

Test API connection to FAIRDOMHub

A. Blejec andrej.blejec@nib.si / National Institute of Biology / Ljubljana, Slovenia

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Available at RPubs

Introduction

I will try to connect to the FAIRDOM SEEK with the collection of their APIs.

The API description with examples is available here.

My final goal is to enable upload/download from pISA-tree to FAIRDOMHub. This feature will be a part of the R package pISAR that will enable use of pISA structure and metafile information for reproducible statistical analyses.

Package RCurl

My first attempt will be the RCurl package avaialbe at CRAN. Additional description is a part of Omegahat project.

```
library(RCurl)
```

```
## Loading required package: bitops
```

Check existence

First check if the base URL exists

```
(FAIRDOMExists = url.exists("https://www.fairdomhub.org/"))
```

```
## [1] TRUE
```

Get personal information

For this I will use the GET \people\{id} from documentation and my ID {808}.

```
person <- getURL("https://www.fairdomhub.org/people/808")
str(person)
```

```
##  chr "<!doctype html>\n\n<html lang=\"en\">\n<head>\n  <meta http-equiv=\"content-type\" content=\"t
```

It returns the HTML content and might not be the optimal way.

Package httr

The use is described in LockeData blog.

This solution uses packages `httr` and `jsonlite` and is based on `libcurl` (the same is used by `RCurl`). Package `httr` has useful vignettes that were a good guide for the example below.

```
library(httr)
library(jsonlite)
```

Get personal information

```
r <- GET("https://www.fairdomhub.org/people/808",
         add_headers(Accept="application/json"))
names(r)

## [1] "url"          "status_code"   "headers"      "all_headers"  "cookies"
## [6] "content"       "date"          "times"        "request"     "handle"

r

## Response [https://www.fairdomhub.org/people/808]
##   Date: 2018-06-11 22:38
##   Status: 200
##   Content-Type: application/vnd.api+json; charset=utf-8
##   Size: 1.49 kB
##   <BINARY BODY>
```

The response is structured and can be investigated with the helper functions. The response has three important parts: status, headers, and body.

Status

Check status

```
http_status(r)
```

```
## $category
## [1] "Success"
##
## $reason
## [1] "OK"
##
## $message
```

```

## [1] "Success: (200) OK"
or just a status code
r$status_code

## [1] 200

```

Body

Body can be accessed with the function `content`

```

cont <- content(r, "text")
cont

## [1] "{\"data\":{\"id\":\"808\",\"type\":\"people\",\"attributes\":{\"avatar\":/people/808/avatars/
length(cont)

## [1] 1

```

This gives us a text version of the JSON object. We can use built-in parsers to get other forms of data. We can parse the content into the R list object

```

cont <- content(r, "parsed", type="application/json")
cont

## $data
## $data$id
## [1] "808"
##
## $data$type
## [1] "people"
##
## $data$attributes
## $data$attributes$avatar
## [1] "/people/808/avatars/473"
##
## $data$attributes$title
## [1] "Andrej Blejec"
##
## $data$attributes$description
## NULL
##
## $data$attributes$first_name
## [1] "Andrej"
##
## $data$attributes$last_name
## [1] "Blejec"
##
## $data$attributes$web_page
## [1] "http://ablejec.nib.si"
##
## $data$attributes$orcid
## [1] "http://orcid.org/0000-0001-7484-6031"
##
## $data$attributes$mbox_sha1sum
## [1] "e37350ada8b1af0c7fe7a6cb61a6a5d35d20f8bd"

```

```

## 
## $data$attributes$phone
## [1] "+38659232789"
##
## $data$attributes$skype_name
## [1] "ablejec"
##
## $data$attributes$expertise
## $data$attributes$expertise[[1]]
## [1] "Bioinformatics"
##
## $data$attributes$expertise[[2]]
## [1] "Computational Statistics"
##
## $data$attributes$expertise[[3]]
## [1] "Mathematical and statistical modeling"
##
## $data$attributes$expertise[[4]]
## [1] "Programming"
##
## $data$attributes$expertise[[5]]
## [1] "R"
##
## $data$attributes$expertise[[6]]
## [1] "Statistics"
##
## $data$attributes$expertise[[7]]
## [1] "Visualization"
##
## 
## $data$attributes$tools
## $data$attributes$tools[[1]]
## [1] "Computational Systems Biology"
##
## $data$attributes$tools[[2]]
## [1] "Data Management"
##
## $data$attributes$tools[[3]]
## [1] "Data integration"
##
## $data$attributes$tools[[4]]
## [1] "R"
##
## $data$attributes$tools[[5]]
## [1] "Statistical tools"
##
## $data$attributes$tools[[6]]
## [1] "Statistics"
##
## 
## $data$attributes$project_positions
## $data$attributes$project_positions[[1]]
## $data$attributes$project_positions[[1]]$project_id
## [1] "57"

```

```

## 
## $data$attributes$project_positions[[1]]$position_id
## [1] "3"
##
## $data$attributes$project_positions[[1]]$position_name
## [1] "Project Coordinator"
##
##
##
## $data$relationships
## $data$relationships$projects
## $data$relationships$projects$data
## $data$relationships$projects$data[[1]]
## $data$relationships$projects$data[[1]]$id
## [1] "56"
##
## $data$relationships$projects$data[[1]]$type
## [1] "projects"
##
##
## $data$relationships$projects$data[[2]]
## $data$relationships$projects$data[[2]]$id
## [1] "57"
##
## $data$relationships$projects$data[[2]]$type
## [1] "projects"
##
##
##
## $data$relationships$institutions
## $data$relationships$institutions$data
## $data$relationships$institutions$data[[1]]
## $data$relationships$institutions$data[[1]]$id
## [1] "189"
##
## $data$relationships$institutions$data[[1]]$type
## [1] "institutions"
##
##
##
## $data$relationships$investigations
## $data$relationships$investigations$data
## $data$relationships$investigations$data[[1]]
## $data$relationships$investigations$data[[1]]$id
## [1] "190"
##
## $data$relationships$investigations$data[[1]]$type
## [1] "investigations"
##
##
##

```

```

## 
## $data$relationships$studies
## $data$relationships$studies$data
## $data$relationships$studies$data[[1]]
## $data$relationships$studies$data[[1]]$id
## [1] "365"
##
## $data$relationships$studies$data[[1]]$type
## [1] "studies"
##
##
## 
## $data$relationships$assays
## $data$relationships$assays$data
## $data$relationships$assays$data[[1]]
## $data$relationships$assays$data[[1]]$id
## [1] "655"
##
## $data$relationships$assays$data[[1]]$type
## [1] "assays"
##
##
## 
## $data$relationships$data_files
## $data$relationships$data_files$data
## $data$relationships$data_files$data[[1]]
## $data$relationships$data_files$data[[1]]$id
## [1] "2165"
##
## $data$relationships$data_files$data[[1]]$type
## [1] "data_files"
##
##
## 
## $data$relationships$models
## $data$relationships$models$data
## list()
##
##
## 
## $data$relationships$sops
## $data$relationships$sops$data
## list()
##
##
## 
## $data$relationships$publications
## $data$relationships$publications$data
## list()
##
##
## 
## $data$relationships$presentations
## $data$relationships$presentations$data

```

```

## list()
##
##
## $data$relationships$events
## $data$relationships$events$data
## list()
##
##
## $data$relationships$documents
## $data$relationships$documents$data
## list()
##
##
## $data$links
## $data$links$self
## [1] "/people/808"
##
##
## $data$meta
## $data$meta$created
## [1] "2016-11-07T22:17:49.000Z"
##
## $data$meta$modified
## [1] "2017-08-01T08:28:42.000Z"
##
## $data$meta$api_version
## [1] "0.1"
##
## $data$meta$uuid
## [1] "f1427a00-8765-0134-769f-549f350973c0"
##
## $data$meta$base_url
## [1] "https://fairdomhub.org"
##
##
## $jsonapi
## $jsonapi$version
## [1] "1.0"

```

The structure of the parsed response

```

names(cont)

## [1] "data"      "jsonapi"
cont$jsonapi

## $version
## [1] "1.0"
names(cont$data)

## [1] "id"          "type"        "attributes"   "relationships"

```

```

## [5] "links"           "meta"
Response elements
cont$data$id

## [1] "808"

List the response content
d <- cont$data
for (nm in names(d)) {
  cat("\n", nm, "-----\n\n")
  print(d[[nm]])
}

## -----
##   id -----
## 
## [1] "808"
##
##   type -----
## 
## [1] "people"
##
##   attributes -----
## 
## $avatar
## [1] "/people/808/avatars/473"
##
## $title
## [1] "Andrej Blejec"
##
## $description
## NULL
##
## $first_name
## [1] "Andrej"
##
## $last_name
## [1] "Blejec"
##
## $web_page
## [1] "http://ablejec.nib.si"
##
## $orcid
## [1] "http://orcid.org/0000-0001-7484-6031"
##
## $mbox_sha1sum
## [1] "e37350ada8b1af0c7fe7a6cb61a6a5d35d20f8bd"
##
## $phone
## [1] "+38659232789"
##
## $skype_name
## [1] "ablejec"
##

```

```

## $expertise
## $expertise[[1]]
## [1] "Bioinformatics"
##
## $expertise[[2]]
## [1] "Computational Statistics"
##
## $expertise[[3]]
## [1] "Mathematical and statistical modeling"
##
## $expertise[[4]]
## [1] "Programming"
##
## $expertise[[5]]
## [1] "R"
##
## $expertise[[6]]
## [1] "Statistics"
##
## $expertise[[7]]
## [1] "Visualization"
##
##
## $tools
## $tools[[1]]
## [1] "Computational Systems Biology"
##
## $tools[[2]]
## [1] "Data Management"
##
## $tools[[3]]
## [1] "Data integration"
##
## $tools[[4]]
## [1] "R"
##
## $tools[[5]]
## [1] "Statistical tools"
##
## $tools[[6]]
## [1] "Statistics"
##
##
## $project_positions
## $project_positions[[1]]
## $project_positions[[1]]$project_id
## [1] "57"
##
## $project_positions[[1]]$position_id
## [1] "3"
##
## $project_positions[[1]]$position_name
## [1] "Project Coordinator"
##

```

```

##
##
##
##   relationships -----
##
## $projects
## $projects$data
## $projects$data[[1]]
## $projects$data[[1]]$id
## [1] "56"
##
## $projects$data[[1]]$type
## [1] "projects"
##
##
## $projects$data[[2]]
## $projects$data[[2]]$id
## [1] "57"
##
## $projects$data[[2]]$type
## [1] "projects"
##
##
##
## $institutions
## $institutions$data
## $institutions$data[[1]]
## $institutions$data[[1]]$id
## [1] "189"
##
## $institutions$data[[1]]$type
## [1] "institutions"
##
##
##
## $investigations
## $investigations$data
## $investigations$data[[1]]
## $investigations$data[[1]]$id
## [1] "190"
##
## $investigations$data[[1]]$type
## [1] "investigations"
##
##
##
## $studies
## $studies$data
## $studies$data[[1]]
## $studies$data[[1]]$id
## [1] "365"

```

```
##  
## $studies$data[[1]]$type  
## [1] "studies"  
##  
##  
##  
##  
## $assays  
## $assays$data  
## $assays$data[[1]]  
## $assays$data[[1]]$id  
## [1] "655"  
##  
## $assays$data[[1]]$type  
## [1] "assays"  
##  
##  
##  
##  
## $data_files  
## $data_files$data  
## $data_files$data[[1]]  
## $data_files$data[[1]]$id  
## [1] "2165"  
##  
## $data_files$data[[1]]$type  
## [1] "data_files"  
##  
##  
##  
##  
## $models  
## $models$data  
## list()  
##  
##  
## $sops  
## $sops$data  
## list()  
##  
##  
## $publications  
## $publications$data  
## list()  
##  
##  
## $presentations  
## $presentations$data  
## list()  
##  
##  
## $events  
## $events$data  
## list()
```

```

## 
## $documents
## $documents$data
## list()
##
## 
## 
##   links -----
## 
## $self
## [1] "/people/808"
## 
## 
##   meta -----
## 
## $created
## [1] "2016-11-07T22:17:49.000Z"
## 
## $modified
## [1] "2017-08-01T08:28:42.000Z"
## 
## $api_version
## [1] "0.1"
## 
## $uuid
## [1] "f1427a00-8765-0134-769f-549f350973c0"
## 
## $base_url
## [1] "https://fairdomhub.org"

```

Person data are in the *attributes* part

Here is the first name of the person with id 808:

```

cont$data$attributes$first_name

## [1] "Andrej"

```

Conclusion

The rudimentary R read connection with FAIRDOMHub can be established using the package `httr`. From here it should be fairly possible to write the wrapper functions for easier usage.

Wrapper functions for GET and content

```

# this function is equivaalent to a call:
# curl -X GET "https://www.fairdomhub.org/what/id" -H "accept: application/json"
#
myGET <- function(what, id,
                    uri="https://www.fairdomhub.org", ... ){
  if(!missing(what)) uri <- paste0(uri, "/", what)
  if(!missing(id)) uri <- paste0(uri, "/", id)
}

```

```

r <- GET(uri,
          add_headers(Accept="application/json"))
cat("Status code:",r$status_code,"\n")
invisible(r)
}

myData <- function(r,
                     part="attributes",
                     type="application/json", ...)
  invisible(content(r,"parsed",type=type)$data[[part]])
}

```

Use of wrapper functions

Get data for person 808

```
# https://www.fairdomhub.org/people/808
r <- myGET("people","808")
```

```
## Status code: 200
r$status_code
```

```
## [1] 200
```

Get the person 808 last name and tools

```
myData(r)$last_name
```

```
## [1] "Blejec"
```

```
myData(r)$tools
```

```
## [[1]]
## [1] "Computational Systems Biology"
##
## [[2]]
## [1] "Data Management"
##
## [[3]]
## [1] "Data integration"
##
## [[4]]
## [1] "R"
##
## [[5]]
## [1] "Statistical tools"
##
## [[6]]
## [1] "Statistics"
```

Get person 808 institution id

```
myData(r,"relationships")$institutions
```

```
## $data
## $data[[1]]
```

```

## $data[[1]]$id
## [1] "189"
##
## $data[[1]]$type
## [1] "institutions"
instID <- myData(r, "relationships")$institutions$data[[1]]$id
instID

## [1] "189"

Get institution `instID` name. The details are in the institutions part.

# https://www.fairdomhub.org/institutions/189
rinst <- myGET("institutions",instID)

## Status code: 200
myData(rinst)$title

## [1] "National Institute of Biology"

Full institution description

catln <- function(...) cat(..., "\n")
d <- sapply(myData(rinst),catln)

## /institutions/189/avatars/568
## National Institute of Biology
## SI
##
## Ljubljana
## Vecna pot 111
## SI-1000 Ljubljana
## Slovenia
## http://www.nib.si

```