"
hs_search(text = "ulysses", limit = 5)
# Search with a custom field for tags
hs_search(custom = list(tags = "todo"))
# use the 'uri.parts' field to find annotations on a given domain (exclude
# the TLD, as this will result in all annotations on sites with, e.g., .org,
# as well.)
hs_search(custom = list(uri.parts = "programminghistorian"))
```

hs_search_all

Retreive all annotation search results as a data frame

Description

Takes the same arguments as hs_search and pages through all available results, formatting the output as a data.frame.

Usage

```
hs_search_all(sort = "updated", order = "asc", uri = NULL, user = NULL,
  text = NULL, any = NULL, custom = list(), pagesize = 200,
  progress = interactive())
```

Arguments

sort	Character. Specify which field the annotations should be sorted by: consumer, created, id, text, updated, uri, user.
order	Character. Specify which order annotations should be sorted by: asc or desc.
uri	Character. Search for annotations of a particular URI, for example www.example.com. URI searches will also find annotations of equivalent URIs. For example if the HTML document at http://www.example.com/document.html? includes a link rel="canonical" href="http://www.example.com/canonical_document.html"> then annotations of http://www.example.com/canonical_document.html will also be included in the search results. Other forms of document equivalence that are supported include rel="alternate" links, DOIs, PDF file IDs, and more.
user	Character. Search for annotations by a particular user. For example, tim will find all annotations by users named tim at any provider, while tim@hypothes.is will only find annotations by tim on hypothes.is
text	Character. Search for annotations whose body text contains some text, for example: foobar.

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any	Character. Search for annotations whose quote, tags, text, uri.parts or user fields match some query text.
custom	A named list of any field in the results returned by hypothes.is as a name, and the search text as values.
pagesize	Integer. How many annotations to retrieve per query. Between 1 and 200. (Default: 200)
progress	Boolean. Should a progress bar be displayed during download?

Value

A dataframe with annotation data.

Examples

```
## Not run:
hs_search_all(text = "arxiv")
## End(Not run)
```

hs_update

Update annotations

Description

Update annotations

Usage

```
hs_update(token, id, uri = NULL, user = NULL, permissions = NULL,
document = NULL, target = NULL, tags = NULL, text = NULL,
custom = NULL)
```

Arguments

token	Character. Your account token, which you can generate at https://hypothes.is/register .
id	Character. A hypothes.is annotation id.
uri	Character. The URI to be annotated.
user	Character. Your user account, normally in the format acct:username@hypothes.is
permissions	A named list with read, update, delete, and admin permissions. Defaults to setting global read permissions (group:world) and setting the user string for update, delete, and admin permissions.
document	A list describing the document. CURRENTLY IGNORED.
target	A list describing the highlight position of the annotation. CURRENTLY IGNORED

open_

tags Character. (optional) Tags to apply to the annotation.

text Character. Text to put in the body of the annotation. This will be coerced into a

character vector of length 1 using paste.

custom Add arbitrary fields to the JSON object submitted to hypothes.is by means of a

named list.

Value

TRUE on successful update.

Source

```
https://h.readthedocs.io/en/latest/api/#update
```

Examples

```
## Not run:
hs_update(user_token, "lDf9rC3EEea6ck-G5kLdXA", text = "Now even more annotate-y!")
## End(Not run)
```

hypothesisr

hypothesisr: Wrapper for the Hypothes.is API

Description

Interact with the API for the web annotation service hypothes.is. Allows users to add, search for, and retrieve annotation data.

open_

Functions for opening an annotation in the system browser

Description

Functions for opening an annotation in the system browser

Usage

```
open_context(id)
open_annotation(id)
open_uri(id)
```

open_

Arguments

id Annotation ID.

Functions

• open_context: Open an annotation in context, displaying the original webpage with an annotation overlay

- open_annotation: Open an annotation on the hypothes.is webpage
- open_uri: Open the original webpage to which the annotation links

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