ctrialsgov: Query Data from U.S. National Library of Medicine's Clinical Trials Database

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Abstract Tools to create and query database from the U.S. National Library of Medicine's Clinical Trials database https://clinicaltrials.gov/. Functions provide access a variety of techniques for searching the data using range queries, categorical filtering, and by searching for full-text keywords. Minimal graphical tools are also provided for interactively exploring the constructed data.

Introduction

Creating Local Data

Before querying the ClinicalTrials.gov data, we need to load a pre-processed version of the data into R. There are three ways to do this. If you have installed a copy of the data set locally into PostGRES, the data can be created from scratch with the following block of code (it will take a couple of minutes to finish):

```
> library(DBI)
> library(RPostgreSQL)

> drv <- dbDriver('PostgreSQL')
> con <- DBI::dbConnect(drv, dbname="aact")
> ctgov_create_data(con)
```

Alternatively, we can download a static version of the data from GitHub and load this into R without needing the setup a local version of the database. This will be cached locally so that it can be re-loaded without downloading each time. To download and load this data, use the following:

```
> ctgov_load_cache()
```

Finally, we can load a small sample dataset (2% of the total) that is included with the package itself using the following:

```
> ctgov_load_sample()
```

This is the version of the data that is used in most of the tests, examples, and in this vignette.

Running Data Queries

The primary function for querying the dataset is called ctgov_query. It can be called after using any of the functions in the previous section. Here are a few examples of how the function works. We will see a few examples here; see the help pages for a complete list of options.

There are a number of fields in the data that use exact matches of categories. Here, for example, we find the interventional studies:

```
> ctgov_query(study_type = "Interventional")
# A tibble: 2,403 Œ 32
  enrollment brief_title official_title
                                      <int> <chr>
                                                           <chr>
1 NCT04999163 2021-12-31 N/A
                                            50 Aortix Ther Aortix Therap
2 NCT05002153 2021-11-30 N/A
                                          300 The Role of The Role of M
3 NCT04472702 2021-11-30 N/A
                                           45 Fluoroscopi Fluoroscopic
4 NCT05032157 2021-11-30 Phase 3
                                          450 A Phase 3 S A Multicenter
5 NCT04471142 2021-11-08 N/A
                                          270 Effectivene Effectiveness
6 NCT04772651 2021-11-01 N/A
                                           108 Mediterrane Mediterranean
7 NCT04390451 2021-11-01 Phase 1
                                            54 Initial Tes Initial Testi
```

```
8 NCT04696861 2021-11-01 N/A
                                                     60 Telehealth Telehealth to
9 NCT03954431 2021-10-31 Phase 1/Phase 2
                                                    100 High-Resolu Study of High
10 NCT04273022 2021-10-31 N/A
                                                     20 Effect of E The Effect of
# with 2,393 more rows, and 26 more variables:
   primary_completion_date <date>, study_type <chr>, rec_status <chr>,
   completion_date <date>, last_update <date>, description <chr>,
   eudract_num <chr>, other_id <chr>, allocation <chr>,
   intervention_model <chr>, observational_model <chr>, primary_purpose <chr>,
   time_perspective <chr>, masking_description <chr>,
   intervention_model_description <chr>, sampling_method <chr>,
   Or, all of the interventional studies that have a primary industry sponsor:
> ctgov_query(study_type = "Interventional", sponsor_type = "Industry")
# A tibble: 640 Œ 32
  enrollment brief_title official_title
                                           <int> <chr>
  <chr>
               <date> <chr>
                                                                      <chr>
                                                 50 Aortix Ther Aortix Therap
1 NCT04999163 2021-12-31 N/A 50 AOFTIX THEI AULTIA THEI UP
2 NCT05032157 2021-11-30 Phase 3 450 A Phase 3 S A Multicenter
3 NCT05029856 2021-10-04 Phase 1/Phase 2 240 Evaluation A Randomized,
1 NCT04999163 2021-12-31 N/A
4 NCT04963179 2021-09-30 N/A
                                                   154 PREvention PREvention of
                                              68 A Study to A Randomized,
100 Study of Al Treatment of
60 Study of HE Phase Ic Clin
64 Renal Funct An Open-label
5 NCT04875975 2021-09-30 Phase 2
6 NCT04909879 2021-09-30 Phase 2
7 NCT04925674 2021-09-29 Phase 1
8 NCT04935177 2021-09-17 Phase 3
9 NCT04956744 2021-08-31 Phase 1
                                                    30 A Study to A Phase 1, Do
10 NCT04920253 2021-08-31 N/A
                                                   180 Real World Real World Ev
# with 630 more rows, and 26 more variables: primary_completion_date <date>,
   study_type <chr>, rec_status <chr>, completion_date <date>,
   last_update <date>, description <chr>, eudract_num <chr>, other_id <chr>,
   allocation <chr>, intervention_model <chr>, observational_model <chr>,
   primary_purpose <chr>, time_perspective <chr>, masking_description <chr>,
    intervention_model_description <chr>, sampling_method <chr>, gender <chr>,
   minimum_age <dbl>, maximum_age <dbl>, population <chr>, criteria <chr>,
```

A few fields have continuous values that can be searched by giving a vector with two values. The results return any values that fall between the lower bound (first value) and the upper bound (second value). Here, we find the studies that have between 40 and 42 patients enrolled in them:

```
> ctgov_query(enrollment_range = c(40, 42))
# A tibble: 125 Œ 32
  nct id
              start_date phase
                                          enrollment brief_title official_title
  <chr>
              <date> <chr>
                                           <int> <chr>
                                                                 <chr>
                                                 42 A Proof of A Proof of Co
1 NCT04188119 2021-09-30 Phase 2
2 NCT04992975 2021-08-31 NA
                                                 40 Brain Iron Brain Iron To
3 NCT05001854 2021-08-31 Phase 2/Phase 3 40 Hemodynamic Evaluation of 4 NCT04749355 2021-08-14 Phase 2 40 Phase 2, Op A Phase 2, Op
5 NCT04648319 2021-04-15 Phase 2
                                               40 A Study of A Pilot Study
6 NCT04744779 2021-03-31 N/A
                                                40 Office Base Effectiveness
7 NCT04841174 2021-03-30 N/A
                                                40 The Effect The Effect of
                                                40 Clinical Ef Effects of Bi
8 NCT04808180 2021-03-25 N/A
9 NCT04746105 2021-02-24 Phase 1
                                                40 A Clinical A Study to Ev
10 NCT04355780 2021-01-08 NA
                                                 40 Immunologic Immunologic F
# with 115 more rows, and 26 more variables: primary_completion_date <date>,
   study_type <chr>, rec_status <chr>, completion_date <date>,
   last_update <date>, description <chr>, eudract_num <chr>, other_id <chr>,
   allocation <chr>, intervention_model <chr>, observational_model <chr>,
   primary_purpose <chr>, time_perspective <chr>, masking_description <chr>,
   intervention_model_description <chr>, sampling_method <chr>, gender <chr>,
   minimum_age <dbl>, maximum_age <dbl>, population <chr>, criteria <chr>,
```

Setting one end of the range to missing avoids searching for that end of the range. For example, the following finds any studies with 1000 or more patients.

```
> ctgov_query(enrollment_range = c(1000, NA))
# A tibble: 204 @ 32
```

```
nct_id
                     start_date phase enrollment brief_title
                                                                                       official_title
    <chr>
                    <date>
                                <chr>
                                                     <int> <chr>
                                                                                       <chr>
                                                        1500 Impact of the M Impact of the Mod
 1 NCT05033782 2021-12-01 NA
2 NCT05033548 2021-10-10 NA 4000 Technology Enable
3 NCT04982614 2021-10-01 Phase 4 1400 HPV Vaccination A Multi-site, Ope
4 NCT05033678 2021-08-16 NA 8000 Implementation Teledermoscopy an
5 NCT04917185 2021-06-30 N/A 1000 EA for PAAS: A Electro-acupunctu
6 NCT04839757 2021-06-03 NA 1400 Dengue Vaccine Preparing for the
7 NCT04889924 2021-06-01 N/A 1666 ALND vs RDT in Axillary Lymph No
8 NCT04472845 2021-03-30 N/A 1018 HYPofractionated
9 NCT04735744 2021-02-15 NA 1315 Evaluation of A Evaluation of All
 2 NCT05033548 2021-10-10 NA
                                                      4000 Technology Enab Technology Enable
 9 NCT04735744 2021-02-15 NA
                                                      1315 Evaluation of A Evaluation of All
10 NCT04626973 2021-01-15 N/A
                                                       3048 Effects of Ezet Effects of Ezetim
# with 194 more rows, and 26 more variables: primary_completion_date <date>,
    study_type <chr>, rec_status <chr>, completion_date <date>,
     last_update <date>, description <chr>, eudract_num <chr>, other_id <chr>,
     allocation <chr>, intervention_model <chr>, observational_model <chr>,
     primary_purpose <chr>, time_perspective <chr>, masking_description <chr>,
    intervention_model_description <chr>, sampling_method <chr>, gender <chr>,
     minimum_age <dbl>, maximum_age <dbl>, population <chr>, criteria <chr>,
```

Similarly, we can give a range of dates. These are given in the form of strings as "YYYY-MM-DD":

```
> ctgov_query(date_range = c("2020-01-01", "2020-02-01"))
# A tibble: 34 Œ 32
                                    enrollment brief_title official_title
  nct_id
               start_date phase
                                         <int> <chr>
   <chr>
               <date>
                          <chr>
                                                                   <chr>
1 NCT04224597 2020-02-01 NA
                                                   48 Evaluation Evaluation of
                                                200 Choroidal C OCTA to Quant
                                          200 Choroidal C OCTA to Quant
25000 Killing Pai Killing Pain
2 NCT04255524 2020-02-01 N/A
3 NCT04336605 2020-02-01 NA
                                               105 The Approac A Clinical Ra
4 NCT04218669 2020-02-01 N/A
                                                  59 Cost-Effect Cost-Effectiv
5 NCT04409613 2020-02-01 N/A
6 NCT04424576 2020-01-31 NA
                                                 60 Ovarian Mor Trajectory of
6 NCT04424576 2020-01-31 NA
7 NCT04115397 2020-01-31 Phase 4
80 Bisphosphon Towards Effic
8 NCT04497064 2020-01-30 NA
585 Breakfast K Breakfast Kno
9 NCT03892785 2020-01-27 Phase 3
                                                 200 MEthotrexat MEthotrexate
10 NCT03710122 2020-01-23 Phase 2/Phase 3 102 Vancomycin A Prospective
# with 24 more rows, and 26 more variables: primary_completion_date <date>,
   study_type <chr>, rec_status <chr>, completion_date <date>,
   last_update <date>, description <chr>, eudract_num <chr>, other_id <chr>,
   allocation <chr>, intervention_model <chr>, observational_model <chr>,
   primary_purpose <chr>, time_perspective <chr>, masking_description <chr>,
   intervention_model_description <chr>, sampling_method <chr>, gender <chr>,
   minimum_age <dbl>, maximum_age <dbl>, population <chr>, criteria <chr>,
```

Finally, we can also search free text fields using keywords. The following for example finds and study that includes the phrase "lung cancer" (ignoring case) in the description field:

```
> ctgov_query(description_kw = "lung cancer")
# A tibble: 59 Œ 32
   nct id
                start_date phase enrollment brief_title
                                                                      official_title
   <chr>
                 <date> <chr> <int> <chr>
                                                                      <chr>
                                           15 To Evaluate the An Open-Labeled,
1 NCT04814056 2021-06-01 Phase 4
1 NCT04629027 2021-03-03 NA
3 NCT04179305 2020-10-25 N/A
4 NCT04452877 2020-08-19 Phase 2
5 NCT04422392 2020-07-13 Phase 2
6 NCT04120454 2020-03-16 Phase 2
7 12-19 Phase 2
7 12-19 Phase 2
7 Modified Versus Randomized Clinic
2 NCT04629027 2021-03-03 NA
9 NCT04151940 2019-09-26 NA
                                               40 PET/CT Changes An Observational
10 NCT04081688 2019-08-21 Phase 1
                                               15 Atezolizumab an A Phase I Trial o
# with 49 more rows, and 26 more variables: primary_completion_date <date>,
   study_type <chr>, rec_status <chr>, completion_date <date>,
    last_update <date>, description <chr>, eudract_num <chr>, other_id <chr>,
    allocation <chr>, intervention_model <chr>, observational_model <chr>,
```

```
# primary_purpose <chr>, time_perspective <chr>, masking_description <chr>,
# intervention_model_description <chr>, sampling_method <chr>, gender <chr>,
# minimum_age <dbl>, maximum_age <dbl>, population <chr>, criteria <chr>,
```

We can search two terms at once as well, by default it finds things that match at least one of the terms:

```
> ctgov_query(description_kw = c("lung cancer", "colon cancer"))
# A tibble: 59 Œ 32
     official_title
                                                           <int> <chr>
                          <date> <chr>
     <chr>
                                                                                                                 <chr>
                                                                       15 To Evaluate the An Open-Labeled,
 1 NCT04814056 2021-06-01 Phase 4
 2 NCT04629027 2021-03-03 NA
                                                                         80 Evaluation Syst Establishment of
3 NCT04179305 2020-10-25 N/A
4 NCT04452877 2020-08-19 Phase 2
5 NCT04422392 2020-07-13 Phase 2
6 NCT04120454 2020-03-16 Phase 2
7 NCT04332367 2019-12-19 Phase 2
6 NCT043209055 2019-12-01 N/A
6 Modified Versus Randomized Clinic
6 NCT04309055 2019-12-01 N/A
6 Modified Versus Randomized Clinic
6 NCT04309055 2019-12-01 N/A
6 NCT04452877 2020-08-19 Phase 2
7 NCT04309055 2019-12-01 N/A
6 NCT04452877 2020-08-19 Phase 2
7 NCT04309055 2019-12-01 N/A
6 NCT04452877 2020-08-19 Phase 2
7 NCT04309055 2019-12-01 N/A
6 NCT04452877 2020-08-19 Phase 2
7 NCT04309055 2019-12-01 N/A
7 NCT04309055 2019-12-01 N/A
7 NCT04309055 2019-12-01 N/A
# with 49 more rows, and 26 more variables: primary_completion_date <date>,
      study_type <chr>, rec_status <chr>, completion_date <date>,
      last_update <date>, description <chr>, eudract_num <chr>, other_id <chr>,
      allocation <chr>, intervention_model <chr>, observational_model <chr>,
       primary_purpose <chr>, time_perspective <chr>, masking_description <chr>,
       intervention_model_description <chr>, sampling_method <chr>, gender <chr>,
      minimum_age <dbl>, maximum_age <dbl>, population <chr>, criteria <chr>,
```

But the 'match_all' flag can be set to search for both terms at the same time (here, that returns no matches):

```
> ctgov_query(description_kw = c("lung cancer", "colon cancer"), match_all = TRUE)
# A tibble: 0 Œ 32
# with 32 variables: nct_id <chr>, start_date <date>, phase <chr>,
# enrollment <int>, brief_title <chr>, official_title <chr>,
primary_completion_date <date>, study_type <chr>, rec_status <chr>,
# completion_date <date>, last_update <date>, description <chr>,
# eudract_num <chr>, other_id <chr>, allocation <chr>,
# intervention_model <chr>, observational_model <chr>, primary_purpose <chr>,
# time_perspective <chr>, masking_description <chr>,
```

Other keyword fields include official_title_kw, source_kw and criteria_kw. Any of the options can be combined as needed.

```
> ctgov_query(
           description_kw = "cancer",
            enrollment_range = c(100, 200),
            date_range = c("2019-01-01", "2020-02-01")
 # A tibble: 5 Œ 32
      nct_id start_date phase enrollment brief_title official_title primary_complet
1 NCT04 2020-01-22 N/A 120 Symptom Ma Improving Sym 2025-10-01 2 NCT04 2020-01-07 Phas 121 Study Eval A Phase 2, Op 2023-07-31 3 NCT04 2020-01-01 NA 100 Extraordin Extraordinary 2022-01-01 4 NCT03 2019-03-12 NA 115 Moderately Moder
      <chr> <date> <chr> <int> <chr> <chr>
                                                                                                                                                                                                          <date>
                                                                                                115 Moderately Moderately Hy 2022-05-31
 5 NCT03 2019-02-20 N/A
                                                                                                 160 The Effect The Effect of 2019-05-31
 # with 25 more variables: study_type <chr>, rec_status <chr>,
          completion_date <date>, last_update <date>, description <chr>,
           eudract_num <chr>, other_id <chr>, allocation <chr>,
          intervention_model <chr>, observational_model <chr>, primary_purpose <chr>,
          time_perspective <chr>, masking_description <chr>,
            intervention_model_description <chr>, sampling_method <chr>, gender <chr>,
            minimum_age <dbl>, maximum_age <dbl>, population <chr>, criteria <chr>,
```

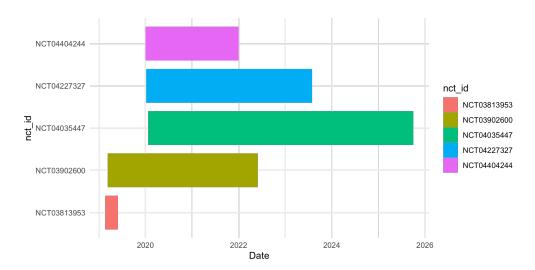


Figure 1: Static Output from the timeline graphic interface.

Finally, we can also pass a current version of the data set to the query function, rather than starting with the full data set. This is useful when you want to combine queries in a more complex way. For example, this is equivalent to the above:

```
> library(dplyr)
> ctgov_query() %>%
   ctgov_query(description_kw = "cancer") %>%
   ctgov_query(enrollment_range = c(100, 200)) %>%
   ctgov_query(date_range = c("2019-01-01", "2020-02-01"))
# A tibble: 5 Œ 32
 nct_id start_date phase enrollment brief_title official_title primary_complet
                <chr> <int> <chr>
  <chr> <date>
                                                <chr>
                                                               <date>
1 NCT04 2020-01-22 N/A
                              120 Symptom Ma Improving Sym 2025-10-01
                            121 Study Eval A Phase 2, Op 2023-07-31
2 NCT04 2020-01-07 Phas
3 NCT04 2020-01-01 NA
                              100 Extraordin Extraordinary 2022-01-01
4 NCT03 2019-03-12 NA
                              115 Moderately Moderately Hy 2022-05-31
5 NCT03 2019-02-20 N/A
                              160 The Effect The Effect of 2019-05-31
 with 25 more variables: study_type <chr>, rec_status <chr>,
   completion_date <date>, last_update <date>, description <chr>,
   eudract_num <chr>, other_id <chr>, allocation <chr>,
   intervention_model <chr>, observational_model <chr>, primary_purpose <chr>,
   time_perspective <chr>, masking_description <chr>,
   intervention_model_description <chr>, sampling_method <chr>, gender <chr>,
   minimum_age <dbl>, maximum_age <dbl>, population <chr>, criteria <chr>,
```

Data Visualization

The package also contains a number of tools for visualizing the output. Here is one example:

```
> ctgov_query(
+    description_kw = "cancer",
+    enrollment_range = c(100, 200),
+    date_range = c("2019-01-01", "2020-02-01")
+ ) %>%
+ ctgov_plot_timeline() +
+ ggplot2::theme_minimal()
```

Text Analysis

Keywords in Context

The function ctgov_kwic highlights all of the occurances of a term within its context (the few words before and after the term occurs). For example, if we want to show the occurances of the term "bladder" in the titles of the interventional trials we can do this:

```
> z <- ctgov_query(study_type = "Interventional")</pre>
> ctgov_kwic("bladder", z[['brief_title']])
ible Local Advanved |Bladder| Cancer
 of Sulforaphane in |Bladder| Cancer Chemoprevent
      Comparison of |Bladder| Filling vs. Non-Fil
tment of Overactive |Bladder|/Urge Incontinence
in the Detection of |Bladder| Cancer in the Surve
A-F Betafood on GallIbladderl and Liver Function.
Non-Muscle Invasive |Bladder| Cancer
iopathic Overactive |Bladder| With Urinary Incont
tion for Overactive |Bladder|
anced or Metastatic |Bladder| Cancer
     Autologous Neo-|Bladder| Construct in Non-Ne
urogenic Overactive |Bladder| and Urge Predominan
AMG 706 on the Gall|bladder| in Advanced Solid T
Men With Overactive |Bladder|.
```

The function also has an option to include a title along with each occurance that is printed alongside each row. Here we will print the NCT id for each trial:

```
> z <- ctgov_query(study_type = "Interventional")</pre>
> ctgov_kwic("bladder", z[['brief_title']], z[['nct_id']])
[NCT04553939] ible Local Advanved |Bladder| Cancer
[NCT03517995] of Sulforaphane in |Bladder| Cancer Chemoprevent
                    Comparison of |Bladder| Filling vs. Non-Fil
[NCT04210479]
[NCT03535857] tment of Overactive |Bladder|/Urge Incontinence
[NCT02560584] in the Detection of |Bladder| Cancer in the Surve
[NCT01981343] A-F Betafood on Gall|bladder| and Liver Function.
[NCT01625260] Non-Muscle Invasive |Bladder| Cancer
[NCT00910845] iopathic Overactive |Bladder| With Urinary Incont
[NCT00912314] tion for Overactive |Bladder|
[NCT00635726] anced or Metastatic |Bladder| Cancer
                  Autologous Neo-|Bladder| Construct in Non-Ne
[NCT00594139]
[NCT00594139] urogenic Overactive |Bladder| and Urge Predominan
[NCT00448786] AMG 706 on the Gall|bladder| in Advanced Solid T
[NCT00282932] Men With Overactive |Bladder|.
```

There are some other options that can be used to change the way that the output is displayed. The default (shown above) prints the results out using the cat function. Other options return the results as a character vector of data frame, which are useful for further post-processing. There is also a flag use_color that prints the term in color rather than with pipes; it looks great in a terminal or RStudio but does not display correctly when knit to HTML.

TF-IDF

We can use a technique called term frequence-inverse document frequency (TF-IDF) to determine the most important words in a collection of of text fields. To implement this in R we will use the ctgov_tfidf function:

```
2
       1 pollution|ms|air|viral|france
3
       2 fmt|diversity|microbiota|weight|gut
4
       3 nerves|landmarks|bony|guidance|knee
5
       4 antihistamines|h1|inadequately|spontaneous|suffering
       5 bandage|seroma|drain|categorical|variables
6
       6 vagal|mediterranean|nerve|diet|depression
8
       7 veterans|peer|whole|steps|structured
       8 suicidal|ideation|telehealth|engagement|counseling
10
       9 bct|ce|breast|structures|cancers
      10 athletes|pathways|exercise|biomarkers|strenuous
11
      11 acetylsalicylic|vessels|artery|affects|acid
12
13
      12 mhealth|90|monitoring|organ|impact
14
      13 cascade|sugar|glucose|sensor|doctors
15
      14 variant|b1351|cov2|sars|b16172
16
      15 scenario|uncertainties|oncological|relating|real
      16 itch|epigenetic|mechanisms|chronic|antagonists
17
18
      17 9vhpv|1526|hiv|living|uninfected
19
      18 dengue|fever|permeability|five|vascular
20
      19 influenza|icu|aspergillosis|eortc|pathogen
21
      20 cannabigerol|cbg|thc|appetite|stimulating
22
      21 antibiotics|decide|how|parent|prescribed
23
      22 counseling|education|behavior|behavioral|his
24
      23 intrauterine|adhesiolysis|leaf|film|named
25
      24 purifiers|cardiopulmonary|indicators|students|air
26
      25 donepezil|french|alzheimers|efficiency|authorities
27
      26 avelumab|checkpoint|breast|immune|aspirin
28
      27 dbs|ps|expectations|pd|preoperative
      28 brentuximab|vedotin|classic|nivolumab|checkpoint
      29 wl|calorie|aas|ba|crc
# with 3,044 more rows
```

The default takes the lower case version of the terms, but (particularly with acronyms) it may be better to preserve the capitalization of the terms. Here is how we can do that in this example:

```
> tfidf <- ctgov_tfidf(z[['description']], tolower = FALSE)</pre>
> print(tfidf, n = 30)
# A tibble: 3,074 Œ 2
     doc terms
   <int> <chr>
       0 heightened|AKI|providing|abdominal|System
 2
       1 pollution|MS|air|viral|France
 3
       2 FMT|diversity|microbiota|weight|gut
 4
       3 nerves|landmarks|bony|guidance|knee
 5
       4 H1|inadequately|spontaneous|suffering|comparison
 6
       5 seroma|drain|categorical|variables|regression
 7
       6 Mediterranean|vagal|nerve|diet|depression
       7 Whole|Veterans|package|Health|mental
 8
 9
       8 suicidal|ideation|telehealth|engagement|counseling
10
       9 BCT|CE|breast|structures|cancers
11
      10 athletes|pathways|exercise|biomarkers|strenuous
12
      11 Acetylsalicylic|Acid|vessels|artery|affects
13
      12 mHealth|90|monitoring|organ|impact
14
      13 sugar|glucose|sensor|doctors|venous
15
      14 variant|CoV2|SARS|Beta|vaccine
16
      15 scenario|uncertainties|oncological|relating|real
17
      16 itch|epigenetic|mechanisms|chronic|antagonists
18
      17 9vHPV|1526|HIV|living|uninfected
19
      18 dengue|fever|permeability|five|vascular
20
      19 influenza|ICU|aspergillosis|EORTC|pathogen
      20 THC|appetite|stimulating|subjective|analgesic
21
22
      21 antibiotics|decide|how|prescribed|pneumonia
23
      22 counseling|education|behavioral|his|behavior
24
      23 intrauterine|adhesiolysis|film|named|adhesion
25
      24 purifiers|cardiopulmonary|air|students|indicators
26
      25 French|Alzheimers|efficiency|controversy|reimbursed
```

```
27 26 checkpoint|Immune|breast|immune|aspirin
28 27 DBS|PS|expectations|PD|preoperative
29 28 vedotin|brentuximab|nivolumab|classic|checkpoint
30 29 WL|calorie|AAs|BA|CRC
# with 3,044 more rows
```

We can also refine the results by including fewer rare terms. The argument min_df specifies the minimal proportion of documents that must contain a term for it to be returned as a keyword; the upper bound can also be specified with the argument max_df.

```
> tfidf <- ctgov_tfidf(z[['description']], min_df = 0.02, max_df = 0.2)</pre>
> print(tfidf, n = 30)
# A tibble: 3,072 Œ 2
     doc terms
   <int> <chr>
       0 injury|support|cardiovascular|performance|feasibility
 1
       1 impact|care|health|risk|better
 2
 3
       2 weight|loss|body|10|least
 4
       3 but | these | compare | two | which
 5
       4 adult|tolerability|chronic|placebo|participants
       5 outcome|multiple|analysis|evaluated|performed
       6 depression|function|assess|efficacy
       7 support|health|level|care|primary
 8
       8 improved|support|high|intervention|care
10
       9 breast|out|being|performed|if
      10 exercise|events|associated|compared|inflammation
11
      11 condition|heart|information|blood|when
12
13
      12 impact|days|objective|outcome|secondary
      13 levels|diabetes|blood|level|device
14
15
      14 vaccine|novel|include|label|open
16
      15 about | new| studies | free | benefit
17
      16 chronic|examine|testing|disorder|while
      17 vaccine|women|among|those|dose
      18 death|treat|syndrome|pilot|evaluation
19
20
      19 pulmonary|incidence|observational|multi|identify
21
      20 alone|combination|assess|effects
22
      21 how|often|children|if|not
      22 diseases|chronic|follow|it|changes
23
      23 novel|aims|controlled|randomized|efficacy
24
25
      24 explore|aims|changes|function|health
      25 non|disease|cognitive|approach|currently
26
27
      26 breast|immune|approximately|called|cancer
      27 postoperative|improvement|result|brain|specific
29
      28 treated|cells|may|cancer|ability
30
      29 prevention|interventions|weight|its|no
  with 3,042 more rows
```

Any number of text fields can be passed to the ctgov_tokens function; all of the fields for a specific trial are pasted together and treated a single block of text.

Document Similarity

Finally, the package also provides a function for producing similarity scores based on the text fields of the studies. Here, we will produce a similarity matrix based on the description field of Interventional, Industry-sponsored, Phase 2 trials.

```
> z <- ctgov_query(
+    study_type = "Interventional", sponsor_type = "Industry", phase = "Phase 2"
+ )
+    scores <- ctgov_text_similarity(z[['description']], min_df = 0, max_df = 0.1)
+    dim(scores)
[1] 147 147</pre>
```

The returned value is a square matrix with one row and one colum for each clinical trial in the set. We can use these scores to find studies that are particularly close to one another in the words used within their descriptions. Here for example we can see five studies that use similar terms in their descriptions:

- > index <- order(scores[,100], decreasing = TRUE)[1:5]</pre>
- > z[['brief_title']][index]
- [1] "AL-38583 Ophthalmic Solution for Allergic Conjunctivitis Associated Inflammation"
- [2] "Safety and Efficacy of BRM421 for Dry Eye Syndrome"
- [3] "Phosphorylcholine PC-mAb Effects in Subjects With Elevated Lipoprotein a"
- [4] "Safety, Clinical Tolerability and Immunogenicity of Increasing Doses of gpASIT+TM"
- [5] "Phase 2 Clinical Trial of CartiLife in the United States"

Further post-processing can be done with the similarity scores, such as spectral clustering and dimensionality reduction.

Case Study

Conclusions

Bibliography

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