

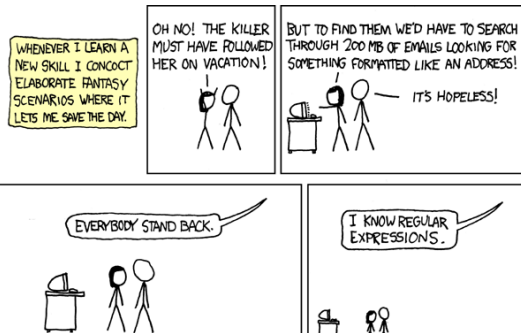
Regular Expressions, Web Scraping, APIs

JSC 370: Data Science II

- Learn how to use an API

Regular Expressions: What is it?

A regular expression (shortened as regex or regexp; also referred to as rational expression) is a sequence of characters that define a search pattern. -- [Wikipedia](#)



Regular Expressions: Why should you care?

We can use Regular Expressions for:

- Validating data fields, email address, numbers, etc.
- Searching text in various formats, e.g., addresses, there are many ways to write an address.
- Replace text, e.g., different spellings, Storm, Stôrm, Stórm to Storm.
- Remove text, e.g., tags from an HTML text, <name>George</name> to George.

Regular Expressions 101: Metacharacters

What makes *regex* special is metacharacters. While we can always use *regex* to match literals like `dog`, `human`, `1999`, we only make use of all *regex* power when using metacharacters:

- `.` Any character except new line
- `^` beginning of the text
- `$` end of the text
- `[regex]` Match a single character in `regex`, e.g.
 - `[0123456789]` Any number
 - `[0-9]` Any number in the range 0-9
 - `[a-z]` Lower-case letters
 - `[A-Z]` Upper-case letters
 - `[a-zA-Z]` Lower or upper case letters.
 - `[a-zA-Z0-9]` Any alpha-numeric
- `[^regex]` Match any except those in `regex`, e.g.
 - `[^0123456789]` Match any except a number
 - `[^0-9]` Match anything except in the range 0-9
 - `[^./]` any except dot, slash, and space.

Regular Expressions 101: Metacharacters (cont. 1)

Ranges, e.g., `0-9` or `a-Z`, are locale- and implementation-dependent, meaning that the range of lower case letters may vary depending on the OS's language. To solve for this problem, you could use [Character classes](#). Some examples:

- `[:lower:]` lower case letters in the current locale, could be `[a-z]`
- `[:upper:]` upper case letters in the current locale, could be `[A-Z]`
- `[:alpha:]` upper and lower case letters in the current locale, could be `[a-zA-Z]`
- `[:digit:]` Digits: `0 1 2 3 4 5 6 7 8 9`
- `[:alnum:]` Alpha numeric characters `[:alpha:]` and `[:digit:]`.
- `[:punct:]` Punctuation characters: `! " # $ % & ' () * + , - . / : ; < = > ? @ [\] ^ _ ` { | } ~`.

For example, in the locale `en_US`, the word `Hóla` IS NOT fully matched by `[a-zA-Z]+`, but IT IS fully matched by `[:alpha:]+`.

Other important Metacharacters:

- `\s` white space, equivalent to `[\r\n\t\f\v]`
- `|` or (logical or).

Regular Expressions 101: Metacharacters (cont. 2)

These usually come together with specifying how many times (repetition):

- `regex?` Zero or one match.
- `regex*` Zero or more matches
- `regex+` One or more matches
- `regex{n, }` At least `n` matches
- `regex{, m}` at most `m` matches
- `regex{n, m}` Between `n` and `m` matches.

Where `regex` is a regular expression

Regular Expressions 101: Metacharacters (cont. 3)

There are other operators that can be very useful,

- `(regex)` Group capture.
- `(?: regex)` Group operation without capture.
- `(?= regex)` Look ahead (match)
- `(?! regex)` Look ahead (don't match)
- `(?<= regex)` Look behind (match)
- `(?<! regex)` Look behind (don't match)

Group captures can be reused with `\1`, `\2`, ..., `\n`.

More (great) information here <https://regex101.com/>

Regular Expressions 101: Examples

Here we are extracting the first occurrence of the following regular expressions (using `stringr::str_extract()`):

regex	Hanna Perez [name] The 年 year was 1999 HaHa, @abc said that GoGo trojans #2020!			
.{5}	Hanna	The 年	HaHa,	GoGo
n{2}	nn			
[0-9]+		1999		2020
\s[a-zA-Z]+\s	Perez	year	said	trojans
\s[:alpha:]+\s	Perez	年	said	trojans
[a-zA-Z]+ [a-zA-Z]+	Hanna Perez	year was	abc said	GoGo trojans
([a-zA-Z]+\s?){2}	Hanna Perez	The	HaHa	GoGo trojans
([a-zA-Z]+\s)\1	nn		HaHa	GoGo
(@ #)[a-z0-9]+			@abc	#2020
(?<=# @)[a-z0-9]+			abc	2020

`[a - z]+` `[name]`

Regular Expressions 101: Examples (cont. 1)

1. `.{5}` Match **any character** (except line end) five times.
2. `n{2}` Match the letter **n** twice.
3. `[0-9]+` Match **any number** at least once
4. `\s[a-zA-Z]+\s` Match a **space, any lower or upper case letter** at least once, and a **space**.
5. `\s[:alpha:]+\s` Same as before but this time .
6. `[a-zA-Z]+ [a-zA-Z-Z]+` Match two sets of letters separated by one space.
7. `([a-zA-Z]+\s?){2}` Match **any lower or upper case letter** at least once, maybe followed by a white space, twice.
8. `([a-zA-Z]+\s)\1` Match **any lower or upper case letter** at least once, and then match the same pattern again.
9. `(@|#[a-z0-9]+)` Match either the **@** or **#** symbol, followed by one or more **lower case letter** or **number**.
10. `(?<=#|@)[a-z0-9]+` Match one or more **lower case letter** or **number** that follows either the **@** or **#** symbol.
11. `\[[a-z]+\]` Match the symbol **[**, at least one **lower case letter**, and the symbol **]**.

Regular Expressions 101: Functions in R

1. Lookup text: `base::grepl()`, `stringr::str_detect()`.
2. Similar to `which()`, which elements are TRUE `base::grep()`, `stringr::str_which()`
3. Replace the first instance: `base::sub()`, `stringr::str_replace()`
4. Replace all instances: `base::gsub()`, `stringr::str_replace_all()`
5. Extract text: `base::regmatches()`, `stringr::str_extract()` and `stringr::str_extract_all()`.

Regular Expressions 101: Functions in R (cont.)

For example, like in Twitter, let's create a regex that matches usernames or hashtags with the following pattern:

```
(@|#)([[:alnum:]]+)
```

Code	@Hanna Perez [name] #html	The @年 year was 1999	HaHa, @abc said that @z
str_detect(text, pattern) or grepl(pattern, text)	TRUE	TRUE	TRUE
str_extract(text, pattern)	@Hanna	@年	@abc
str_extract_all(text, pattern)	[@Hanna, #html]	[@年]	[@abc, @z]
str_replace(text, pattern, "\1justinbieber")	@justinbieber Perez [name] #html	The @justinbieber year was 1999	HaHa, @justinbieber said that @z
str_replace_all(text, pattern, "\1justinbieber")	@justinbieber Perez [name] #justinbieber	The @justinbieber year was 1999	HaHa, @justinbieber said that @justinbieber

Note: While it is not showing in the table, the group replacement was escaped, i.e., `\\1` instead of `\1` in the code.

Data

This week we will use a dataset consisting of medical transcriptions from <https://www.mtsamples.com/>. See the readme on the [course git](#). The dataset consists of 4999 rows and 6 columns: “X”, “description”, “medical_specialty”, “sample_name”, “transcription” and “keywords”.

```
library(data.table)
library(stringr)

fn <- "mtsamples.csv"
if (!file.exists(fn))
  download.file(
    url = "https://github.com/JSC370/jsc370-2023/blob/main/data/medical_transcriptions/mtsamples"
    destfile = fn
  )
mtsamples <- fread(fn, sep = ",", header = TRUE)
names(mtsamples)

## [1] "V1"          "description"  "medical_specialty"
## [4] "sample_name" "transcription" "keywords"
```

Regex to Lookup Text: Tumor

We would like to see if this is a tumor-related entry. For that we can search through the "description" using `grepl` in the following code:

```
# How many entries contain the word tumor
mtsamples[grepl("tumor", description, ignore.case = TRUE), .N]

## [1] 67

# Generating a column tagging tumor
mtsamples[, tumor_related := grepl("tumor", description, ignore.case = TRUE)]

# Taking a look at a few examples
mtsamples[tumor_related == TRUE, .(description)][1:3,]

##
## 1: Transurethral resection of a medium bladder tumor (TURBT), left lat
## 2: Transurethral resection of the bladder tumor (TURB
## 3: Cystoscopy, transurethral resection of medium bladder tumor (4.0 cm in diameter), and direct blad
```

Notice the `ignore.case = TRUE`. This is equivalent to transforming the text to lower case using `tolower()` before passing the text to the regular expression function.

Regex Lookup text: Pronoun of the patient

Now, let's try to guess the pronoun of the patient. To do so, we could tag by using the words *he, his, him, they, them, theirs, ze, hir, hers, she, her* (see [this article on sexist text](#)):

```
mtsamples[, pronoun := str_extract(
  string = tolower(transcription),
  pattern = "he|his|him|they|them|theirs|ze|hir|hirs|she|hers|her"
)]
mtsamples[1:10, pronoun]
```

```
## [1] "his" "his" "his" "ze" "he" "he" "he" "he" "he" "ze"
```

```
mtsamples[, table(pronoun, useNA = "always")]
```

```
## pronoun
##   he him hir his she them ze <NA>
## 2558   6  14 934  46  13  43   68
```

What is the problem with this approach?

Regex Lookup text: Pronoun of the patient (cont. 1)

For this we use the following regular expression:

```
(?<=\W|^)(he|his|him|they|them|theirs|ze|hir|hirs|she|hers|her)(?=\W|$)
```

Bit by bit this is:

- `(?<=regex)` lookback search.
 - `\W` any non alpha numeric character, this is equivalent to `^[a\W:]`, | or
 - `^` the beginning of the text,
- `he|his|him...` any of these words,
- `(?=regex)` followed by,
 - `\W` any non alpha numeric character, this is equivalent to `^[a\W:]`, | or
 - `$` the end of the text.

```
mtsamples[, pronoun := str_extract(  
  string = tolower(transcription),  
  pattern = "(?<=\W|^)(he|his|him|they|them|theirs|ze|hir|hirs|she|hers|her)(?=\W|$)"  
)]  
mtsamples[1:10, pronoun]
```

```
## [1] "she" "he" "he" NA NA "she" "she" NA NA NA
```

Regex Lookup text: Pronoun of the patient (cont. 2)

```
mtsamples[, table(pronoun, useNA = "always")]
```

```
## pronoun  
##   he  her  him  his  she  them  they  <NA>  
##  767  394   29  361  870   18   67 1176
```

Regex Extract Text: Type of Cancer

- Imagine now that you need to see the types of cancer mentioned in the data.
- For simplicity, let's assume that, if specified, it is in the form of `TYPE cancer`, i.e. single word.
- We are interested in the word before cancer, how can we capture this?

Regex Extract Text: Type of Cancer (cont 1.)

We can just try to **extract** the phrase "[some word] cancer", in particular, we could use the following regular expression

```
[[:alnum:]-_]{4,}\\s*cancer
```

Where

- `[[:alnum:]-_]{4,}` captures any alphanumeric character, including `-` and `_`. Furthermore, for this match to work there must be at least 4 characters,
- `\\s*` captures 0 or more white-spaces, and
- `cancer` captures the word cancer:

```
mtsamples[, cancer_type := str_extract(tolower(keywords), "[[:alnum:]-_]{4,}\\s*cancer")]
mtsamples[, table(cancer_type)]
```

```
## cancer_type
##      anal cancer      bladder cancer      breast cancer      colon cancer
##              1              6              16              12
## endometrial cancer esophageal cancer      lung cancer      ovarian cancer
##              5              1              8              1
##      papillary cancer      prostate cancer      uterine cancer
##              2              14              4
```

Fundamentals of Web Scrapping

What?

Web scraping, web harvesting, or web data extraction is data scraping used for extracting data from websites --
[Wikipedia](#)

How?

- The [rvest](#) R package provides various tools for reading and processing web data.
- Under-the-hood, `rvest` is a wrapper of the [xml2](#) and [httr](#) R packages.

(in the case of [dynamic websites](#), take a look at [selenium](#)))

Web scraping raw HTML: Example

We would like to capture the table of COVID-19 death rates per country directly from Wikipedia.

```
library(rvest)
library(xml2)

# Reading the HTML table with the function xml2::read_html
covid <- read_html(
  x = "https://en.wikipedia.org/wiki/COVID-19_pandemic_death_rates_by_country"
)

# Let's see the output
covid

## {html_document}
## <html class="client-nojs vector-feature-language-in-header-enabled vector-feature-language-in-main-page-disabled vector-feature-search-enabled vector-feature-l10n-enabled vector-feature-media-autoplay-enabled vector-feature-table-of-contents-enabled vector-feature-sticky-header-enabled vector-feature-page-not-found vector-feature-pinned-sections-enabled">
## [1] <head>\n<meta http-equiv="Content-Type" content="text/html; charset=UTF-8 ...
## [2] <body class="skin-vector skin-vector-search-vue vector-toc-pinned mediawiki ...
```

Web scraping raw HTML: Example (cont 1.)

- We want to get the HTML table that shows up in the doc. To do this, we can use the function `xml2::xml_find_all()` and `rvest::html_table()`
- The first will locate the place in the document that matches a given **XPath** expression.
- [XPath](#), XML Path Language, is a query language to select nodes in a XML document.
- A nice tutorial can be found [here](#)
- Modern Web browsers make it easy to use XPath!

Live Example! (inspect elements in [Google Chrome](#), [Mozilla Firefox](#), [Internet Explorer](#), and [Safari](#))

Web scraping with xml2 and the rvest package (cont. 2)

Now that we know what is the path, let's use that and extract

```
table <- xml2::xml_find_all(covid, xpath = "/html/body/div[3]/div[3]/div[5]/div[1]/table[2]")
table <- html_table(table) # This returns a list of tables
#head(table)
table

## list()
```


Web APIs

What?

A Web API is an application programming interface for either a web server or a web browser. -- [Wikipedia](#)

Some examples include: [twitter API](#), [facebook API](#), [Gene Ontology API](#)

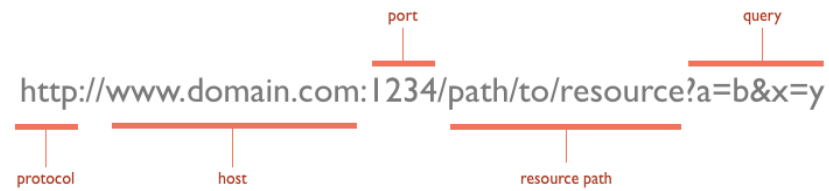
How?

You can request data, the **GET method**, post data, the **POST method**, and do many other things using the [HTTP protocol](#).

How in R?

For this part, we will be using the `httr()` package, which is a wrapper of the `curl()` package, which in turn provides access to the `curl` library that is used to communicate with APIs.

Web APIs with curl



Structure of a URL (source: ["HTTP: The Protocol Every Web Developer Must Know - Part 1"](#))

Web APIs with curl

Under-the-hood, the `httr` (and thus `curl`) sends request somewhat like this

```
curl -X GET https://google.com -w "%{content_type}\n%{http_code}\n"
```

A get request (`-X GET`) to `https://google.com`, which also includes (`-w`) the following: `content_type` and `http_code`:

```
<HTML><HEAD><meta http-equiv="content-type" content="text/html; charset=utf-8">
<TITLE>301 Moved</TITLE></HEAD><BODY>
<H1>301 Moved</H1>
The document has moved
<A HREF="https://www.google.com/">here</A>.
</BODY></HTML>
text/html; charset=UTF-8
301
```

We use the `httr` R package to make life easier.

Web API Example 1: Gene Ontology

- We will make use of the [Gene Ontology API](#).
- We want to know what genes (human or not) are **involved in** the function **antiviral innate immune response** (go term [GO:0140374](#)), looking only at those annotations that have evidence code [ECO:0000006](#) (experimental evidence):

```
library(httr)
go_query <- GET(
  url   = "http://api.geneontology.org/",
  path  = "api/bioentity/function/GO:0140374/genes",
  query = list(
    evidence           = "ECO:0000006",
    relationship_type = "involved_in"
  ),
  # May need to pass this option to curl to allow to wait for at least
  # 60 seconds before returning error.
  config = config(
    connecttimeout = 60
  )
)
```

We could have also passed the full URL directly...

Web API Example 1: Gene Ontology (cont. 1)

Let's take a look at the curl call:

```
curl -X GET "http://api.geneontology.org/api/bioentity/function/G0:0140374/genes?evidence=ECO%3A0000000"
```

What `httr::GET()` does:

```
> go_query$request
## <request>
## GET http://api.geneontology.org/api/bioentity/function/G0:0140374/genes?evidence=ECO%3A0000000
## Output: write_memory
## Options:
## * useragent: libcurl/7.58.0 r-curl/4.3 httr/1.4.1
## * connecttimeout: 60
## * httpget: TRUE
## Headers:
## * Accept: application/json, text/xml, application/xml, */*
```

Web API Example 1: Gene Ontology (cont. 2)

Let's take a look at the response:

```
go_query
```

```
## Response [http://api.geneontology.org/api/bioentity/function/G0:0140374/genes?evidence=ECO%3A0000006&]
##   Date: 2023-02-13 16:16
##   Status: 200
##   Content-Type: application/json
##   Size: 74.5 kB
## {"associations": [{"id": "4d4749094d47493a31313030353235094e6d62720909474f3a3..."}
```

Remember the codes:

- 1xx: Information message
- 2xx: Success
- 3xx: Redirection
- 4xx: Client error
- 5xx: Server error

Web API Example 1: Gene Ontology (cont. 3)

We can extract the results using the `httr::content()` function

```
dat <- content(go_query)
dat <- lapply(dat$associations, function(a) {
  data.frame(
    Gene      = a$subject$id,
    taxon_id  = a$subject$taxon$id,
    taxon_label = a$subject$taxon$label
  )
})
dat <- do.call(rbind, dat)
str(dat)

## 'data.frame':   61 obs. of  3 variables:
## $ Gene      : chr  "MGI:1100525" "MGI:2441706" "MGI:2385051" "MGI:1099786" ...
## $ taxon_id  : chr  "NCBITaxon:10090" "NCBITaxon:10090" "NCBITaxon:10090" "NCBITaxon:10090" ...
## $ taxon_label: chr  "Mus musculus" "Mus musculus" "Mus musculus" "Mus musculus" ...
```

Web API Example 1: Gene Ontology (cont. 4)

The structure of the result will depend on the API. In this case, the output was a JSON file, so the content function returns a list in R. In other scenarios it could return an XML object (we will see more in the lab)

```
knitr::kable(head(dat),  
  caption = "Genes experimentally annotated with the function\  
  **antiviral innate immune response** (GO:0140374)"  
)
```

Table: Genes experimentally annotated with the function **antiviral innate immune response** (GO:0140374)

Gene	taxon_id	taxon_label
MGI:1100525	NCBITaxon:10090	Mus musculus
MGI:2441706	NCBITaxon:10090	Mus musculus
MGI:2385051	NCBITaxon:10090	Mus musculus
MGI:1099786	NCBITaxon:10090	Mus musculus
MGI:1099786	NCBITaxon:10090	Mus musculus
MGI:1913565	NCBITaxon:10090	Mus musculus

Web API Example 2: Using Tokens

- Sometimes, APIs are not completely open, you need to register.
- The API may require to login (user+password), or pass a token.
- In this example, I'm using a token which I obtained [here](#)
- You can find information about the [National Centers for Environmental Information](#) API [here](#)

Web API Example 2: Using Tokens (cont. 1)

- The way to pass the token will depend on the API service.
- Some require authentication, others need you to pass it as an argument of the query, i.e., directly in the URL.
- In this case, we pass it on the header.

```
stations_api <- GET(  
  url      = "https://www.ncdc.noaa.gov",  
  path     = "cdo-web/api/v2/stations",  
  config = add_headers(  
    token = "[YOUR TOKEN HERE]"  
  ),  
  query    = list(limit = 1000)  
)
```

This is equivalent to using the following query

```
curl --header "token: [YOUR TOKEN HERE]" \  
https://www.ncdc.noaa.gov/cdo-web/api/v2/stations?limit=1000
```

Note: This won't run, you need to get your own token

Web API Example 2: Using Tokens (cont. 2)

Again, we can recover the data using the `content()` function:

```
ans <- content(stations_api)
ans$results[[1]]
## $elevation
## [1] 139
##
## $mindate
## [1] "1948-01-01"
##
## $maxdate
## [1] "2014-01-01"
##
## $latitude
## [1] 31.5702
##
## $name
## [1] "ABBEVILLE, AL US"
##
## $datacoverage
## [1] 0.8813
##
## $id
## [1] "COOP:010008"
```

Web API Example 3: HHS health recommendation

Here is a last example. We will use the Department of Health and Human Services API for "[...] demographic-specific health recommendations" (details at health.gov)

```
health_advises <- GET(  
  url = "https://health.gov/",  
  path = "myhealthfinder/api/v3/myhealthfinder.json",  
  query = list(  
    lang = "en",  
    age = "32",  
    sex = "male",  
    tobaccoUse = 0  
  ),  
  config = c(  
    add_headers(accept = "application/json"),  
    config(connecttimeout = 60)  
  )  
)
```

Web API Example 3: HHS health recommendation (cont. 1)

Let's see the response

health_advises

```
## Response [https://health.gov/myhealthfinder/api/v3/myhealthfinder.json?lang=en&age=32&sex=male&tobacco=0]
##   Date: 2023-02-13 16:16
##   Status: 200
##   Content-Type: application/json
##   Size: 359 kB
## {
##   "Result": {
##     "Error": "False",
##     "Total": 18,
##     "Query": {
##       "ApiVersion": "3",
##       "ApiType": "myhealthfinder",
##       "TopicId": null,
##       "ToolId": null,
##       "CategoryId": null,
##     }
##   }
## }
```

Web API Example 3: HHS health recommendation (cont. 2)

```
# Extracting the content
health_advises_ans <- content(health_advises)

# Getting the titles
txt <- with(health_advises_ans$Result$Resources, c(
  sapply(all$Resource, "[", "Title"),
  sapply(some$Resource, "[", "Title"),
  sapply(`You may also be interested in these health topics:`$Resource, "[", "Title")
))
cat(txt, sep = "; ")
```

Hepatitis C Screening: Questions for the Doctor; Protect Yourself from Seasonal Flu; Talk with Your Doctor About Depression; Drink Alcohol Only in Moderation; Get Vaccines to Protect Your Health (Adults Ages 19 to 49); Get Tested for HIV; Get Your Blood Pressure Checked; Quit Smoking; Talk with Your Doctor About Drug Misuse; Watch Your Weight; Eat Healthy; Testing for Syphilis: Questions for the Doctor; Protect Yourself from Hepatitis B; Testing for Latent Tuberculosis: Questions for the Doctor; Manage Stress; Alcohol Use: Conversation Starters; Get Active; Quitting Smoking: Conversation Starters

Summary

- We learned about regular expressions with the package **stringr** (a wrapper of **stringi**)
- We can use regular expressions to detect (`str_detect()`), replace (`str_replace()`), and extract (`str_extract()`) expressions.
- We looked at web scraping using the **rvest** package (a wrapper of **xml2**).
- We extracted elements from the HTML/XML using `xml_find_all()` with XPath expressions.
- We also used the `html_table()` function from **rvest** to extract tables from HTML documents.
- We took a quick review on Web APIs and the Hyper-text-transfer-protocol (HTTP).
- We used the **httr** R package (wrapper of **curl**) to make GET requests to various APIs
- We even showed an example using a token passed via the `header`.
- Once we got the responses, we used the `content()` function to extract the message of the response.

Detour on CURL options

Sometimes you will need to change the default set of options in CURL. You can checkout the list of options in `curl::curl_options()`. A common hack is to extend the time-limit before dropping the connection, e.g.:

Using the **Health IT** API from the US government, we can obtain the **Electronic Prescribing Adoption and Use by County** (see docs [here](#))

The problem is that it usually takes longer to get the data, so we pass the config option `connecttimeout` (which corresponds to the flag `--connect-timeout`) in the curl call (see next slide)

Detour on CURL options (cont.)

```
ans <- httr::GET(
  url = "https://dashboard.healthit.gov/api/open-api.php",
  query = list(
    source = "AHA_2008-2015.csv",
    region = "California",
    period = 2015
  ),
  config = config(
    connecttimeout = 60
  )
)

> ans$request
# <request>
# GET https://dashboard.healthit.gov/api/open-api.php?source=AHA_2008-2015.csv&region=California
# Output: write_memory
# Options:
# * useragent: libcurl/7.58.0 r-curl/4.3 httr/1.4.1
# * connecttimeout: 60
# * httpget: TRUE
# Headers:
# * Accept: application/json, text/xml, application/xml, */*
```

Regular Expressions: Email validation

This is the official regex for email validation implemented by [RFC 5322](#)

```
(?:[a-z0-9!#$%&'*/+=?^_`{|}~-]+(?:\\. [a-z0-9!#$%&'*/+=?^_`{|}~-]+)*|"(?:[\x01-\x08
\x0b\x0c\x0e-\x1f\x21\x23-\x5b\x5d-\x7f]|\\[\x01-\x09\x0b\x0c\x0e-\x7f])*")@(?:(?
: [a-z0-9](?:[a-z0-9-]*[a-z0-9])?\.)+[a-z0-9](?:[a-z0-9-]*[a-z0-9])?|\[(?:(?:(2(5[
0-5]| [0-4][0-9])|1[0-9][0-9]| [1-9]?[0-9]))\.\.){3}(?: (2(5[0-5]| [0-4][0-9])|1[0-9][0
-9]| [1-9]?[0-9])|[a-z0-9-]*[a-z0-9]: (?:[\x01-\x08\x0b\x0c\x0e-\x1f\x21-\x5a\x53-\x
7f]|\\[\x01-\x09\x0b\x0c\x0e-\x7f]))+)\])
```

See the corresponding post in [StackOverflow](#)