

# Package ‘studyUser’

March 24, 2021

**Title** studyUser

**Version** 1.36.0-1

**API version** v0.1

**Description** This is the API for layer 3 of the Roche pRed project. This swagger page describes the studyUser API endpoints for ODM. These are typically used to find and retrieve study metadata. Before carrying out any API calls you will need an API token. API tokens can be obtained under your profile within the Genestack software. Further instructions can be found [here](https://odm-user-guide.readthedocs.io/en/latest/doc-odm-user-guide/getting-a-genestack-api-token.html). To try out calls in this swagger page: 1. Click the 'Authorize' button below to enter your API token 2. Scroll to the 'Parameters' section for the method you wish to try out and click the 'Try it out' button 3. Enter parameter values that you wish to try 4. Scroll to the bottom of the Parameters section and click the 'Execute' bar that appears The server response will be in the section that follows.

**Depends** R (>= 3.3.2)

**License** Proprietary

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

## R topics documented:

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StudySPoTApi\_get\_study

*Retrieve a single study object by ID (accession)*

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## Description

Retrieve a single study object by ID (accession)

**Usage**

```
StudySPoTApi_get_study(id, returned_metadata_fields, ...)
```

**Arguments**

**id** Unique identifier (accession) of the object.

**returned\_metadata\_fields**  
The parameter defines amount of metadata attributes to return: 1. `template` - return metadata attributes according to applied template, if object doesn't have applied template default template will be used. This is the default for User endpoints. 2. `all` - return all metadata attributes with values and null attributes, if they are present in the applied template. This is the default for Curator endpoints.

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```
StudySPoTApi_get_study_by_version
```

*Retrieve a single study object by ID (accession)*

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**Description**

Retrieve a single study object by ID (accession)

**Usage**

```
StudySPoTApi_get_study_by_version(id, version, returned_metadata_fields, ...)
```

**Arguments**

**id** Unique identifier (accession) of the object.

**version** Unique version of the object.

**returned\_metadata\_fields**  
The parameter defines amount of metadata attributes to return: 1. `template` - return metadata attributes according to applied template, if object doesn't have applied template default template will be used. This is the default for User endpoints. 2. `all` - return all metadata attributes with values and null attributes, if they are present in the applied template. This is the default for Curator endpoints.

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StudySPoTApi_get_study_versions
<i>Retrieve a list of object versions by ID</i>

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**Description**

Retrieve a list of object versions by ID

**Usage**

StudySPoTApi\_get\_study\_versions(id, ...)

**Arguments**

id	Unique identifier (accession) of the object.
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StudySPoTApi_search_studies
<i>List or search for study metadata objects</i>

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**Description**

List or search for study metadata objects

**Usage**

```
StudySPoTApi_search_studies(  
  filter,  
  query,  
  search_specific_terms,  
  returned_metadata_fields,  
  sort,  
  page_limit,  
  page_offset,  
  ...  
)
```

**Arguments**

filter	Filter by study metadata (key-value metadata pair(s)). E.g. "Study Source"=ArrayExpress
query	Search for study objects via a full-text query over all study metadata fields. E.g. "RNA-Seq of human dendritic cells"

**search\_specific\_terms**

If the full-text query term is present in an ODM dictionary, enabling this parameter will modify the query to include child terms of the full-text query. For example, the search query "Body fluid" can be expanded to include the term "Blood" (a child term of "Body fluid") so files containing either "Body fluid" or "Blood" in their metadata will be returned in the search results. The parent-child relationship is defined by the key "broaders" in the dictionary. If the full query term is not present in a dictionary then this parameter has no effect.

**returned\_metadata\_fields**

The parameter defines amount of metadata attributes to return: 1. `template` - return metadata attributes according to applied template, if object doesn't have applied template default template will be used. This is the default for User endpoints. 2. `all` - return all metadata attributes with values and null attributes, if they are present in the applied template. This is the default for Curator endpoints.

**sort**

Attribute to sort by, with optional ascending/descending marker, of the form "[+|-]<key\_name>". No marker or "+" indicates ascending sort, and "-" indicates descending sort. \*Default:\* sort by ID in ascending order.

**page\_limit**

Maximum number of results to return. This value must be between 0 and 100 (inclusive).

**page\_offset**

Show the page `pageOffset+1` results from the start of the results. E.g. 100 will show a page of results starting from the 101st result. The default value is 0.