

Package ‘REDCapR’

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Title Interaction between R and REDCap

Description Encapsulates functions to streamline calls from R

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URL <http://ouhsc.edu/bbmc/>

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R topics documented:

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REDCapR	<i>REDCapR</i>
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Description

REDCapR

redcap_read

*Read records from a REDCap project.***Description**

This function uses REDCap’s **API** to select and return data.

Usage

```
redcap_read(redcap_uri, token, records = NULL, records_collapsed = NULL,
  fields = NULL, fields_collapsed = NULL,
  export_data_access_groups = FALSE, raw_or_label = "raw", verbose = TRUE,
  cert_location = NULL)
```

Arguments

redcap_uri	The URI of the REDCap project. Required.
token	The user-specific string that serves as the password for a project. Required.
records	An array, where each element corresponds to the ID of a desired record. Optional.
records_collapsed	A single string, where the desired ID values are separated by commas. Optional.
fields	An array, where each element corresponds a desired project field. Optional.
fields_collapsed	A single string, where the desired field names are separated by commas. Optional.
export_data_access_groups	A boolean value that specifies whether or not to export the “redcap_data_access_group” field when data access groups are utilized in the project. Default is FALSE. See the details below.
raw_or_label	A string (either 'raw' or 'label' that specifies whether to export the raw coded values or the labels for the options of multiple choice fields. Default is 'raw'.
verbose	A boolean value indicating if messages should be printed to the R console during the operation. Optional.
cert_location	If present, this string should point to the location of the cert files required for SSL verification. If the value is missing or NULL, the server’s identity will be verified using a recent CA bundle from the cURL website . See the details below. Optional.

Details

I like how **PyCap** creates a ‘project’ object with methods that read and write from REDCap. However this isn’t a style that R clients typically use. I like the logic that it’s associated with a particular REDCap project that shouldn’t change between calls. As a compromise, I think I’ll wrap the uri, token, and cert location into a single S4 object that’s passed to these methods. It will make these calls take less space.

The ‘REDCapR’ package includes a recent version of the **Bundle of CA Root Certificates** from the official **cURL site**. This version is used by default, unless the ‘cert_location’ parameter is given another location.

If you do not pass in this `export_data_access_groups` value, it will default to `FALSE`. The following is from the API help page for version 5.2.3: This flag is only viable if the user whose token is being used to make the API request is **not** in a data access group. If the user is in a group, then this flag will revert to its default value.

Value

Currently, a list is returned with the following elements,

1. `data`: an R data frame of the desired records and columns.
2. `raw_csv`: the text of comma separated values returned by REDCap through `RCurl`.
3. `records_collapsed`: the desired records IDs, collapsed into a single string, separated by commas.
4. `fields_collapsed`: the desired field names, collapsed into a single string, separated by commas.
5. `elapsed_seconds`: the duration of the function.
6. `status_message`: a boolean value indicating if the operation was apparently successful.

Author(s)

Will Beasley

References

The official documentation can be found on the ‘API Examples’ page on the REDCap wiki (<https://iwg.devguard.com/trac/redcap/wiki/ApiExamples>). A user account is required.

The official [cURL site](#) discusses the process of using SSL to verify the server being connected to.

Examples

```
## Not run:
library(REDCapR) #Load the package into the current R session.
uri <- "https://bbmc.ouhsc.edu/redcap/api/"
token <- "9A81268476645C4E5F03428B8AC3AA7B"
#Return all records and all variables.
ds_all_rows_all_fields <- redcap_read(redcap_uri=uri, token=token)$data

#Return only records with IDs of 1 and 3
desired_records_v1 <- c(1, 3)
ds_some_rows_v1 <- redcap_read(
  redcap_uri=uri,
  token=token,
  records=desired_records_v1
)$data

#Return only the fields recordid, first_name, and age
desired_fields_v1 <- c("recordid", "first_name", "age")
ds_some_fields_v1 <- redcap_read(
  redcap_uri=uri,
  token=token,
  fields=desired_fields_v1
)$data

## End(Not run)
```

redcap_read_batch	<i>Read records from a REDCap project in subsets, and stacks them together before returning a data.frame.</i>
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Description

From an external perspective, this function is similar to [redcap_read](#). The internals differ in that `read_read_batch` retrieves subsets of the data, and then combines them before returning (among other objects) a single `data.frame`. This function can be more appropriate than [redcap_read](#) when returning large datasets that could tie up the server.

Usage

```
redcap_read_batch(batch_size = 100L, interbatch_delay = 0, redcap_uri,
  token, records = NULL, records_collapsed = NULL, fields = NULL,
  fields_collapsed = NULL, export_data_access_groups = FALSE,
  raw_or_label = "raw", verbose = TRUE, cert_location = NULL)
```

Arguments

<code>batch_size</code>	The maximum number of subject records a single batch should contain. The default is 100.
<code>interbatch_delay</code>	The number of seconds the function will wait before requesting a new subset from REDCap. The default is 0.5 seconds.
<code>redcap_uri</code>	The URI of the REDCap project. Required.
<code>token</code>	The user-specific string that serves as the password for a project. Required.
<code>records</code>	An array, where each element corresponds to the ID of a desired record. Optional.
<code>records_collapsed</code>	A single string, where the desired ID values are separated by commas. Optional.
<code>fields</code>	An array, where each element corresponds a desired project field. Optional.
<code>fields_collapsed</code>	A single string, where the desired field names are separated by commas. Optional.
<code>export_data_access_groups</code>	A boolean value that specifies whether or not to export the “ <code>redcap_data_access_group</code> ” field when data access groups are utilized in the project. Default is <code>FALSE</code> . See the details below.
<code>raw_or_label</code>	A string (either <code>'raw'</code> or <code>'label'</code>) that specifies whether to export the raw coded values or the labels for the options of multiple choice fields. Default is <code>'raw'</code> .
<code>verbose</code>	A boolean value indicating if messages should be printed to the R console during the operation. Optional.
<code>cert_location</code>	If present, this string should point to the location of the cert files required for SSL verification. If the value is missing or <code>NULL</code> , the server’s identity will be verified using a recent CA bundle from the cURL website . See the details below. Optional.

Details

Specifically, it internally uses multiple calls to `redcap_read` to select and return data. Initially, only primary key is queried through the REDCap API. The long list is then subsetting into partitions, whose sizes are determined by the `batch_size` parameter. REDCap is then queried for all variables of the subset's subjects. This is repeated for each subset, before returning a unified `data.frame`.

The function allows a delay between calls, which allows the server to attend to other users' requests.

Value

Currently, a list is returned with the following elements,

1. `data`: an R `data.frame` of the desired records and columns.
2. `raw_csv`: the text of comma separated values returned by REDCap through `RCurl`.
3. `records_collapsed`: the desired records IDs, collapsed into a single string, separated by commas.
4. `fields_collapsed`: the desired field names, collapsed into a single string, separated by commas.
5. `elapsed_seconds`: the duration of the function.
6. `status_message`: a boolean value indicating if the operation was apparently successful.

Author(s)

Will Beasley

References

The official documentation can be found on the 'API Examples' page on the REDCap wiki (<https://iwg.devguard.com/trac/redcap/wiki/ApiExamples>). A user account is required.

The official [cURL site](#) discusses the process of using SSL to verify the server being connected to.

Examples

```
## Not run:
library(REDCapR) #Load the package into the current R session.
uri <- "https://bbmc.ouhsc.edu/redcap/api/"
token <- "9A81268476645C4E5F03428B8AC3AA7B"
redcap_read_batch(batch_size=2, redcap_uri=uri, token=token)

## End(Not run)
```

<code>validate_for_write</code>	<i>Inspect a <code>data.frame</code> to anticipate problems before writing to a REDCap project.</i>
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Description

This set of functions inspect a `data.frame` to anticipate problems before writing with REDCap's [API](#).

Usage

```
validate_for_write( df )  
  
validate_no_logical( df )  
  
validate_no_uppercase( df )
```

Arguments

df	The data.frame containing the dataset used to update the REDCap project. Required.
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Details

All functions listed in the Usage section above inspect a specific aspect of the dataset. The `validate_for_read()` function executes all these individual validation checks. It allows the client to check everything with one call.

Value

A data.frame, where each potential violation is a row. The two columns are:

1. `field_name`: The name of the data.frame that might cause problems during the upload.
2. `field_index`: The position of the field. (For example, a value of '1' indicates the first column, while a '3' indicates the third column.)
3. `concern`: A description of the problem potentially caused by the field.
4. `suggestion`: A *potential* solution to the concern.

Author(s)

Will Beasley

Examples

```
df <- data.frame(  
  recordid = 1:4,  
  flag_logical = c(TRUE, TRUE, FALSE, TRUE),  
  flag_Uppercase = c(4, 6, 8, 2)  
)  
validate_for_write(df = df)
```

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