rbison vignette - wrapper to the USGS BISON API

About the package

rbison is an R package to search and retrieve data from the USGS BISON service. rbison wraps R code around the BISON API to allow you to talk to the BISON database from R.

BISON has occurrence data for the US only.

BISON is a node of the Global Biodiversity Information Facility (GBIF) - i.e., you can get data that's available in BISON via GBIF instead if you want.

Info

See here for API docs for the BISON API.

Quick start

Install rbison

```
# install.packages('devtools'); library(devtools); install_github('rbison',
# 'ropensci')
library(rbison)
library(sp)
library(plyr)
library(httr)
library(gridExtra)
library(gridExtra)
library(ggplot2)
library(maps)
library(rgeos)
library(rgdal)
```

Notice that the function bisonmap automagically selects the map extent to plot for you, being one of the contiguous lower 48 states, or the lower 48 plus AK and HI, or a global map

If some or all points outside the US, a global map is drawn, and throws a warning. . You may want to make sure the occurrence lat/long coordinates are correct.

get data

```
out <- bison(species = "Helianthus annuus", count = 10)</pre>
```

inspect summary

bison_data(out)

total observation fossil specimen literature unknown living centroid 1 4424 13 102 1303 1106 1892 8 1106

map occurrences

bisonmap(out)



Figure 1: plot of chunk unnamed-chunk-1 $\,$

All points within the US (including AK and HI)

${\rm get}\ {\rm data}$

```
out <- bison(species = "Bison bison", count = 600)</pre>
```

inspect summary

bison_data(out)

total observation fossil specimen unknown 1 781 38 4 722 17

map occurrences

bisonmap(out)



Figure 2: plot of chunk six

All points within the contiguous 48 states

get data

```
out <- bison(species = "Aquila chrysaetos", count = 600)</pre>
```

inspect summary

bison_data(out)

total observation fossil specimen literature unknown centroid 1 41780 39334 30 1664 118 634 904

map occurrences

bisonmap(out)



Figure 3: plot of chunk nine

With any data returned from a bison call, you can choose to plot county or state level data

Counties - using last data call for Aquila

bisonmap(out, tomap = "county")



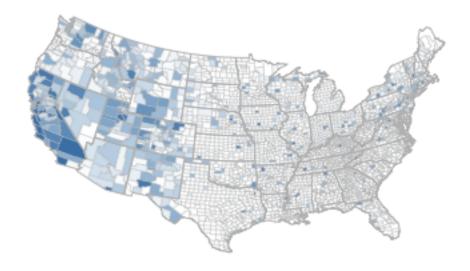


Figure 4: plot of chunk ten

States - using last data call for Aquila

```
bisonmap(out, tomap = "state")
```

Constrain search with county IDs or bounding boxes

Constrain search to a certain county. Check out this site to get state and county fips codes. Fips codes are like so: First two digits are the state code - last three are the county code. For example the $\theta 6$ in 06037 is the state of California, and the $\theta 37$ is the Los Angeles county.

```
out <- bison(species = "Helianthus annuus", countyFips = 6037)
# Inspect summary
bison_data(out)

total observation fossil specimen literature unknown centroid
1 23 1 3 12 1 6 1

# By default, the query only returned 10 records
bison_data(out, "data_df")</pre>
```



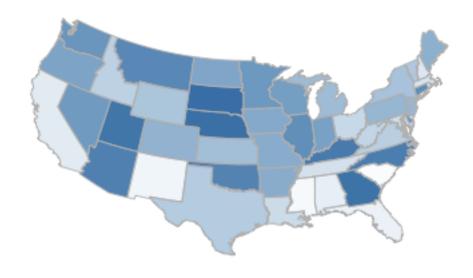


Figure 5: plot of chunk eleven

2 687026030 Helianthus annuus -118.2 34.09 3 40886954 Helianthus annuus -118.9 34.80 4 40886955 Helianthus annuus -118.9 34.80 5 67195635 Helianthus annuus -118.8 34.70 6 67195791 Helianthus annuus -118.5 34.30 7 67195582 Helianthus annuus -118.4 34.04 8 40886953 Helianthus annuus -118.1 34.70 9 40886951 Helianthus annuus -118.1 34.70 10 40886950 Helianthus annuus -118.1 34.70 provider 1 USDA PLANTS 2 iNaturalist.org 3 US National Plant Germplasm System 4 US National Plant Germplasm System 5 < NA> 6 < NA> 7		id		name	longitude	latitude	
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7 <na></na>	5				<na></na>		
	6				<na></na>		
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9 US National Plant Germplasm System	9	US Nation	al Plant Ge	rmplasm	System		
10 US National Plant Germplasm System	10	US Nation	al Plant Ger	rmplasm	System		

Or specify county by its actual name - probably much easier.

```
# Inspect summary
bison_data(out)
 total observation fossil specimen literature unknown centroid
    23
                 1
                        3
                                12
                                            1
# By default, the query only returned 10 records
bison_data(out, "data_df")
          id
                         name longitude latitude
  832095838 Helianthus annuus
                                 -118.4
                                           33.39
  687026030 Helianthus annuus
                                 -118.2
                                           34.09
                                 -118.9
                                           34.80
3
   40886954 Helianthus annuus
4
   40886955 Helianthus annuus
                                 -118.9
                                           34.80
   67195635 Helianthus annuus
                                 -118.8
                                           34.70
5
6
   67195791 Helianthus annuus
                                 -118.5
                                           34.30
   67195582 Helianthus annuus
7
                                 -118.4
                                           34.04
   40886953 Helianthus annuus
                                 -118.1
                                           34.70
                                           34.70
9
   40886951 Helianthus annuus
                                 -118.1
10 40886950 Helianthus annuus
                                 -118.1
                                           34.70
                            provider
1
                         USDA PLANTS
2
                     iNaturalist.org
3
  US National Plant Germplasm System
  US National Plant Germplasm System
5
                                <NA>
6
                                <NA>
7
                                <NA>
  US National Plant Germplasm System
  US National Plant Germplasm System
10 US National Plant Germplasm System
bison will help you if you spell the name wrong, or use a partial name. The results are not printed below,
but you would get a prompt asking you to pick between the two counties that start with Los.
bison(species = "Helianthus annuus", county = "Los")
Constrain search to a amorphous area. Check out the Wikipedia page here for an in depth look at
the options, terminology, etc.
# Inspect summary
bison_data(out)
 total literature centroid
1
     1
                1
                         1
```

out <- bison(species = "Helianthus annuus", county = "Los Angeles")

```
# The data
bison_data(out, "data_df")
         id
                         name longitude latitude
                                                    provider
1 832098012 Helianthus annuus
                                -110.7
                                          38.99 USDA PLANTS
Constrain search to a certain aoibbox. An aoibbox uses the format minx,miny,maxx,maxy.
out <- bison(species = "Helianthus annuus", aoibbox = "-120.31,35.81,-110.57,40.21")
# Inspect summary
bison_data(out)
  total observation fossil specimen literature unknown centroid
    145
                  6
                        8
                                34
                                           25
                                                    72
                                                             25
# The data, by default, the query only returned 10 records
bison_data(out, "data_df")
           id
                          name longitude latitude
                                                     provider
1 1800986531 Helianthus annuus
                                  -112.1
                                            37.18
                                                        BISON
2 1800986532 Helianthus annuus
                                  -112.1
                                            37.18
                                                        BISON
3 1800986533 Helianthus annuus
                                  -112.1
                                            37.18
                                                        BISON
4 1800986534 Helianthus annuus
                                  -112.1
                                            37.18
                                                        BISON
5 1800986645 Helianthus annuus
                                            37.13
                                  -111.8
                                                         BISON
6 1800986646 Helianthus annuus
                                  -111.8
                                            37.13
                                                        BISON
  832096828 Helianthus annuus
7
                                  -118.4
                                            38.54 USDA PLANTS
8 832098025 Helianthus annuus
                                  -111.8
                                            38.75 USDA PLANTS
   832098012 Helianthus annuus
                                  -110.7
                                            38.99 USDA PLANTS
10 832098018 Helianthus annuus
                                  -113.1
                                            39.05 USDA PLANTS
```