

Package ‘EJAM’

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Title EJAM Environmental Justice Analysis Multisite tool

Version 2.1.1

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Description Tools for summarizing environmental and demographic indicators (such as those in EJScreen) for residents living near any one of a number of specific sites. It uses quad tree search/indexing of block locations, data.table, parallel processing to provide very fast identification of nearby blocks, distances, and aggregation of indicators within each distance. It can be used as a web app, with the user interface provided by the shiny R package.

URL <https://github.com/USEPA/EJAM>

Depends R (>= 2.10),
shiny (>= 1.7.2),
EJAMblockdata,
EJAMfrsdata,
EJAMbatch.summarizer,
EJAMejscreenapi

Imports attempt,
config (>= 0.3.1),
data.table,
DBI,
doSNOW,
DT,
EJAMejscreenapi,
foreach,
ggplot2,
glue,
golem (>= 0.3.3),
htmltools,
leaflet,
magrittr,
openxlsx,
pdist,
pkgload,
readxl,
RMySQL,

SearchTrees,
shinyBS,
shinycssloaders,
shinyjs,
sf,
sp,
tidyverse

Remotes github::USEPA/EJAMblockdata,
github::USEPA/EJAMfrsdata,
github::USEPA/EJAMejscreenapi,
github::USEPA/EJAMbatch.summarizer,
github::USEPA/EJAMejscreendata

Suggests knitr,
rmarkdown,
spelling,
testthat (>= 3.0.0)

Config/testthat/edition 3

Encoding UTF-8

LazyData true

Language en-US

VignetteBuilder knitr

RoxygenNote 7.2.3

Roxygen list(markdown = TRUE)

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.onLoad	<i>Creates index to all US blocks (internal point lat lon) at package load</i>
---------	--

Description

Creates index to all US blocks (internal point lat lon) at package load

Usage

```
.onLoad(libname, pkgname)
```

Arguments

libname	na
pkgname	na

all.equal_functions	<i>helper function for checking possibly different versions of a function with same name in 2 packages</i>
---------------------	--

Description

helper function for checking possibly different versions of a function with same name in 2 packages

Usage

```
## S3 method for class 'equal_functions'  
all(fun = "latlon_infer", package1 = "EJAM", package2 = "EJAMejscreenapi")
```

Arguments

fun	quoted name of function, like "latlon_infer"
package1	quoted name of package, like "EJAM"
package2	quoted name of package, like "EJAMejscreenapi"

Value

TRUE or FALSE

See Also

[dupenames\(\)](#)

app_run_EJAM	<i>Launch the Shiny Application in RStudio</i>
--------------	--

Description

launch Shiny web app from RStudio

Usage

```
app_run_EJAM(  
  onStart = NULL,  
  options = list(),  
  enableBookmarking = NULL,  
  uiPattern = "/",  
  ...  
)
```

Arguments

onStart	A function that will be called before the app is actually run. This is only needed for shinyAppObj, since in the shinyAppDir case, a global.R file can be used for this purpose.
options	Named options that should be passed to the runApp call (these can be any of the following: "port", "launch.browser", "host", "quiet", "display.mode" and "test.mode"). You can also specify width and height parameters which provide a hint to the embedding environment about the ideal height/width for the app.
enableBookmarking	Can be one of "url", "server", or "disable". The default value, NULL, will respect the setting from any previous calls to enableBookmarking() . See enableBookmarking() for more information on bookmarking your app.
uiPattern	A regular expression that will be applied to each GET request to determine whether the ui should be used to handle the request. Note that the entire request path must match the regular expression in order for the match to be considered successful.
...	arguments to pass to golem_opts. See ?golem::get_golem_options for more details.

Details

app_run_EJAM() is like [run_app\(\)](#) app_run_EJAMejscreenapi() is like [EJAMejscreenapi::run_app\(\)](#)
 app_run_EJAMbatch.summarizer() is like [EJAMbatch.summarizer::run_app\(\)](#)

app_server

EJAM app server

Description

EJAM app server

Usage

```
app_server(input, output, session)
```

Arguments

input, output, session

Internal parameters for shiny. DO NOT REMOVE.

bgpts	<i>lat lon of popwtd center of blockgroup, and count of blocks per block group</i>
-------	--

Description

This is just a list of US block groups and how many blocks are in each... It also has the lat and lon roughly of each blockgroup

Details

The point used for each bg is the Census 2020 population weighted mean of the blocks' internal points. It gives an approximation of where people live and where each bg is, which is useful for some situations.

As of 10/2022 it is the EJScreen 2.1 version of data, which uses ACS 2016-2020 and Census 2020. it has all US States, DC, PR, but not "AS" "GU" "MP" "VI"

How lat lon were estimated:

```
# proxistat::bg.pts had a lat/lon internal point for each us block group for Census 2010.
# that had been used to include those lat/lon in ejsscreen::bg21, for convenience.
```

```
> head(proxistat::bg.pts)
      FIPS    aland  awater    lat    lon
1 010950302024 14969994 15040133 34.42668 -86.2437
2 010950306002  6751877 16610261 34.31763 -86.34399
```

```
# Now, for Census 2020 blocks, create pop wtd centroids lat lon for each block group ####
# using EJAMblockdata::blockwts and EJAMblockdata::blockpoints
```

```
bgpts_blocks <- copy(blockpoints) # not essential but ok to make sure we do not change blockpoints i
# all.equal(bgpts$blockid , blockwts$blockid)
bgpts_blocks[ , bgid := blockwts$bgid]
bgpts_blocks[ , blockwt := blockwts$blockwt]
# get pop wtd mean of lat, and same for lon, by bgid
bgpts <- bgpts_blocks[ , lapply(.SD, FUN = function(x) stats::weighted.mean(x, w = blockwt, na.rm =
rm( bgpts_blocks)
# add the bgfips column, so it has bgfips, bgid, lat, lon
# all.equal(bgpts$bgid,bgid2fips$bgid)
bgpts[ , bgfips := bgid2fips$bgfips]
# setnames(bgpts, 'bgfips', 'FIPS')
```

```
# BUT NOTE this census2020 block table has PR but lacks "AS" "GU" "MP" "VI" ####
# > uniqueN(EJAMblockdata::blockid2fips[,substr(blockfips,1,2)])
# [1] 52
# length(unique(EJAMEjscreendata::EJSCREEN_Full_with_AS_CNMI_GU_VI$ST_ABBREV))
# [1] 56
# dim(bgejam)
# [1] 242,940    155
# dim(bg22)
# [1] 242,335    157
```

```
#
# so how do we get latlon for bg in as/gu/mp/vi ? #####

# view those block group points on a map (plot only a subset which is enough)
sam <- sample(seq_along(bgpts$bgid),5000)
plot(bgpts$lon[sam], bgpts$lat[sam], pch = '.')

# view one state, florida, where 12 are the 1st 2 digits of the FIPS:
# bgpts[bgid2fips[substr(bgfips,1,2) == '12', ], on = 'bgid']
xx='12'
mystate <- bgpts[bgid2fips[substr(bgfips, 1, 2) == xx, ], on = 'bgid'][, .(lon,lat)]
plot(mystate, pch = '.')
rm(mystate, xx)
```

How blockcounts were done:

```
library(EJAMblockdata)
library(data.table)
bg_blockcounts <- blockwts[, .(blockcount = uniqueN(.SD)), by=bgid]
sum(bg_blockcounts$blockcount == 1)
# [1] 1874 blockgroups have only 1 block
sum(bg_blockcounts$blockcount == 1000) the max is 1000 blocks in a bg
# # [1] 22
round(100*table(bg_blockcounts[blockcount <20, blockcount]) / nrow(bg_blockcounts) ,1)
# about 1 to 3
# 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19
# 0.8 1.2 1.3 1.4 1.5 2.1 2.2 2.4 2.6 2.8 2.8 3.0 3.0 2.9 3.0 2.9 2.8 2.7 2.5
all.equal(bgpts$bgid, bg_blockcounts$bgid)
bgpts[, blockcount := bg_blockcounts$blockcount]
dim(bgpts)
# 242335 x 5
usethis::use_data(bgpts) # saved for EJAM package
```

bg_cenpop2020

data.table with all US Census 2020 block groups, Census 2020 population count, and lat/lon of Census2020-population-weighted centroid of block group

Description

data.table with all US Census 2020 block groups, Census 2020 population count, and lat/lon of Census2020-population-weighted centroid of block group

Details

also see `attributes(bg_cenpop2020)` for source URL and date

See Also

[bgpts blockgroupstats](#)

bg_from_county	<i>Analyze US Counties as if they were sites, to get EJ indicators summary for each county</i>
----------------	--

Description

Analyze US Counties as if they were sites, to get EJ indicators summary for each county

Usage

```
bg_from_county(fips)
```

Arguments

fips County FIPS vector as character not numeric values

Value

data.table with all pairs of county fips - bgid, and a unique siteid assigned to each county

Examples

```
bg_from_county(c('01001','72153'))
# Largest US Counties by ACS Population Totals:
blockgroupstats[ , .(ST = ST[1], countypop = sum(pop)),
  by=.(FIPS = substr(bgfips,1,5))[order(-countypop),][1:20, .(
    CountyPopulation = prettyNum(countypop, big.mark = ","), FIPS, ST)]
```

blockgroupstats	<i>EJScreen demographic and enviromental indicators for Census block groups</i>
-----------------	---

Description

The EJScreen dataset (demographic, environmental, EJ indicators), plus more demographic sub-groups.

Details

As of 10/2022 it is the EJScreen 2.1 version of data, which uses ACS 2016-2020. EJScreen 2.1 was released October 2022. As of 4/2022 it was the EJScreen 2.0 version of data, which used ACS 2015-2019. EJScreen 2.0 was released 2/18/2022 (raw data download avail 2/22/2022).

NOTE: It also has the race/ethnic subgroups that add up to minority or people of color.

Each year this could be created as for the latest version. See `attributes(blockgroupstats)` It is also available in a similar form via the `ejscreen` package on github, and `EJAMejscreendata::EJSCREEN_Full_with_AS_CN` but there are differences in which columns are kept.

It is a `data.table` of US Census blockgroups (not blocks). With PR, 242,335 rows, approx 175 columns. See <https://www.epa.gov/ejscreen>

column names include `bgfips`, `bgid` (for join to `blockwt$bgid`), `pop`, `pctlowinc`, `pcthis`, etc.

see source code and notes in `EJAM/inst/notes_datasets/` which has `create_blockgroupstats()`

See maybe the notes on cleaning up and changing the dataset starting from `ejscreen::bg22plus`

<code>counties_as_sites</code>	<i>Analyze US Counties as if they were sites, to get EJ indicators summary for each county</i>
--------------------------------	--

Description

Analyze US Counties as if they were sites, to get EJ indicators summary for each county

Usage

```
counties_as_sites(fips)
```

Arguments

<code>fips</code>	County FIPS vector as character not numeric values
-------------------	--

Value

`data.table` with all pairs of county fips - bgid, and a unique siteid assigned to each county

Examples

```
bg_from_county(c('01001', '72153'))
# Largest US Counties by ACS Population Totals:
blockgroupstats[, .(ST = ST[1], countypop = sum(pop)),
  by=.(FIPS = substr(bgfips,1,5))[order(-countypop),][1:20, .(
  CountyPopulation = prettyNum(countypop, big.mark = ","), FIPS, ST)]
```

datapack	<i>See info about the data sets in one or more packages Wrapper for data() - just shows info in console and silently returns a data.frame</i>
----------	---

Description

See info about the data sets in one or more packages Wrapper for data() - just shows info in console and silently returns a data.frame

Usage

```
datapack(pkg = ejampackages, len = 30)
```

Arguments

pkg	a character vector giving the package(s) to look in for data sets
len	Only affects what is printed to console - specifies the number of characters to limit Title to, making it easier to see in the console.

Value

data.frame with Item and Title as columns

Examples

```
datapack("datasets")
datapack("MASS")
datapack(ejampackages)
```

distance_by_group	<i>distance_by_group</i> Get average distance for ONE demographic group versus everyone else
-------------------	--

Description

distance_by_group Get average distance for ONE demographic group versus everyone else

Usage

```
distance_by_group(
  results_bybg_people,
  demogvarname = "Demog.Index",
  demoglabel = demogvarname
)
```

Arguments

results_bybg_people	data.table from doaggregate()\$results_bybg_people
demogvarname	e.g., "pctlowinc"
demoglabel	e.g., "Low Income Residents"

Details

Note on Avg Distance and range of distances in each Demog group, & %D as function of distance:

We have info on each blockgroup near each site, which means some small % of those bgs are duplicated in this table:

```
results_bybg_people
```

Mostly we want overall (not by site) to know avg and cum distrib of distances in each demog, (and also %D as a function of continuous distance),

and for those stats we would want to take only unique blockgroups from here, using the shorter distance, so the distribution of distances does not doublecount people.

But we might also want to see that distribution of distances by D for just 1 site?

And we might also want to see the %D as a function of continuous distance at just 1 site?

So to retain flexibility doaggregate() reports all instances of blockgroup-site pairings.

Value

list of 2 numbers: avg_distance_for_group and avg_distance_for_nongroup

See Also

[distance_by_group_plot\(\)](#) [distance_by_groups\(\)](#)

distance_cdf_by_group_plot

distance_cdf_by_group_plot - SLOW - needs to be optimized Plot a graphic showing cumulative shares of ONE demographic group that are within each distance

Description

distance_cdf_by_group_plot - SLOW - needs to be optimized Plot a graphic showing cumulative shares of ONE demographic group that are within each distance

Usage

```
distance_cdf_by_group_plot(
  results_bybg_people,
  radius_miles = round(max(x$distance_min_avgperson, na.rm = T), 1),
  demogvarname = "Demog.Index",
  demoglabel = demogvarname,
  color1 = "red",
  color2 = "black"
)
```

Arguments

results_bybg_people	data.table from doaggregate()\$results_bybg_people
radius_miles	miles radius that was max distance analyzed
demogvarname	name of column in results_bybg_people, e.g., "pctlowinc"
demoglabel	friendly text name for labelling graphic, like "Low income residents"
color1	color like "red" for demographic group of interest
color2	color like "gray" for everyone else

Value

invisibly returns full table of sorted distances of blockgroups, cumulative count of demog group at that block group's distance, and cumulative count of everyone else in that block group

See Also

[distance_by_group\(\)](#) [getblocksnearbyviaQuadTree\(\)](#) for examples

distance_via_surfacdistance
convert surface distance to actual distance

Description

```
\preformatted{
  Just a simple formula:
  earthRadius_miles <- 3959
  angle_rad <- x/earthRadius_miles
  # Calculate radius * cord length
  return( earthRadius_miles * 2*sin(angle_rad/2) )
}
```

Usage

```
distance_via_surfacdistance(x)
```

Arguments

x	surface distance in miles
---	---------------------------

doaggregate	<i>Summarize indicators in each buffer (given the blocks in each buffer and indicators for each block)</i>
-------------	--

Description

This updated 2023 code takes a set of facilities and the set of blocks that are near each, (as identified previously, in other code that has identified which blocks are nearby) and combines those with indicator scores for block groups.

Usage

```
doaggregate(
  sites2blocks,
  sites2states_or_latlon = NA,
  countcols = NULL,
  popmeancols = NULL,
  calculatedcols = NULL,
  testing = FALSE,
  include_ejindexes = FALSE,
  updateProgress = NULL,
  ...
)
```

Arguments

sites2blocks	data.table of distances in miles between all sites (facilities) and nearby Census block internal points, with columns siteid, blockid, distance, created by get-blocks-nearby function. See sites2blocks_example dataset in package, as input to this function
sites2states_or_latlon	data.table or just data.frame, with columns siteid (each unique one in sites2blocks) and ST (2-character State abbreviation) or lat and lon
countcols	character vector of names of variables to aggregate within a buffer using a sum of counts, like, for example, the number of people for whom a poverty ratio is known, the count of which is the exact denominator needed to correctly calculate percent low income.
popmeancols	character vector of names of variables to aggregate within a buffer using population weighted mean.
calculatedcols	character vector of names of variables to aggregate within a buffer using formulas that have to be specified.
testing	used while testing this function
include_ejindexes	not yet implemented
updateProgress	progress bar function used for shiny app
...	more to pass to another function? Not used currently.

Details

For all examples, see [getblocksnearbyviaQuadTree\(\)](#)

This function aggregates the blockgroup scores to create a summary of each indicator, as a raw score and US percentile and State percentile, in each buffer (i.e., near each facility):

- **SUMS OF COUNTS:** for population count, or number of households or Hispanics, etc.
- **POPULATION-WEIGHTED MEANS:** for Environmental indicators.
EJ Indexes: These could be in theory recalculated via formula, but the way EJScreen does this is apparently finding the pop wtd mean of EJ Index raw scores, not the EJ Index formula applied to the summarized demographic score and aggregated envt number.
- **CALCULATED BY FORMULA:** Buffer or overall score calculated via formulas using aggregated counts, such as percent low income = sum of counts low income / sum of counts of denominator, which in this case is the count of those for whom the poverty ratio is known.
- **LOOKED UP:** Aggregated scores are converted into percentile terms via lookup tables (US or State version).

This function requires the following as data lazy loaded for example from EJAMblockdata package:

- blockwts: data.table with these columns: blockid , bgid, blockwt
- quaddata, and blockquadtree: data.table and quad tree, for indexes of block points (and local-tree that is created when package is loaded)
- EJAM::blockgroupstats - A data.table (such as EJScreen demographic and environmental data by blockgroup?)

See Also

[ejamit getblocksnearby\(\)](#)

dupenames	<i>helper function to look at several packages to spot conflicting exported names See what objects (functions or data) are exported by a given (installed) package</i>
-----------	--

Description

helper function to look at several packages to spot conflicting exported names See what objects (functions or data) are exported by a given (installed) package

Usage

```
dupenames(
  pkg = EJAM::ejampackages,
  sortbypkg = FALSE,
  compare.functions = TRUE
)
```

Arguments

pkg	one or more package names as vector of strings. If "all" it checks all installed pkgs, but takes very very long potentially.
sortbypkg	If TRUE, just returns same thing but sorted by package name
compare.functions	If TRUE, sends to console inf about whether body and formals of the functions are identical between functions of same name from different packages. Only checks the first 2 copies, not any additional ones (where 3+ pkgs use same name)

Details

This can help find duplicates/conflicts within source code and make sure they are on search path, for when renaming / moving functions/packages

Value

data.frame with columns Package, Object name (or NA if no dupes)

See Also

[all.equal_functions\(\)](#)

ejamit	<i>Get complete EJ analysis numbers (demographic and environmental indicators) for a list of locations</i>
--------	--

Description

This is the main function in EJAM for users who want to use EJAM from RStudio. See `help("EJAM")`

Usage

```
ejamit(
  sitepoints,
  cutoff = 3,
  maxcutoff = 31.07,
  avoidorphans = TRUE,
  quadtree,
  ...
)
```

Arguments

sitepoints	see getblocksnearbyviaQuadTree() or other such functions
cutoff	see getblocksnearbyviaQuadTree() or other such functions
maxcutoff	see getblocksnearbyviaQuadTree() or other such functions
avoidorphans	see getblocksnearbyviaQuadTree() or other such functions
quadtree	User does not need to provide this parameter. Already created when EJAM package is loaded, this is a large quadtree object created from the SearchTree package example: <code>SearchTrees::createTree(EJAMblockdata::quaddata, treeType = "quad", dataType = "point")</code>
...	see getblocksnearbyviaQuadTree_Clustered() or other such functions

See Also

[getblocksnearby\(\)](#) [doaggregate\(\)](#)

Examples

```
## Not run:

# All in one step, using functions not shiny app:
out <- ejamit(testpoints_100_dt, 2, quadtree=localtree)

# do not specify sitepoints and it will prompt you for a file,
# if in RStudio in interactive mode!
out <- ejamit(cutoff = 3)

# Specify facilities or sites as points for test data,
# use 1000 test facility points from the R package
testsites <- testpoints_1000_dt
# use facility points in an excel or csv file
testsites <- latlon_from_anything(
  "./inst/testdata/testpoints_207_sites_with_signif_violations_NAICS_326_ECHO.csv")
# use facility points from a random sample of EPA-regulated facilities
testsites <- EJAMfrsdata::frs[sample(1:nrow(EJAMfrsdata::frs), 1e3),] # this is slow

# Specify max distance from sites to look at (residents within X miles of site point)
radius <- 3.1 # miles

# Get summaries of all indicators near a set of points
out <- ejamit(testsites, radius)
# out <- ejamit("myfile.xlsx", 3.1)

# out2 <- EJAMejscreenapi::ejscreenit(testpoints_1000[1:3,])

# View results overall
round(t(out$results_overall), 3.1)

# View plots
# plot_distance_avg_by_group(out)
# plot_distance_cdf_by_group(out)

# View maps
mapfast(out$results_bysite, radius = 3.1)

# view results at a single site
t(out$results_bysite[1, ])
t(out$results_bysite[out$results_bysite$siteid == 2, ])

# if doing just 1st step of ejamit()
# get distance between each site and every nearby Census block
s2b <- testdata_sites2blocks
s2b <- getblocksnearby(testsites, cutoff = radius)
s2b <- getblocksnearbyviaQuadTree(testsites, cutoff = radius)
summarize_blockcount(s2b)

# if doing just 2d step of ejamit()
# get summaries of all indicators based on table of distances
out <- doaggregate(s2b, testsites) # this works now and is simpler
```

```
# if localtree had to be built from block point data again
localtree_example = SearchTrees::createTree(EJAMblockdata::quaddata, treeType = "quad", dataType = "point")

## End(Not run)
```

ejumpackages	<i>list of names of EJAM-related R packages</i>
--------------	---

Description

list of names of EJAM-related R packages

fipsbg_from_anyfips	<i>fipsbg_from_anyfips convert any FIPS codes to the FIPS of all the blockgroups that are among or within or containing those FIPS This is a way to get a list of blockgroups, specified by state/county/tract or even block. Takes a vector of one or more FIPS that could be State (2-digit), County (5-digit), Tract (11-digit), or blockgroup (12 digit), or even block (15-digit fips). Returns unique vector of FIPS of all US blockgroups (including DC and Puerto Rico) that contain any specified blocks, are equal to any specified blockgroup fips, or are contained within any provided tract/county/state FIPS. <code>stateinfo\$STmatch(unique(substr(blockgroupstats\$bgfips,1,2)), stateinfo\$FIPS.ST)</code></i>
---------------------	--

Description

fipsbg_from_anyfips convert any FIPS codes to the FIPS of all the blockgroups that are among or within or containing those FIPS This is a way to get a list of blockgroups, specified by state/county/tract or even block. Takes a vector of one or more FIPS that could be State (2-digit), County (5-digit), Tract (11-digit), or blockgroup (12 digit), or even block (15-digit fips). Returns unique vector of FIPS of all US blockgroups (including DC and Puerto Rico) that contain any specified blocks, are equal to any specified blockgroup fips, or are contained within any provided tract/county/state FIPS. `stateinfo$STmatch(unique(substr(blockgroupstats$bgfips,1,2)), stateinfo$FIPS.ST)`

Usage

```
fipsbg_from_anyfips(fips)
```

Arguments

fips	vector of US FIPS codes, as character or numeric, with or without their leading zeroes, each with as many characters
------	--

Value

vector of blockgroup FIPS (or NA values) that may be much longer than the vector of fips passed to this function.

See Also

[fips_lead_zero\(\)](#)

Examples

```
# all blockgroups in one state
blockgroupstats[,.N,by=substr(bgfips,1,2)]
length(fipsbg_from_anyfips("72"))
# all blockgroups in this one county
fipsbg_from_anyfips(30001)
# all blockgroups that contain any of these 6 blocks (just one bg)
fipsbg_from_anyfips( blockid2fips$blockfips[1:6])
# 2 counties
fipsbg_from_anyfips(c(36009,36011))
```

fips_lead_zero	<i>fips_lead_zero Add leading zeroes to fips codes if missing, replace with NA if length invalid Note it does NOT VALIDATE FIPS - It does NOT check if FIPS is valid other than checking its length seems OK, i.e., it might be a state, county, tract, blockgroup, or block FIPS code.</i>
----------------	---

Description

fips_lead_zero Add leading zeroes to fips codes if missing, replace with NA if length invalid Note it does NOT VALIDATE FIPS - It does NOT check if FIPS is valid other than checking its length seems OK, i.e., it might be a state, county, tract, blockgroup, or block FIPS code.

Usage

```
fips_lead_zero(fips)
```

Arguments

fips vector of numeric or character US FIPS codes

Value

vector of same length

Examples

```
fips_lead_zero(c(1,"01",1234,"1234","12345",123456))
```

format_gt_table	<i>format_gt_table</i>
-----------------	------------------------

Description

format_gt_table

Usage

```
format_gt_table(
  df,
  type,
  my_cell_color = "#dce6f0",
  my_border_color = "#0070c0"
)
```

Arguments

df,	a data frame with 6 columns (var_names, value, state_avg, state_pctile, usa_avg, usa_pctile), and one row per indicator
type,	string - one of 'demog', 'envt'
my_cell_color,	color for filling in background of table cells, can be given as string ('blue') or hex code ('#0070c0')
my_border_color,	color for table borders and boundaries, can be given as string ('blue') or hex code ('#0070c0')

Value

a 'gt'-style table with formatting to closely match EJScreen standard report formatting

frs_from_naics	<i>Use NAICS code or industry title text search to see FRS facility registry services data on those EPA-regulated sites</i>
----------------	---

Description

Use NAICS code or industry title text search to see FRS facility registry services data on those EPA-regulated sites

Usage

```
frs_from_naics(naics_code_or_name, ...)
```

Arguments

naics_code_or_name	
...	passed to naics_from_any()

Value

relevant rows of the data.table called `frs`, which has column names that are "lat" "lon" "REGISTRY_ID" "PRIMARY_NAME" "NAICS" "PGM_SYS_ACRNMS"

See Also

`siteid_from_naics()` `naics_from_any()`

Examples

```
frs_from_naics("uranium")
mapfast(frs_from_naics(naics_from_any("nuclear")$code))
naics_from_any("silver")
naics_from_name("silver")
naics_from_any(212222 )
frs_from_naics(212222)
siteid_from_naics(212222)
latlon_from_naics(212222)
```

<code>frs_from_program</code>	<i>Use EPA Program acronym like TRIS to see FRS facility registry services data on those EPA-regulated sites</i>
-------------------------------	--

Description

Get data.table based on given FRS Program System CATEGORY. Find all FRS sites in a program like RCRAINFO, TRIS, or others.

Usage

```
frs_from_program(program)
```

Arguments

`program` vector of one or more EPA Program names used by FRS

Value

relevant rows of the data.table called `frs`, which has column names that are "lat" "lon" "REGISTRY_ID" "PRIMARY_NAME" "NAICS" "PGM_SYS_ACRNMS"

Examples

```
mapfast(latlon_from_program("CAMDBS"))
#p1="TRIS"; p2="EIS";
#analyze.stuff::overlaps(
#  latlon_from_program(p1)$REGISTRY_ID,
#  latlon_from_program(p2)$REGISTRY_ID,
#  ab_names = c(p1,p2)
#)
```

frs_from_programid	<i>Use EPA Program ID to see FRS facility registry services data on those EPA-regulated sites</i>
--------------------	---

Description

Use EPA Program ID to see FRS facility registry services data on those EPA-regulated sites

Usage

```
frs_from_programid(programid)
```

Arguments

siteid vector of one or more EPA Program ID codes used by FRS

Value

relevant rows of the data.table called [frs](#), which has column names that are "lat" "lon" "REGISTRY_ID" "PRIMARY_NAME" "NAICS" "PGM_SYS_ACRNMS"

Examples

```
x=frs_from_programid(testids_program_sys_id)
x
mapfast(x)
```

frs_from_regid	<i>Use registry ID to see FRS facility registry services data on those EPA-regulated sites</i>
----------------	--

Description

Use registry ID to see FRS facility registry services data on those EPA-regulated sites

Usage

```
frs_from_regid(siteid)
```

Arguments

siteid vector of one or more EPA Registry ID codes used by FRS

Value

relevant rows of the data.table called [frs](#), which has column names that are "lat" "lon" "REGISTRY_ID" "PRIMARY_NAME" "NAICS" "PGM_SYS_ACRNMS"

Examples

```
frs_from_siteid(testids_registry_id)
```

frs_from_siteid	<i>Use registry ID to see FRS facility registry services data on those EPA-regulated sites</i>
-----------------	--

Description

Use registry ID to see FRS facility registry services data on those EPA-regulated sites

Usage

```
frs_from_siteid(siteid)
```

Arguments

siteid	vector of one or more EPA Registry ID codes used by FRS
--------	---

Value

relevant rows of the data.table called [frs](#), which has column names that are "lat" "lon" "REG-ISTRY_ID" "PRIMARY_NAME" "NAICS" "PGM_SYS_ACRNMS"

Examples

```
frs_from_siteid(testids_registry_id)
```

frs_from_sitename	<i>Use site name text search to see FRS facility registry services data on those EPA-regulated sites VERY SLOW search within PRIMARY_NAME of facilities for matching text</i>
-------------------	---

Description

Use site name text search to see FRS facility registry services data on those EPA-regulated sites VERY SLOW search within PRIMARY_NAME of facilities for matching text

Usage

```
frs_from_sitename(sitenames, ignore.case = TRUE, fixed = FALSE)
```

Arguments

sitenames	one or more strings in a vector, which can be regular expressions or query for exact match using fixed=TRUE
ignore.case	logical, search is not case sensitive by default (unlike grepl() default)
fixed	see grepl() , if set to TRUE it looks for only exact matches

Value

relevant rows of the data.table called [frs](#), which has column names that are "lat" "lon" "REG-ISTRY_ID" "PRIMARY_NAME" "NAICS" "PGM_SYS_ACRNMS"

Examples

```
## Not run:
# very slow
x=frs_from_sitename
nrow(x)
head(x)

## End(Not run)
```

frs_is_valid	<i>Validate FRS uploads</i>
--------------	-----------------------------

Description

Check for proper FRS facility id in uploaded data

Usage

```
frs_is_valid(frs_upload)
```

Arguments

frs_upload upload frs converted to data frame

Value

boolean value (valid or not valid)

getblocksnearby	<i>Fast way to find nearby points - finds distance to each Census block centroid nearby</i>
-----------------	---

Description

Given a set of points and a specified radius (cutoff), this function quickly finds all the US Census blocks near each point. For each point, it uses the specified cutoff distance and finds the distance to every block within the circle defined by the radius (cutoff). Each block is defined by its Census-provided internal point, by latitude and longitude.

Each point can be the location of a regulated facility or other type of site, and the blocks are a high-resolution source of information about where residents live.

Finding which blocks have their internal points in a circle provides a way to quickly estimate what fraction of a block group is inside the circular buffer more accurately and more quickly than areal apportionment of block groups would provide.

Usage

```
getblocksnearby(
  sitepoints,
  cutoff = 3,
  maxcutoff = 31.07,
  avoidorphans = TRUE,
  quadtree,
  parallel = FALSE,
  ...
)
```

Arguments

sitepoints	data.table with columns siteid, lat, lon giving point locations of sites or facilities around which are circular buffers
cutoff	miles radius, defining circular buffer around site point
maxcutoff	miles distance (max distance to check if not even 1 block point is within cutoff)
avoidorphans	logical Whether to avoid case where no block points are within cutoff, so if TRUE, it keeps looking past cutoff to find nearest one within maxcutoff.
quadtree	(a pointer to the large quadtree object) created from the SearchTree package example: SearchTrees::createTree(EJAMblockdata::quaddata, treeType = "quad", dataType = "point") Takes about 2-5 seconds to create this each time it is needed. It is automatically created when the package is loaded via the .onLoad() function
...	passed to getblocksnearbyviaQuadTree() or other such functions

Details

For all examples, see [ejamit\(\)](#) `getblocksnearby()` is a wrapper redirecting to the right version like [getblocksnearbyviaQuadTree\(\)](#) Census block "internal points" are actually what it looks for, like a centroid. The blocks are pre-indexed for the whole USA, via the data object quadtree or localtree

See Also

[ejamit\(\)](#) [getblocksnearbyviaQuadTree\(\)](#) [getblocksnearbyviaQuadTree_Clustered\(\)](#) [getblocksnearbyviaQuadTree_Clustered_parallel\(\)](#)

getblocksnearby2	<i>Key buffering function - wrapper redirecting to the right version of getblocksnearby()</i>
------------------	---

Description

Key buffering function - wrapper redirecting to the right version of `getblocksnearby()`

Usage

```
getblocksnearby2(
  sitepoints,
  cutoff = 3,
  maxcutoff = 31.07,
  avoidorphans = TRUE,
  quadtree = is.null,
  ...
)
```

Arguments

sitepoints	see getblocksnearbyviaQuadTree() or other such functions
cutoff	see getblocksnearbyviaQuadTree() or other such functions
maxcutoff	see getblocksnearbyviaQuadTree() or other such functions
avoidorphans	see getblocksnearbyviaQuadTree() or other such functions
quadtree	a large quadtree object created from the SearchTree package example: SearchTrees::createTree(EJAM, treeType = "quad", dataType = "point")
...	see getblocksnearbyviaQuadTree_Clustered() or other such functions

Details

For all examples, see [ejamit\(\)](#)

Like [getblocksnearby\(\)](#) but tries to handle localtree and quadtree parameter differently

- not sure how to check if they are in the right environment.

See Also

[getblocksnearby_and_doaggregate\(\)](#) [getblocksnearby\(\)](#) [getblocksnearbyviaQuadTree\(\)](#)
[getblocksnearbyviaQuadTree_Clustered\(\)](#) [getblocksnearbyviaQuadTree2\(\)](#)

getblocksnearbyviaQuadTree

Find nearby blocks using Quad Tree data structure for speed, NO PARALLEL PROCESSING

Description

Given a set of points and a specified radius (cutoff), this function quickly finds all the US Census blocks near each point. For each point, it uses the specified cutoff distance and finds the distance to every block within the circle defined by the radius (cutoff). Each block is defined by its Census-provided internal point, by latitude and longitude.

Each point can be the location of a regulated facility or other type of site, and the blocks are a high-resolution source of information about where residents live.

Finding which blocks have their internal points in a circle provides a way to quickly estimate what fraction of a block group is inside the circular buffer more accurately and more quickly than areal apportionment of block groups would provide.

Usage

```
getblocksnearbyviaQuadTree(
  sitepoints,
  cutoff = 3,
  maxcutoff = 31.07,
  avoidorphans = TRUE,
  report_progress_every_n = 500,
  quadtree
)
```

Arguments

sitepoints	data.table with columns siteid, lat, lon giving point locations of sites or facilities around which are circular buffers
cutoff	miles radius, defining circular buffer around site point
maxcutoff	miles distance (max distance to check if not even 1 block point is within cutoff)
avoidorphans	logical Whether to avoid case where no block points are within cutoff, so if TRUE, it keeps looking past cutoff to find nearest one within maxcutoff.
report_progress_every_n	Reports progress to console after every n points, mostly for testing, but a progress bar feature might be useful unless this is super fast.
quadtree	(a pointer to the large quadtree object) created from the SearchTree package example: SearchTrees::createTree(EJAMblockdata::quaddata, treeType = "quad", dataType = "point") Takes about 2-5 seconds to create this each time it is needed. It is automatically created when the package is loaded via the <code>.onLoad()</code> function

See Also

[ejamit\(\)](#) [getblocksnearby\(\)](#) [getblocksnearbyviaQuadTree\(\)](#) [getblocksnearbyviaQuadTree_Clustered\(\)](#)
[getblocksnearbyviaQuadTree2\(\)](#)

getblocksnearbyviaQuadTree2

Find nearby blocks using Quad Tree data structure for speed, NO PARALLEL PROCESSING

Description

This should be almost identical to `getblocksnearbyviaQuadTree()`, but it uses `f2`, a copy of `sitepoints`, and more importantly it pulls some code out of the for loop and uses a vectorized approach. Given a set of points and a specified radius (`cutoff`), this function quickly finds all the US Census blocks near each point. For each point, it uses the specified cutoff distance and finds the distance to every block within the circle defined by the radius (`cutoff`). Each block is defined by its Census-provided internal point, by latitude and longitude.

Each point can be the location of a regulated facility or other type of site, and the blocks are a high-resolution source of information about where residents live.

Finding which blocks have their internal points in a circle provides a way to quickly estimate what fraction of a block group is inside the circular buffer more accurately and more quickly than areal apportionment of block groups would provide.

Usage

```
getblocksnearbyviaQuadTree2(
  sitepoints,
  cutoff = 3,
  maxcutoff = 31.07,
  avoidorphans = TRUE,
  report_progress_every_n = 500,
  quadtree
)
```

Arguments

sitepoints	data.table with columns siteid, lat, lon giving point locations of sites or facilities around which are circular buffers
cutoff	miles radius, defining circular buffer around site point
maxcutoff	miles distance (max distance to check if not even 1 block point is within cutoff)
avoidorphans	logical Whether to avoid case where no block points are within cutoff, so if TRUE, it keeps looking past cutoff to find nearest one within maxcutoff.
report_progress_every_n	Reports progress to console after every n points, mostly for testing, but a progress bar feature might be useful unless this is super fast.
quadtree	(a pointer to the large quadtree object) created from the SearchTree package example: SearchTrees::createTree(EJAMblockdata::quaddata, treeType = "quad", dataType = "point") Takes about 2-5 seconds to create this each time it is needed. It can be automatically created when the package is loaded via the .onLoad() function

See Also

[getblocksnearbyviaQuadTree_Clustered\(\)](#) [getblocksnearbyviaQuadTree\(\)](#)

Examples

```
localtree_example = SearchTrees::createTree(EJAMblockdata::quaddata, treeType = "quad", dataType = "point")
x = getblocksnearby2(testpoints_1000_dt, quadtree = localtree_example)
```

getblocksnearbyviaQuadTree3

Find nearby blocks using Quad Tree data structure for speed, NO PARALLEL PROCESSING

Description

Given a set of points and a specified radius (cutoff), this function quickly finds all the US Census blocks near each point. For each point, it uses the specified cutoff distance and finds the distance to every block within the circle defined by the radius (cutoff). Each block is defined by its Census-provided internal point, by latitude and longitude.

Each point can be the location of a regulated facility or other type of site, and the blocks are a high-resolution source of information about where residents live.

Finding which blocks have their internal points in a circle provides a way to quickly estimate what fraction of a block group is inside the circular buffer more accurately and more quickly than areal apportionment of block groups would provide.

Usage

```
getblocksnearbyviaQuadTree3(
  sitepoints,
  cutoff = 3,
  maxcutoff = 31.07,
  avoidorphans = TRUE,
  report_progress_every_n = 500,
  quadtree
)
```

Arguments

sitepoints	data.table with columns siteid, lat, lon giving point locations of sites or facilities around which are circular buffers
cutoff	miles radius, defining circular buffer around site point
maxcutoff	miles distance (max distance to check if not even 1 block point is within cutoff)
avoidorphans	logical Whether to avoid case where no block points are within cutoff, so if TRUE, it keeps looking past cutoff to find nearest one within maxcutoff.
report_progress_every_n	Reports progress to console after every n points, mostly for testing, but a progress bar feature might be useful unless this is super fast.
quadtree	(a pointer to the large quadtree object) created from the SearchTree package example: SearchTrees::createTree(EJAMblockdata::quaddata, treeType = "quad", dataType = "point") Takes about 2-5 seconds to create this each time it is needed. It is automatically created when the package is loaded via the <code>.onLoad()</code> function

See Also

[ejamit\(\)](#) [getblocksnearby\(\)](#) [getblocksnearbyviaQuadTree\(\)](#) [getblocksnearbyviaQuadTree_Clustered\(\)](#) [getblocksnearbyviaQuadTree2\(\)](#)

getblocksnearbyviaQuadTree_Clustered

find nearby blocks using Quad Tree data structure for speed, CLUSTERED FOR PARALLEL PROCESSING

Description

Uses packages [parallel](#) and [snow](#). `parallel::makePSOCKcluster` is an enhanced version of `snow::makeSOCKcluster` in package `snow`. It runs Rscript on the specified host(s) to set up a worker process which listens on a socket for expressions to evaluate, and returns the results (as serialized objects).

Usage

```
getblocksnearbyviaQuadTree_Clustered(
  sitepoints,
  cutoff,
  maxcutoff,
  avoidorphans,
  CountCPU = 1,
  quadtree
)
```

Arguments

sitepoints	data.table with columns LAT, LONG
cutoff	miles distance (check what this actually does)
maxcutoff	miles distance (check what this actually does)
avoidorphans	logical
CountCPU	for parallel processing via makeCluster() and doSNOW::registerDoSNOW()
quadtree	index of all US blocks like localtree

Details

For all examples, see [getblocksnearbyviaQuadTree\(\)](#)

Uses indexgridsize and quaddata variables that come from global environment (but should pass to this function rather than assume in global env?)

See Also

[getblocksnearby_and_doaggregate\(\)](#) [getblocksnearby\(\)](#) [getblocksnearbyviaQuadTree\(\)](#)
[getblocksnearbyviaQuadTree_Clustered\(\)](#) [getblocksnearbyviaQuadTree2\(\)](#)

get_blockpoints_in_shape

*find blocks that are in a polygon, using internal point of block - WORK
 IN PROGRESS *****

Description

This is like getblocksnearby() but for a polygonal buffer area instead of a circular buffer.

Usage

```
get_blockpoints_in_shape(polys, addedbuffermiles = 0)
```

Arguments

polys	polygons (FORMAT???)
addedbuffermiles	width of optional buffering to add to edges, in miles

Value

Polygons with joined block points info. *****

get_shape_buffered	<i>add buffer around shape</i>
--------------------	--------------------------------

Description

add buffer around shape

Usage

```
get_shape_buffered(shape, radius, ...)
```

Arguments

shape	spatial object like areas at high risk or areas with facilities to be analyzed
radius	width of buffer to add to shape. (in case dist is a units object, it should be convertible to arc_degree if x has geographic coordinates, and to st_crs(x)\$units otherwise)
...	passed to st_buffer()

Details

Just a wrapper for [sf::st_buffer\(\)](#)

latlon_as.numeric	<i>Strip non-numeric characters from a vector</i>
-------------------	---

Description

Remove all characters other than minus signs, decimal points, and numeric digits

Usage

```
latlon_as.numeric(x)
```

Arguments

x	vector of something that is supposed to be numbers like latitude or longitude and may be a character vector because there were some other characters like tab or space or percent sign or dollar sign
---	---

Details

Useful if latitude or longitude vector has spaces, tabs, etc. CAUTION - Assumes stripping those out and making it numeric will fix whatever problem there was and end result is a valid set of numbers. Inf etc. are turned into NA values. Empty zero length string is turned into NA without warning. NA is left as NA. If anything other than empty or NA could not be interpreted as a number, it returns NA for those and offers a warning.

Value

numeric vector same length as x

See Also

latlon_df_clean() latlon_infer() latlon_is.valid() latlon_as.numeric()

Examples

```
latlon_as.numeric(c("-97.179167000000007", "-94.0533", "-95.152083000000005"))
latlon_as.numeric(~3:3)
latlon_as.numeric(c(1:3, NA))
latlon_as.numeric(c(1, 'asdf'))
latlon_as.numeric(c(1, ''))
latlon_as.numeric(c(1, '', NA))
latlon_as.numeric(c('aword', '$b'))
latlon_as.numeric(c('-10.5%', '<5', '$100'))
latlon_as.numeric(c(Inf, 1))
```

latlon_df_clean

Find and clean up latitude and longitude columns in a data.frame

Description

Utility to identify lat and lon columns, renaming and cleaning them up.

Usage

```
latlon_df_clean(df)
```

Arguments

df data.frame With columns lat and lon or names that can be interpreted as such

Details

Tries to figure out which columns seem to have lat lon values, renames those in the data.frame.
Cleans up lat and lon values (removes extra characters, makes numeric)

Value

Returns the same data.frame but with relevant colnames changed to lat and lon, and invalid lat or lon values cleaned up if possible or else replaced with NA

See Also

Used by [latlon_from_anything\(\)](#). Uses [latlon_infer\(\)](#) [latlon_is.valid\(\)](#) [latlon_as.numeric\(\)](#)

Examples

```
# x <- latlon_df_clean(x)
```

latlon_from_anything *Flexibly get lat/lon from file, data.frame, data.table, or lat/lon vectors*

Description

Try to figure out if user provided latitude / longitude as vectors, data.frame, file, or interactively pick file.

Usage

```
latlon_from_anything(x, y)
```

Arguments

x	If missing and interactive mode in RStudio, prompts user for file. Otherwise, this can be a filename (csv or xlsx, with path), or data.frame/ data.table/ matrix, or vector of longitudes (in which case y must be the latitudes). Note that even though it is called latlon_etc the lon is x and comes before the lat among parameters x,y File or data.frame/data.table/matrix must have columns called lon and lat, or something that can be inferred to be that by latlon_infer()
y	If x is a vector of longitudes, y must be the latitudes. Ignored otherwise.

Details

This function, `latlon_from_anything()`

relies on

`EJAMbatch.summarizer::read_csv_or_xl()` = `EJAMejscreenapi::read_csv_or_xl()` and

`latlon_df_clean()` which in turn uses `latlon_infer()` `latlon_as.numeric()` `latlon_is.valid()`

`EJAMejscreenapi::read_and_clean_points()`

would be the most general / flexible broadest way to get points, but is still work in progress

is similar to what is done by `latlon_from_anything()`

except it also uses these functions:

`EJAMfrsdata::latlon_from_siteid()`

`EJAMfrsdata::latlon_from_programid()` but not `_from_naics()` ?

Value

A data.frame that has at least columns lon and lat (and others if they were in x)

See Also

`EJAMbatch.summarizer::read_csv_or_xl()` `latlon_df_clean()`

Examples

```

if (interactive()) {
pts <- latlon_from_anything()
}
latlon_from_anything(system.file("testdata/Sample12.xlsx",
  package="EJAMejscreenapi"))
latlon_from_anything(system.file("testdata/testpoints_05.csv",
  package="EJAMejscreenapi"))
latlon_from_anything(testpoints_50[1:6,] )
latlon_from_anything(testpoints_50[1:6, c('lat','lon')] )
latlon_from_anything(x=testpoints_50$lon[1:6], y=testpoints_50$lat[1:6] )

```

latlon_infer	<i>guess which columns have lat and lon based on aliases like latitude, FacLat, etc.</i>
--------------	--

Description

guess which columns have lat and lon based on aliases like latitude, FacLat, etc.

Usage

```
latlon_infer(mycolnames)
```

Arguments

mycolnames e.g., colnames(x) where x is a data.frame from read.csv

Value

returns all of mycolnames except replacing the best candidates with lat and lon

See Also

latlon_df_clean() latlon_infer() latlon_is.valid() latlon_as.numeric()

Examples

```

latlon_infer(c('trilat', 'belong', 'belong')) # warns if no alias found,
# but doesnt warn of dupes in other terms, just preferred term.
latlon_infer(c('a', 'LONG', 'Longitude', 'lat')) # only the best alias is converted/used
latlon_infer(c('a', 'LONGITUDE', 'Long', 'Lat')) # only the best alias is converted/used
latlon_infer(c('a', 'longing', 'Lat', 'lat', 'LAT')) # case variants of preferred are
# left alone only if lowercase one is found
latlon_infer(c('LONG', 'long', 'lat')) # case variants of a single alias are
# converted to preferred word (if pref not found), creating dupes! warn!
latlon_infer(c('LONG', 'LONG')) # dupes of an alias are renamed and still are dupes! warn!
latlon_infer(c('lat', 'lat', 'Lon')) # dupes left as dupes but warn!

```

latlon_is.valid	<i>Validate latitudes and longitudes</i>
-----------------	--

Description

Check each latitude and longitude value to see if they are NA or outside expected numeric ranges (based on approx ranges of lat lon seen among block internal points dataset) lat must be between 17.5 and 71.5, and lon must be (between -180 and -65) OR (between 172 and 180)

Usage

```
latlon_is.valid(lat, lon)
```

Arguments

lat	vector of latitudes in decimal degrees
lon	numeric vector of longitudes in decimal degrees, same length

Value

logical vector, one element per lat lon pair (location)

See Also

latlon_df_clean() latlon_infer() latlon_is.valid() latlon_as.numeric()

Examples

```
## Not run:
table(latlon_is.valid(lat = EJAMblockdata::blockpoints$lat, lon = EJAMblockdata::blockpoints$lon))
##      TRUE
## 8,174,955

## End(Not run)
```

lat_aliases, lon_aliases	<i>Synonyms for lat and lon</i>
--------------------------	---------------------------------

Description

lists of synonyms for "latitude" and "longitude" used when guessing which column is what in user-provided tables of coordinates

map_facilities	<i>map_facilities</i>
----------------	-----------------------

Description

make a leaflet map of uploaded points

Usage

```
map_facilities(mypoints, rad = 3, highlight = FALSE, clustered)
```

Arguments

mypoints,	data frame of uploaded points
rad,	a size for drawing each circle (buffer search radius)
highlight,	a logical for whether to highlight overlapping points (defaults to FALSE)
clustered,	a vector of T/F values for each point, indicating if they overlap with another

Value

a leaflet map with circles, circleMarkers, and basic popup

metadata_add	<i>helper function for package to set attributes of a dataset</i>
--------------	---

Description

This can be used annually to update some datasets in a package. It just makes it easier to set a few metadata attributes similarly for a number of data elements, for example, to add new or update existing attributes.

Usage

```
metadata_add(x, metadata)
```

Arguments

x	dataset (or any object) whose metadata you want to update or create
metadata	must be a named list, so that the function can do this for each i: attr(x, which=names(metadata) i) <- metadata [i]

Value

returns x but with new or altered attributes

See Also

```
metadata_check()
```

Examples

```
x <- data.frame(a=1:10,b=1001:1010)
metadata <- list(
  census_version = 2020,
  acs_version = '2016-2020',
  acs_releasedate = '3/17/2022',
  ejsscreen_version = '2.1',
  ejsscreen_releasedate = 'October 2022',
  ejsscreen_pkg_data = 'bg22'
)
x <- metadata_add(x, metadata)
attributes(x)
x <- metadata_add(x, list(status='final'))
attr(x,'status')
```

metadata_check

helper function in updating the package metadata

Description

Quick and dirty helper during development, to check all the attributes of all the data files in relevant packages. It loads unloaded packages as needed, which you might not want it to do, but it is not coded to be able to check attributes without doing that.

Usage

```
metadata_check(
  packages = NULL,
  which = c("census_version", "acs_version", "acs_releasedate", "ACS",
    "ejsscreen_version", "ejsscreen_releasedate", "ejsscreen_pkg_data", "year", "released"),
  loadifnotloaded = TRUE
)
```

Arguments

packages	Optional. e.g. 'EJAMejsscreendata', or can be a vector of character strings, and if not specified, default is to report on all packages with EJ as part of their name, like EJAMblockdata or ejsscreenapi
which	Optional vector (not list) of strings, the attributes. Default is some typical ones used in EJAM-related packages currently.
loadifnotloaded	Optional to control if func should temporarily attach packages not already loaded.

NAICS	<i>named list of all NAICS code numbers and industry name for each</i>
-------	--

Description

named list of all NAICS code numbers and industry name for each

Details

see <https://naics.com>

See Also

[naicstable](#) [naics_from_any\(\)](#) [naics_categories\(\)](#) [NAICS](#)

naics2children	<i>See NAICS codes queried plus all children of any of those Used by naics_find()</i>
----------------	---

Description

See NAICS codes queried plus all children of any of those Used by naics_find()

Usage

```
naics2children(codes, allcodes = EJAM::NAICS)
```

Arguments

codes	vector of numerical or character
allcodes	Optional (already loaded with package) - dataset with all the codes

Details

start with shortest (highest level) codes. since tied for nchar, these branches have zero overlap, so do each. for each of those, get its children = all rows where parentcode == substr(allcodes, 1, nchar(parentcode)) put together list of all codes we want to include so far. now for the next longest set of codes in original list of codes, do same thing. etc. until did it for 5 digit ones to get 6digit children. take the unique(allthat) table(nchar(as.character(NAICS))) 2 3 4 5 6 17 99 311 709 1057

Value

vector of codes and their names

See Also

[naics_find\(\)](#) [NAICS](#)

Examples

```

naics2children(211)
naics_find(211, exactnumber=TRUE)
naics_find(211, exactnumber=TRUE, add_children = TRUE)
NAICS[211][1:3] # wrong
NAICS[NAICS == 211]
NAICS["211 - Oil and Gas Extraction"]

```

naicstable	<i>data.table with all NAICS code numbers and industry name for each</i>
------------	--

Description

data.table with all NAICS code numbers and industry name for each

Details

see <https://naics.com>

See Also

[naics_from_any\(\)](#) [naics_categories\(\)](#) [NAICS](#) [naics_findwebscrape\(\)](#)

naics_categories	<i>See the names of industrial categories and their NAICS code Easy way to list the 2-digit NAICS (17 categories), or other level</i>
------------------	---

Description

See the names of industrial categories and their NAICS code Easy way to list the 2-digit NAICS (17 categories), or other level

Usage

```
naics_categories(digits = 2, dataset = EJAM::NAICS)
```

Arguments

digits default is 2, for 2-digits NAICS, the top level, but could be up to 6.
dataset Should default to the dataset called NAICS, installed with this package. see [NAICS](#) Check attr(NAICS, 'year')

Details

Also see <https://www.naics.com/search/>

There are this many NAICS codes roughly by number of digits in the code:

```
table(nchar(NAICS))
```

```
2 3 4 5 6
```

```
17 99 311 709 1057
```

See <https://www.census.gov/naics/>

See Also

[naics_find NAICS](#)

Examples

```
naics_categories()
```

naics_download	script to download NAICS file with code and name of sector
----------------	--

Description

See source code. Mostly just a short script to get the 2017 or 2022 codes and names. See <'https://www.census.gov/naics/

Usage

```
naics_download(  
  year = 2017,  
  urlpattern = "https://www.census.gov/naics/YYYYNAICS/2-6%20digit_YYYY_Codes.xlsx",  
  destfile = paste0("~/Downloads/", year, "NAICS.xlsx")  
)
```

Arguments

year	which vintage of NAICS codes to use, 2012, 2017, or 2022
urlpattern	full url of xlsx file to use, but with YYYY instead of year
destfile	full path and name of file to save as locally

Value

names list with year as an attribute

naics_find	<i>OBSOLETE- Search for an industrial sector in the list of NAICS codes, see subsectors THIS IS NOW REPLACED WITH CLEARER FUNCTIONS to get naicstable now, i.e., EJAMfrsdata::naics_from_any()</i>
------------	--

Description

OBSOLETE- Search for an industrial sector in the list of NAICS codes, see subsectors THIS IS NOW REPLACED WITH CLEARER FUNCTIONS to get naicstable now, i.e., EJAMfrsdata::naics_from_any()

Usage

```
naics_find(  
  query,  
  add_children = FALSE,  
  naics_dataset = NULL,  
  ignore.case = TRUE,  
  exactnumber = TRUE,  
  search_on_naics_website = FALSE  
)
```


Arguments

query	a single word or phrase such as "chemical manufacturing" or "cement"
add_children	default is FALSE, so it does NOT children (subcategories) of those that match the query.
naics_dataset	Should default to the dataset NAICS, installed with this package. see NAICS
ignore.case	default TRUE, ignoring whether query is upper or lower case
exactnumber	if TRUE, only return the exact match to (each) queried number (NAICS code). Otherwise it can give unexpected results like finding 92811 when you queried 811.
search_on_naics_website	if TRUE (not default), returns URL of webpage at naics.com with info on the sector

Details

Just a utility, quick way to view NAICS industrial sectors that contain queried word or phrase, but can also see all the subcategories within the matching one.

See <https://www.census.gov/naics/> NOTE: By default, this shows NAICS that match the text query, and also can include all the children NAICS even if they do not match based on text query. So it first finds NAICS that match the text (or code) query via `grep()`, and then can also include all subcategories within those categories.

So `naics_find('soap', add_children=TRUE)` shows "325612 - Polish and Other Sanitation Good Manufacturing", and others, not just "3256 - Soap, Cleaning Compound, and Toilet Preparation Manufacturing", because 3256 matches 'soap' and 325612 is a subcategory of 3256.

The format of NAICS, the `naics_dataset`, in this package is `dput(NAICS1:4)c(11 - Agriculture, Forestry, Fishing = 11, 111 - Crop Production = 111, 1111 - Oilseed and Grain Farming = 1111, 11111 - Soybean Farming = 11111)`

See Also

[naics_categories](#) [NAICS](#) [naics_findwebscrape\(\)](#) [get_facility_info_via_ECHO](#) function [naics_url_of_code\(\)](#) [url_naics.com\(\)](#)

Examples

```
naics_find(8111, exactnumber = FALSE)
naics_find(8111, exactnumber = TRUE)
naics_find(8111, exactnumber = TRUE, add_children = TRUE)

naics_find("paper")
naics_find("cement | concrete")
naics_find("pulp", add_children = FALSE)
naics_find("pulp", add_children = TRUE)
naics_find("asdfasdf", add_children = TRUE)
naics_find("asdfasdf", add_children = FALSE)

EJAMfrsdata::frs[EJAMfrsdata::frs$REGISTRY_ID %in% unlist(
  EJAMfrsdata::siteid_from_naics(
    EJAM::naics_find("pulp", add_children = TRUE))[, "REGISTRY_ID"], 1:5]

EJAMejscreenapi::mapfast(EJAMfrsdata::frs[EJAMfrsdata::frs$REGISTRY_ID %in% unlist(
  EJAMfrsdata::siteid_from_naics(EJAM::naics_find("pulp"))[, "REGISTRY_ID"],  ])
```

```

naics_find(211, exactnumber=TRUE)
naics_find(211, exactnumber=TRUE, add_children = TRUE)
naics2children(211)
NAICS[211][1:3] # wrong
NAICS[NAICS == 211]
NAICS["211 - Oil and Gas Extraction"]

```

naics_findwebscrape	<i>for query term, show list of roughly matching NAICS, scraped from web This finds more than just naics_from_any() does, since that needs an exact match but this looks at naics.com website which lists various aliases for a sector.</i>
---------------------	---

Description

for query term, show list of roughly matching NAICS, scraped from web This finds more than just [naics_from_any\(\)](#) does, since that needs an exact match but this looks at naics.com website which lists various aliases for a sector.

Usage

```
naics_findwebscrape(query)
```

Arguments

query	text like "gasoline" or "copper smelting"
-------	---

Value

data.frame of info on what was found, naics and title

See Also

[naics_from_any\(\)](#) [url_naics.com\(\)](#)

Examples

```

# naics_from_any("copper smelting")
# naics_from_any("copper smelting", website_scrape=TRUE)
# browseURL(naics_from_any("copper smelting", website_url=TRUE) )

url_naics.com("copper smelting")
## Not run:
naics_findwebscrape("copper smelting")
browseURL(url_naics.com("copper smelting"))
browseURL(naics_url_of_code(326))

## End(Not run)

```

naics_url_of_code	<i>Get URL for page with info about industry sector(s) by NAICS See naics.com for more information on NAICS codes</i>
-------------------	---

Description

Get URL for page with info about industry sector(s) by NAICS See [naics.com](https://www.naics.com) for more information on NAICS codes

Usage

```
naics_url_of_code(naics)
```

Arguments

naics	vector of one or more NAICS codes, like 11,"31-33",325
-------	--

Value

vector of URLs as strings like <https://www.naics.com/six-digit-naics/?v=2017&code=22>

naics_validation	<i>Validate NAIC uploads</i>
------------------	------------------------------

Description

Validates and prepares echo uploads

Usage

```
naics_validation(NAICS_enter, NAIC_select)
```

Arguments

NAICS	upload validate missing and/or improper inputs
-------	--

Value

boolean value (valid or not valid)

pctile_from_raw_lookup

Find approx wtd percentiles in lookup table that is in memory

Description

This is used with a data.frame that is a lookup table used to convert a raw indicator value to a percentile - US, Region, or State percentile.

Usage

```
pctile_from_raw_lookup(
  myvector,
  varname.in.lookup.table,
  lookup = usastats,
  zone
)
```

Arguments

myvector	Numeric vector, required. Values to look for in the lookup table.
varname.in.lookup.table	Character element, required. Name of column in lookup table to look in to find interval where a given element of myvector values is.
lookup	Either lookup must be specified, or a lookup table called us must already be in memory. This is the lookup table data.frame with a PCTILE column and column whose name is the value of varname.in.lookup.table
zone	Character element (or vector as long as myvector), optional. If specified, must appear in a column called REGION within the lookup table. For example, it could be 'NY' for New York State.

Details

This could be recoded to be more efficient - could use data.table The data.frame lookup table must have a field called "PCTILE" that has quantiles/percentiles and other column(s) with values that fall at those percentiles. EJAM::usastats, EJAM::statstats, EJAM::regionstats are such lookup tables. This function accepts lookup table (or uses one called us if that is in memory), and finds the number in the PCTILE column that corresponds to where a specified value (in myvector) appears in the column called varname.in.lookup.table. The function just looks for where the specified value fits between values in the lookup table and returns the approximate percentile as found in the PCTILE column. If the value is between the cutpoints listed as percentiles 89 and 90, it returns 89, for example. If the value is exactly equal to the cutpoint listed as percentile 90, it returns percentile 90. If the value is less than the cutpoint listed as percentile 0, which should be the minimum value in the dataset, it still returns 0 as the percentile, but with a warning that the value checked was less than the minimum in the dataset.

Value

By default, returns numeric vector length of myvector.

plot_distance_cdf_by_group

*plot_distance_cdf_by_group - SLOW - needs to be optimized CDF
Line Plots of cumulative share of each demographic group, within
each distance Each groups distribution of distances*

Description

plot_distance_cdf_by_group - SLOW - needs to be optimized CDF Line Plots of cumulative share of each demographic group, within each distance Each groups distribution of distances

Usage

```
plot_distance_cdf_by_group(
  results_bybg_people,
  radius_miles = round(max(x$distance_min_avgperson, na.rm = T), 1),
  demogvarname = c(namez$d, namez$d_subgroups),
  demoglabel = NULL,
  colorlist = colors()[1:length(demogvarname)],
  coloroverall = "gray"
)
```

Arguments

results_bybg_people	data.table from doaggregate()\$results_bybg_people
radius_miles	miles radius that was max distance analyzed
demogvarname	names of columns in results_bybg_people, e.g., "pctlowinc"
demoglabel	friendly text names for labelling graphic, like "Low income residents"
colorlist	colors like "red" etc. for the demographic groups of interest
coloroverall	color like "gray" for everyone as a whole

Value

invisibly returns full table of sorted distances of blockgroups, cumulative count of demog groups at that block group's distance

See Also

[distance_by_groups\(\)](#) [ejamit\(\)](#) for examples

```
plot_distance_mean_by_group
```

*plot_distance_mean_by_group Barplot of Average Proximity to sites,
by Demographic Group (vs everyone else)*

Description

Note ratio shown is ratio of distance among others to distance in group, so values below 1 mean the given demographic group lives closer to facilities.

Usage

```
plot_distance_mean_by_group(  
  results_bybg_people,  
  demogvarname = c(namez$d, namez$d_subgroups),  
  demoglabel = NULL,  
  graph = TRUE  
)
```

Arguments

```
results_bybg_people      data.table from doaggregate()$results_bybg_people  
demogvarname             vector of column names like "pctlowinc" etc.  
demoglabel               vector of labels like "Low Income Residents" etc.
```

Value

data.frame with group, ratio, avg_distance_for_group, avg_distance_for_nongroup

See Also

[distance_by_group\(\)](#)
[distance_by_group_plot\(\)](#)

```
popweightedsums
```

Get population weighted sums of indicators

Description

Get population weighted sums of indicators

Usage

```
popweightedsums(data, fieldnames, fieldnames_out, scaling, popname = "POP100")
```

Arguments

data	data.table with demographic and/or environmental data
fieldnames	vector of terms like pctmin, traffic.score, pm, etc.
fieldnames_out	optional, should be same length as fieldnames
scaling	number to multiply raw values by to put in right units like percent 0-100 vs 0.0-1.0
popname	name of column with population counts to use for weighting

```
prep_EJAM_for_excel
```

Format batch results for excel See EJAM code related to this also!

Description

Format batch results for excel See EJAM code related to this also!

Usage

```
prep_EJAM_for_excel(
  df,
  hyperlink_cols = NULL,
  heatmap_colnames = NULL,
  heatmap_cuts = c(80, 90, 95),
  heatmap_colors = c("yellow", "orange", "red")
)
```

Arguments

df	data.frame, table of batch buffer results
hyperlink_cols	vector of names of columns in df to get treated as hyperlinks in excel
heatmap_colnames	vector of names of columns in df to apply conditional formatting to, by coloring like a heatmap.
heatmap_cuts	vector of color names for heatmap bins, same length as heatmap_cuts, where first color is for those \geq 1st cutpoint, but $< 2d$, second color is for those $\geq 2d$ cutpoint but $< 3d$, etc.
heatmap_colors	vector of colors corresponding to cuts

Value

A workbook via openxlsx::writeData() ready to be saved via openxlsx::saveWorkbook()

```
proximity.score.in.miles
```

convert EJScreen proximity scores to miles per site instead of sites per kilometer Shows US percentiles if no arguments used

Description

convert EJScreen proximity scores to miles per site instead of sites per kilometer Shows US percentiles if no arguments used

Usage

```
proximity.score.in.miles(scoresdf = NULL)
```

Arguments

scoresdf	data.frame of simple proximity scores like for tsdf, rmp, npl but not traffic.score or npdes one since those are weighted and not just count per km
----------	---

```
proxistat2
```

*Calculate a proximity score for every blockgroup Indicator of proximity of each blockgroups to some set of facilities or sites. Proximity score is sum of (1/d) where each d is distance of a given site in km, summed over all sites within 5km, as in EJScreen. *** Still need area of each block to fix this func proxistat2()*

Description

Calculate a proximity score for every blockgroup Indicator of proximity of each blockgroups to some set of facilities or sites. Proximity score is sum of (1/d) where each d is distance of a given site in km, summed over all sites within 5km, as in EJScreen.

*** Still need area of each block to fix this func proxistat2()

Usage

```
proxistat2(pts, cutoff = 8.04672, quadtree)
```

Arguments

pts	data.table of lat lon
cutoff	distance max, in miles, default is 5km (8.04672 miles) which is the EJScreen max search range for proximity scores
quadtree	must be localtree from EJAM::

Value

data.table with proximityscore, bgfips, lat, lon, etc.

Examples

```
# pts <- testpoints_50
# x <- proxistat2(pts = pts[1:1000,], quadtree = localtree)
#
# summary(x$proximityscore)
# # analyze.stuff::pctiles(x$proximityscore)
# plot(x$lon, x$lat)
# tops = x$proximityscore > 500 & !is.infinite(x$proximityscore) & !is.na(x$proximityscore)
# points(x$lon[tops], x$lat[tops], col="red")
```

regionstats

data.table of 100 percentiles and means for each EPA Region.

Description

data.table of 100 percentiles and means for each EPA Region (> 1,000 rows) for all the block groups in that zone (e.g., block groups in [blockgroupstats](#)) for a set of indicators such as percent low income. Each column is one indicator (or specifies the percentile).

This should be similar to the lookup tables in the gdb on the FTP site of EJScreen.

run_app

Launch the Shiny Application in RStudio

Description

launch Shiny web app from RStudio

Usage

```
run_app(
  onStart = NULL,
  options = list(),
  enableBookmarking = "server",
  uiPattern = "/",
  ...
)
```

Arguments

onStart	A function that will be called before the app is actually run. This is only needed for shinyAppObj, since in the shinyAppDir case, a global.R file can be used for this purpose.
options	Named options that should be passed to the runApp call (these can be any of the following: "port", "launch.browser", "host", "quiet", "display.mode" and "test.mode"). You can also specify width and height parameters which provide a hint to the embedding environment about the ideal height/width for the app.

enableBookmarking	Can be one of "url", "server", or "disable". The default value, NULL, will respect the setting from any previous calls to <code>enableBookmarking()</code> . See <code>enableBookmarking()</code> for more information on bookmarking your app.
uiPattern	A regular expression that will be applied to each GET request to determine whether the ui should be used to handle the request. Note that the entire request path must match the regular expression in order for the match to be considered successful.
...	arguments to pass to <code>golem_opts</code> . See <code>?golem::get_golem_options</code> for more details.

Details

`app_run_EJAM()` is like `EJAM::run_app()` `app_run_EJAMejscreenapi()` is like `EJAMejscreenapi::run_app()`
`app_run_EJAMbatch.summarizer()` is like `EJAMbatch.summarizer::run_app()`

sitepoints_example	<i>data.table of points as example of sitepoints for EJAM</i>
--------------------	---

Description

data.table of points as example of sitepoints for EJAM

sites2blocks_example	<i>data.table of output of <code>getblocknearby()</code>, each row is a unique site-block-distance</i>
----------------------	--

Description

data.table of output of `getblocknearby()`, each row is a unique site-block-distance

speedtest	<i>speedtest Runs EJAM analysis for several radius values for many site-points, recording how long each step took.</i>
-----------	--

Description

speedtest Runs EJAM analysis for several radius values for many sitepoints, recording how long each step took.

Usage

```
speedtest(
  n = 10,
  sitepoints = NULL,
  weighting = "frs",
  radius = c(1, 3.106856, 5, 10, 20)[1],
  do_batch.summarize = TRUE,
  logging = TRUE,
  logfolder = getwd(),
  logfilename = "log_n_datetime.txt",
  honk_when_ready = TRUE
)
```

Arguments

<code>n</code>	optional, how many random points to test, or set to 0 to interactively pick file of points in RStudio (n is ignored if sitepoints provided)
<code>sitepoints</code>	optional, (use if you do not want random points) data.frame of points or path/file with points, where columns are lat and lon in decimal degrees
<code>weighting</code>	optional, if using random points, how to weight them, such as facilities, people, or blockgroups. see testpoints_n()
<code>radius</code>	optional, one or more radius values in miles to use in creating circular buffers when findings residents nearby each of sitepoints
<code>do_batch.summarize</code>	optional, if want to also do that function
<code>logging</code>	logical optional, whether to save log file with timings of steps
<code>logfolder</code>	optional, name of folder for log file
<code>logfilename</code>	optional, name of log file to go in folder
<code>honk_when_ready</code>	optional, self-explanatory

Value

EJAM results similar to as from the web app [ejamit\(\)](#)

<code>stateinfo</code>	<i>data.frame of state abbreviations and state names (50+DC+PR; not AS, GU, MP, VI, UM)</i>
------------------------	---

Description

52 rows and 5 variables: ST is the 2-letter abbreviation, statename is the State name (and ftpname is the name as used on Census FTP site).

Details

column names: "ST" "statename" "ftpname" "FIPS.ST" "REGION"

Some datasets lack PR. (72) Many datasets lack these: AS, GU, MP, VI (codes "60" "66" "69" "78")
Almost all datasets lack UM. (74)

72 PR	Puerto Rico
66 GU	Guam
69 MP	Northern Mariana Islands
78 VI	U.S. Virgin Islands
74 UM	U.S. Minor Outlying Islands

```
stateinfo <- structure(list( ST = c("AL", "AK", "AZ", "AR", "CA", "CO", "CT", "DE", "DC", "FL",
"GA", "HI", "ID", "IL", "IN", "IA", "KS", "KY", "LA", "ME", "MD", "MA", "MI", "MN", "MS",
"MO", "MT", "NE", "NV", "NH", "NJ", "NM", "NY", "NC", "ND", "OH", "OK", "OR", "PA",
"RI", "SC", "SD", "TN", "TX", "UT", "VT", "VA", "WA", "WV", "WI", "WY", # "AS", "GU",
"MP", "VI" # "UM", ##### U.S. Minor Outlying Islands # "US", "PR"),
```

```
statename = c("Alabama", "Alaska", "Arizona", "Arkansas", "California", "Colorado", "Connecti-
cut", "Delaware", "District of Columbia", "Florida", "Georgia", "Hawaii", "Idaho", "Illinois", "Indi-
ana", "Iowa", "Kansas", "Kentucky", "Louisiana", "Maine", "Maryland", "Massachusetts", "Michi-
gan", "Minnesota", "Mississippi", "Missouri", "Montana", "Nebraska", "Nevada", "New Hamp-
shire", "New Jersey", "New Mexico", "New York", "North Carolina", "North Dakota", "Ohio",
"Oklahoma", "Oregon", "Pennsylvania", "Rhode Island", "South Carolina", "South Dakota", "Ten-
nessee", "Texas", "Utah", "Vermont", "Virginia", "Washington", "West Virginia", "Wisconsin",
"Wyoming", # "American Samoa", "Guam", "Northern Mariana Islands", "U.S. Virgin Islands",
# "U.S. Minor Outlying Islands", # "United States", "Puerto Rico"),
```

```
ftpname = c("Alabama", "Alaska", "Arizona", "Arkansas", "California", "Colorado", "Connecti-
cut", "Delaware", "DistrictOfColumbia", "Florida", "Georgia", "Hawaii", "Idaho", "Illinois", "Indi-
ana", "Iowa", "Kansas", "Kentucky", "Louisiana", "Maine", "Maryland", "Massachusetts", "Michi-
gan", "Minnesota", "Mississippi", "Missouri", "Montana", "Nebraska", "Nevada", "NewHamp-
shire", "NewJersey", "NewMexico", "NewYork", "NorthCarolina", "NorthDakota", "Ohio", "Okla-
homa", "Oregon", "Pennsylvania", "RhodeIsland", "SouthCarolina", "SouthDakota", "Tennessee",
"Texas", "Utah", "Vermont", "Virginia", "Washington", "WestVirginia", "Wisconsin", "Wyoming",
# NA, NA, NA, NA, # NA, ##### U.S. Minor Outlying Islands # "UnitedStates", "PuertoRico"),
```

```
FIPS.ST = c("01", "02", "04", "05", "06", "08", "09", "10", "11", "12", "13", "15", "16", "17", "18",
"19", "20", "21", "22", "23", "24", "25", "26", "27", "28", "29", "30", "31", "32", "33", "34", "35",
"36", "37", "38", "39", "40", "41", "42", "44", "45", "46", "47", "48", "49", "50", "51", "53", "54",
"55", "56", # "60", "66", "69", "78", # "74", ##### U.S. Minor Outlying Islands # NA, ##### US
"72"), REGION = c(4, 10, 9, 6, 9, 8, 1, 3, 3, 4, 4, 9, 10, 5, 5, 7, 7, 4, 6, 1, 3, 1, 5, 5, 4, 7, 8, 7, 9,
1, 2, 6, 2, 4, 8, 5, 6, 10, 3, 1, 4, 8, 4, 6, 8, 1, 3, 10, 3, 5, 8, # NA, NA, NA, NA, # NA, ##### U.S.
Minor Outlying Islands # NA, # US 2) ), row.names = c(NA, -52L), class = "data.frame")
```

statestats

*data.frame of 100 percentiles and means for each US State and PR
and DC.*

Description

data.frame of 100 percentiles and means for each US State and PR and DC (approx 5,300 rows) for all the block groups in that zone (e.g., block groups in [blockgroupstats](#)) for a set of indicators such as percent low income. Each column is one indicator (or specifies the percentile).

This should be similar to the lookup tables in the gdb on the FTP site of EJScreen, except it also has data for the demographic race/ethnicity subgroups. For details on how the table was made, see [/EJAM/data-raw/usastats_subgroups.R](#)

statestats_query	<i>convenient way to see mean, pctiles of Env or Demog indicators from lookup table</i>
------------------	---

Description

convenient way to see mean, pctiles of Env or Demog indicators from lookup table

Usage

```
statestats_query(
  ST = sort(unique(EJAM::statestats$REGION)),
  varnames = c(EJAM::names_e, EJAM::names_d),
  PCTILES = NULL,
  dig = 2
)
```

Arguments

ST	vector of state abbreviations, or USA
varnames	names of columns in lookup table, like "proximity.rmp"
PCTILES	vector of percentiles 0-100 and/or "mean"
dig	digits to round to

Examples

```
## Not run:
## in USA overall, see mean and key percentiles for all demog and envt indicators
usastats_query() # or statestats_query('us') # can say us or US or USA or usa etc.
usastats_query(PCTILES = 'mean')
usastats_means() # same but nicer looking format in console
usastats_means(dig=4)
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_d)]
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_e)]
## in 1 state, see mean and key percentiles for all demog and envt indicators
statestats_query('MD')
## in 1 state, see mean and key percentiles for just demog indicators
statestats_queryd('MD')
## 1 indicator in 1 state, see a few key percentiles and mean
statestats_query('MD','proximity.tsdf')
## mean of 1 indicator for each state
statestats_query(varnames = 'proximity.tsdf')
## using full blockgroup dataset, not lookup tables of percentiles,
```

```

blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_d, names_e)]
## see all total counts (not just US means),
## demographics including subgroups,
## but not environmental indicators.
t(round(EJAMbatch.summarizer::ustotals2(bg = EJAM::blockgroupstats),2))
t(blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_e, names_d)])

## End(Not run)

```

statestats_queryd	<i>convenient way to see mean, pctlles of DEMOG indicators from lookup table</i>
-------------------	--

Description

convenient way to see mean, pctlles of DEMOG indicators from lookup table

Usage

```

statestats_queryd(
  ST = sort(unique(EJAM::statestats$REGION)),
  varnames = EJAM::names_d,
  PCTILES = NULL,
  dig = 2
)

```

Arguments

ST	vector of state abbreviations, or USA
varnames	names of columns in lookup table, like "proximity.rmp"
PCTILES	vector of percentiles 0-100 and/or "mean"
dig	digits to round to

Examples

```

## Not run:
## in USA overall, see mean and key percentiles for all demog and envt indicators
usastats_query() # or statestats_query('us') # can say us or US or USA or usa etc.
usastats_query(PCTILES = 'mean')
usastats_means() # same but nicer looking format in console
usastats_means(dig=4)
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_d)]
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_e)]
## in 1 state, see mean and key percentiles for all demog and envt indicators
statestats_query('MD')
## in 1 state, see mean and key percentiles for just demog indicators
statestats_queryd('MD')
## 1 indicator in 1 state, see a few key percentiles and mean
statestats_query('MD','proximity.tsdf')
## mean of 1 indicator for each state
statestats_query(varnames = 'proximity.tsdf')

```

```
## using full blockgroup dataset, not lookup tables of percentiles,
blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_d, names_e)]
##   see all total counts (not just US means),
##   demographics including subgroups,
##   but not environmental indicators.
t(round(EJAMbatch.summarizer::ustotals2(bg = EJAM::blockgroupstats),2))
t(blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_e, names_d)])

## End(Not run)
```

statestats_querye	<i>convenient way to see mean, pctiles of ENVIRONMENTAL indicators from lookup table</i>
-------------------	--

Description

convenient way to see mean, pctiles of ENVIRONMENTAL indicators from lookup table

Usage

```
statestats_querye(
  ST = sort(unique(EJAM::statestats$REGION)),
  varnames = EJAM::names_e,
  PCTILES = NULL,
  dig = 2
)
```

Examples

```
## Not run:
## in USA overall, see mean and key percentiles for all demog and envt indicators
usastats_query() # or statestats_query('us') # can say us or US or USA or usa etc.
usastats_query(PCTILES = 'mean')
usastats_means() # same but nicer looking format in console
usastats_means(dig=4)
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_d)]
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_e)]
## in 1 state, see mean and key percentiles for all demog and envt indicators
statestats_query('MD')
## in 1 state, see mean and key percentiles for just demog indicators
statestats_queryd('MD')
## 1 indicator in 1 state, see a few key percentiles and mean
statestats_query('MD','proximity.tsdf')
## mean of 1 indicator for each state
statestats_query(varnames = 'proximity.tsdf')
## using full blockgroup dataset, not lookup tables of percentiles,
blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_d, names_e)]
##   see all total counts (not just US means),
##   demographics including subgroups,
##   but not environmental indicators.
t(round(EJAMbatch.summarizer::ustotals2(bg = EJAM::blockgroupstats),2))
t(blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_e, names_d)])
```

```
## End(Not run)
```

states_infer	<i>states_infer</i> Get cleaned table of US State etc. by siteid, from lat/lon, or from FIPS
--------------	--

Description

states_infer Get cleaned table of US State etc. by siteid, from lat/lon, or from FIPS

Usage

```
states_infer(x)
```

Arguments

x data.frame or data.table with either ST column or lat and lon columns, or FIPS, and optionally a column with siteid or column called n

Value

data.frame with unique siteid, ST, etc.

See Also

[state_from_latlon\(\)](#) [state_from_fips\(\)](#)

states_shapefile	<i>US States boundaries 2020 shapefile from TIGER</i>
------------------	---

Description

This is used to figure out which state contains each point (facility/site).

Details

This is used by [state_from_latlon\(\)](#) to find which state is associated with each point that the user wants to analyze. That is needed to report indicators in the form of State-specific percentiles (e.g., a score that is at the 80th percentile within Texas). It is created by the package via `EJAM/data-raw/make_states_shapefile.R` Created roughly as follows:

```
setwd("~/../R/mysource/EJAM/data-raw") download.file("https://www.census.gov/cgi-bin/geo/shapefiles/index.php?y=2020")
destfile = "tl_2020_us_state.zip" ) dir.create("./shp") unzip("tl_2020_us_state.zip", exdir = "./shp")
states_shapefile <- sf::st_read("./shp", quiet = FALSE) class(states_shapefile)
```

See Also

[state_from_latlon\(\)](#) [get_blockpoints_in_shape\(\)](#)

state_from_blockid	<i>state_from_blockid</i> given vector of blockids, get state abbreviation of each
--------------------	--

Description

state_from_blockid given vector of blockids, get state abbreviation of each

Usage

```
state_from_blockid(blockid)
```

Arguments

blockid	vector of blockid values as from EJAM::blockpoints
---------	--

Value

vector of ST info like AK, CA, DE, etc.

Examples

```
state_from_blockid(c(8174952, blockpoints$blockid[5:6]))
```

state_from_blocktable	<i>state_from_blocktable</i> given data.table with blockid column, get state abbreviation of each
-----------------------	---

Description

state_from_blocktable given data.table with blockid column, get state abbreviation of each

Usage

```
state_from_blocktable(dt_with_blockid)
```

Arguments

dt_with_blockid	
-----------------	--

Value

vector of ST info like AK, CA, DE, etc.

Examples

```
state_from_blocktable(blockpoints[45:49,])
```

state_from_fips	<i>state_from_fips</i>
-----------------	------------------------

Description

state_from_fips

Usage

```
state_from_fips(fips)
```

Arguments

fips	Census FIPS codes vector, numeric or char, 2-digit, 5-digit, etc. OK
------	--

Value

vector of 2-character state abbreviations like CA,MD,TX,OH

state_from_latlon	<i>find what state is where each point is located Takes 3 seconds to find state for 1k points, so a faster alternative would be useful</i>
-------------------	--

Description

find what state is where each point is located Takes 3 seconds to find state for 1k points, so a faster alternative would be useful

Usage

```
state_from_latlon(lat, lon, states_shapefile = EJAM::states_shapefile)
```

Arguments

lat	latitudes vector
lon	longitudes vector
shapefile	shapefile of US States, in package already

Value

Returns data.frame: ST, statename, FIPS.ST, REGION, n as many rows as elements in lat or lon

See Also

[states_shapefile](#) [get_blockpoints_in_shape\(\)](#) [states_infer\(\)](#)

ST_by_site_from_sites2blocks

Get State that each site is in, from a table of siteid, blockid, distance

Description

Find the 2-character State abbreviation for each site. This is for when you need to know the state each site is in, to be able to report state percentiles, but you do not have the original list of siteid lat/lon or State info. This can infer the State each site is located in, based on the state of the nearest block (and its parent blockgroup).

Usage

```
ST_by_site_from_sites2blocks(sites2blocks)
```

Arguments

sites2blocks data.table or data.frame, like [sites2blocks_example](#), from [getblocksnearby\(\)](#) that has columns siteid and blockid and distance

Value

data.table with columns siteid, ST

Examples

```
ST_by_site_from_sites2blocks(sites2blocks_example)
## Not run:
fname = './inst/testdata/testpoints_207_sites_with_signif_violations_NAICS_326_ECHO.csv'
x = ST_by_site_from_sites2blocks(
  getblocksnearby( latlon_from_anything(fname), quadtree = localtree))
y = read_csv_or_xl(fname)
x$ST == y$FacState

## End(Not run)
```

summarize_blockcount *Get summary stats on counts of blocks (unique vs doublecounted) near sites*

Description

Get summary stats on counts of blocks (unique vs doublecounted) near sites

Usage

```
summarize_blockcount(x)
```

Arguments

x The output of [getblocksnearby\(\)](#) like sites2blocks_example

Value

A list of stats

Examples

```
summarize_blockcount(sites2blocks_example)
```

```
summarize_blocks_per_site
```

Get summary stats on counts of blocks near various sites

Description

Tells you # of blocks near avg site, how many sites have only 1 block nearby, or have <30 nearby, etc.

Usage

```
summarize_blocks_per_site(x, varname = "siteid")
```

Arguments

x	The output of <code>getblocksnearby()</code>
varname	colname of variable in data.table x that is the one to summarize by

Value

invisibly, a list of stats

```
summarize_sites_per_block
```

Get summary stats on how many sites are near various blocks (residents)

Description

Get summary stats on how many sites are near various blocks (residents)

Usage

```
summarize_sites_per_block(x, varname = "blockid")
```

Arguments

x	The output of <code>getblocksnearby()</code> like <code>sites2blocks_example</code>
varname	colname of variable in data.table x that is the one to summarize by

Value

invisibly, a list of stats

testdata_sites2blocks	<i>data.table of output of <code>getblocknearby()</code>, each row is a unique site-block-distance</i>
-----------------------	--

Description

data.table of output of `getblocknearby()`, each row is a unique site-block-distance

Details

```
library(EJAM) testdata_sites2blocks <- sites2blocks_example usethis::use_data(testdata_sites2blocks)
```

testpoints_1000_dt	<i>Random test points data.table with columns lat lon site</i>
--------------------	--

Description

Random test points data.table with columns lat lon site

testpoints_100_dt	<i>Random test points data.table with columns lat lon site</i>
-------------------	--

Description

Random test points data.table with columns lat lon site

testpoints_n	<i>Random points on the map Get data.table of Random Points (lat lon) for Testing/ Benchmarking/ Demos, weighted in various ways. The weighting can be specified so that each point reflects the average EPA-regulated facility, blockgroup, block, place on the map, or US resident.</i>
--------------	---

Description

Random points on the map Get data.table of Random Points (lat lon) for Testing/ Benchmarking/ Demos, weighted in various ways. The weighting can be specified so that each point reflects the average EPA-regulated facility, blockgroup, block, place on the map, or US resident.

Usage

```
testpoints_n(
  n = 10,
  weighting = c("frs", "pop", "area", "bg", "block"),
  dt = TRUE,
  ST_of_blockgroup = NULL
)
```

Arguments

n	Number of points needed (sample size)
weighting	word indicating how to weight the random points (some synonyms are allowed, in addition to those shown here): <ul style="list-style-type: none"> • pop or people = Average Person: random person among all US residents (block point of residence per 2020 Census) • frs or facility = Average Facility: random EPA-regulated facility from actives in Facility Registry Services (FRS) • bg = Average Blockgroup: random US Census block group (internal point like a centroid) • block = Average Block: random US Census block (internal point like a centroid) • area or place = Average Place: random point on a map (internal point of avg blockgroup weighted by its square meters size)
dt	logical, whether to return a data.table (DEFAULT) instead of normal data.frame
ST_of_blockgroup	optional, can be a character vector of 2 letter State abbreviations to pick from only some States, but this only works if the weighting is "bg" or "area" (or synonyms) right now.

Value

data.frame or data.table with columns lat, lon in decimal degrees, and any other columns that are in the table used (based on weighting)

Examples

```
## Not run:
n=2
for (d in c(TRUE,FALSE)) {
  for (w in c('frs', 'pop', 'area', 'bg', 'block')) {
    cat("n=",n," weighting=",w, " dt=",d,"\n\n")
    print(testpoints_n(n,w,d))
    cat('\n')
  }
}

## End(Not run)
```

test_regid

test data, vector of EPA FRS Registry ID numbers

Description

test data, vector of EPA FRS Registry ID numbers

Details

For testing, e.g.,

```
frs_from_siteid(test_regid)
```

```
mapfast( frs_from_regid(test_regid) )
```

url_bookmark_save	<i>url_bookmark_save save bookmarked EJScreen session (map location and indicator)</i>
-------------------	--

Description

url_bookmark_save save bookmarked EJScreen session (map location and indicator)

Usage

```
url_bookmark_save(..., file = "ejscreenbookmark.json")
```

Arguments

...	passed to url_bookmark_text()
file	path and name of .json file you want to save locally

Details

WORK IN PROGRESS - NOT USED AS OF EARLY 2023. You can use this function to create and save a json file that is a bookmark for a specific place/ map view/ data layer in EJScreen. You can later pull up that exact map in EJScreen by launching EJScreen, clicking Tools, Save Session, Load from File.

***Units are not lat lon: "spatialReference":{"latestWkid":3857,"wkid":102100

Note: (1) The number of sessions that can be saved depends on the browser cache size. (2) Session files, if saved, are available from the default Downloads folder on your computer. (3) Users should exercise caution when saving sessions that may contain sensitive or confidential data.

Value

URL for 1 bookmarked EJScreen map location and variable displayed on map

url_bookmark_text	<i>url_bookmark_text</i> URL for 1 bookmarked EJScreen session (map location and indicator)
-------------------	---

Description

url_bookmark_text URL for 1 bookmarked EJScreen session (map location and indicator)

Usage

```
url_bookmark_text(
  x = c(-13232599.1784247, -13085305.0249191),
  y = c(3970069.24597194, 4067373.582979),
  name = "BookmarkedEJScreenMap",
  title = "Socioeconomic Indicators",
  renderField = "B_UNEMPPCT",
  pctlevel = "nation",
  xmin = 1.1 * min(x),
  xmax = 0.9 * min(x),
  ymin = 0.9 * min(y),
  ymax = 1.1 * min(y),
  urlrest =
    "https://geopub.epa.gov/arcgis/rest/services/ejscreen/ejscreen_v2022_with_AS_CNMI_GU_VI/MapSe
  )
```

Arguments

x	vector of approx topleft, bottomright longitudes in some units EJScreen uses? Units are not lat lon: "spatialReference": "latestWkid": 3857, "wkid": 102100
y	vector of approx topleft, bottomright latitudes in some units EJScreen uses? Units are not lat lon: "spatialReference": "latestWkid": 3857, "wkid": 102100
name	Your name for the map bookmark
title	Your name for the map like Socioeconomic Indicators or Pollution and Sources
renderField	name of variable shown on map, like B_UNEMPPCT for map color bins of percent unemployed or B_PTRAFF for traffic indicator
pctlevel	nation or state
xmin	calculated bounding box for map view
xmax	calculated bounding box for map view
ymin	calculated bounding box for map view
ymax	calculated bounding box for map view
urlrest	Just use the default but it changes each year

Details

WORK IN PROGRESS - NOT USED AS OF EARLY 2023. You can use this function to create and save a json file that is a bookmark for a specific place/ map view/ data layer in EJScreen. You can later pull up that exact map in EJScreen by launching EJScreen, clicking Tools, Save Session, Load from File.

Note: (1) The number of sessions that can be saved depends on the browser cache size. (2) Session files, if saved, are available from the default Downloads folder on your computer. (3) Users should exercise caution when saving sessions that may contain sensitive or confidential data.

Value

URL for 1 bookmarked EJScreen map location and variable displayed on map

See Also

[url_bookmark_save\(\)](#)

Examples

```
## Not run:
url_bookmark_text()
url_bookmark_save(
  x=c(-10173158.179197036, -10128824.702791695),
  y=c(3548990.034736070, 3579297.316451102),
  file="./mysavedejscreensession1.json")

## End(Not run)
```

url_getacs_epaquery	<i>experimental/ work in progress: get ACS data via EPA API (for <200 places)</i>
---------------------	--

Description

uses ACS2019 rest services ejscreen ejquery MapServer 7

Documentation of format and examples of input parameters:

https://geopub.epa.gov/arcgis/sdk/rest/index.html#/Query_Map_Service_Layer/02ss0000000r000000/

Usage

```
url_getacs_epaquery(
  objectIds = 1:3,
  servicenumber = 7,
  outFields = NULL,
  returnGeometry = FALSE,
  justurl = FALSE,
  ...
)
```

Arguments

objectIds	see API
servicenumber	see API
outFields	see API. eg "STCNTRBG", "TOTALPOP", "PCT_HISP",
returnGeometry	see API
justurl	if TRUE, returns url instead of default making API request
...	passed to url_getacs_epaquery_chunked()

Value

table

Examples

```
url_getacs_epaquery(justurl=TRUE)
```

```
url_getacs_epaquery_chunked
```

experimental/ work in progress: in chunks, get ACS data via EPA API

Description

experimental/ work in progress: in chunks, get ACS data via EPA API

Usage

```
url_getacs_epaquery_chunked(
  objectIds = 1:3,
  servicenumber = 7,
  outFields = NULL,
  returnGeometry = FALSE,
  justurl = FALSE,
  chunksize = 200,
  ...
)
```

Arguments

objectIds	see API
servicenumber	see API
outFields	see API
returnGeometry	see API
justurl	see API
chunksize	eg 200 for chunks of 200 each request
...	passed to url_getacs_epaquery()

Value

table

Examples

```
#
#\dontrun {
# x <- list() # chunked chunks. best not to ask for all these:
# x[[1]] <- url_getacs_epaquery_chunked( 1:1000, chunksize = 100)
# x[[2]] <- url_getacs_epaquery_chunked(1001:5000, chunksize = 100)
# xall <- do.call(rbind, x)
#}
```

`url_get_eparest_chunked_by_id`*experimental/ work in progress: in chunks, get ACS data or Block weights nearby via EPA API*

Description

experimental/ work in progress: in chunks, get ACS data or Block weights nearby via EPA API

Usage

```
url_get_eparest_chunked_by_id(objectIds, chunksize = 200, ...)
```

Arguments

<code>objectIds</code>	see API
<code>chunksize</code>	see API
<code>...</code>	passed to <code>url_getacs_epaquery()</code>

Value

a table

<code>url_get_via_url</code>	<i>helper function work in progress: GET json via url of ejscreen ejquery map services</i>
------------------------------	--

Description

helper function work in progress: GET json via url of ejscreen ejquery map services

Usage

```
url_get_via_url(url)
```

Arguments

<code>url</code>	the url for an EJScreen ejquery request
------------------	---

Value

json

url_naics.com	<i>Get URL for page with info about industry sectors by text query term See naics.com for more information on NAICS codes</i>
---------------	---

Description

Get URL for page with info about industry sectors by text query term See [naics.com](#) for more information on NAICS codes

Usage

```
url_naics.com(query, as_html = FALSE, linktext)
```

Arguments

query	string query term like "gasoline" or "copper smelting"
as_html	Whether to return as just the urls or as html hyperlinks to use in a DT::datatable() for example
linktext	used as text for hyperlinks, if supplied and as_html=TRUE

Value

URL as string

usastats	<i>data.frame of 100 percentiles and means</i>
----------	--

Description

data.frame of 100 percentiles and means (about 100 rows) in the USA overall, across all locations (e.g., block groups in [blockgroupstats](#)) for a set of indicators such as percent low income. Each column is one indicator (or specifies the percentile).

This should be similar to the lookup tables in the gdb on the FTP site of EJScreen, except it also has data for the demographic race/ethnicity subgroups. For details on how the table was made, see [/EJAM/data-raw/usastats_subgroups.R](#)

usastats_means	<i>convenient way to see USA MEANS of ENVIRONMENTAL and DEMOGRAPHIC indicators from lookup table</i>
----------------	--

Description

convenient way to see USA MEANS of ENVIRONMENTAL and DEMOGRAPHIC indicators from lookup table

Usage

```
usastats_means(...)
```

Arguments

... Arguments passed on to [usastats_query](#)

varnames names of columns in lookup table, like "proximity.rmp"

PCTILES vector of percentiles 0-100 and/or "mean"

usastats_query	<i>convenient way to see USA mean, pctiles of Env and Demog indicators from lookup table</i>
----------------	--

Description

convenient way to see USA mean, pctiles of Env and Demog indicators from lookup table

Usage

```
usastats_query(
  varnames = c(EJAM::names_e, EJAM::names_d),
  PCTILES = NULL,
  dig = 2
)
```

Arguments

varnames names of columns in lookup table, like "proximity.rmp"

PCTILES vector of percentiles 0-100 and/or "mean"

@dig how many digits to round to

Examples

```
## Not run:
## in USA overall, see mean and key percentiles for all demog and envt indicators
usastats_query() # or statestats_query('us') # can say us or US or USA or usa etc.
usastats_query(PCTILES = 'mean')
usastats_means() # same but nicer looking format in console
usastats_means(dig=4)
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_d)]
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_e)]
## in 1 state, see mean and key percentiles for all demog and envt indicators
statestats_query('MD')
## in 1 state, see mean and key percentiles for just demog indicators
statestats_queryd('MD')
## 1 indicator in 1 state, see a few key percentiles and mean
statestats_query('MD','proximity.tsdf')
## mean of 1 indicator for each state
statestats_query(varnames = 'proximity.tsdf')
## using full blockgroup dataset, not lookup tables of percentiles,
blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_d, names_e)]
## see all total counts (not just US means),
## demographics including subgroups,
## but not environmental indicators.
t(round(EJAMbatch.summarizer::ustotals2(bg = EJAM::blockgroupstats),2))
t(blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_e, names_d)])

## End(Not run)
```

usastats_queryd	<i>convenient way to see USA mean, pctiles of DEMOGRAPHIC indicators from lookup table</i>
-----------------	--

Description

convenient way to see USA mean, pctiles of DEMOGRAPHIC indicators from lookup table

Usage

```
usastats_queryd(varnames = EJAM::names_d, PCTILES = NULL, dig = 2)
```

Arguments

varnames	names of columns in lookup table, like "proximity.rmp"
PCTILES	vector of percentiles 0-100 and/or "mean"
@dig	how many digits to round to

Examples

```
## Not run:
## in USA overall, see mean and key percentiles for all demog and envt indicators
usastats_query() # or statestats_query('us') # can say us or US or USA or usa etc.
usastats_query(PCTILES = 'mean')
```

```

usastats_means() # same but nicer looking format in console
usastats_means(dig=4)
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_d)]
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_e)]
## in 1 state, see mean and key percentiles for all demog and envt indicators
statestats_query('MD')
## in 1 state, see mean and key percentiles for just demog indicators
statestats_queryd('MD')
## 1 indicator in 1 state, see a few key percentiles and mean
statestats_query('MD','proximity.tsdf')
## mean of 1 indicator for each state
statestats_query(varnames = 'proximity.tsdf')
## using full blockgroup dataset, not lookup tables of percentiles,
blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_d, names_e)]
## see all total counts (not just US means),
## demographics including subgroups,
## but not environmental indicators.
t(round(EJAMbatch.summarizer::ustotals2(bg = EJAM::blockgroupstats),2))
t(blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_e, names_d)])

## End(Not run)

```

usastats_querye	<i>convenient way to see USA mean, pctiles of ENVIRONMENTAL indicators from lookup table</i>
-----------------	--

Description

convenient way to see USA mean, pctiles of ENVIRONMENTAL indicators from lookup table

Usage

```
usastats_querye(varnames = EJAM::names_e, PCTILES = NULL, dig = 2)
```

Arguments

varnames	names of columns in lookup table, like "proximity.rmp"
PCTILES	vector of percentiles 0-100 and/or "mean"
@dig	how many digits to round to

Examples

```

## Not run:
## in USA overall, see mean and key percentiles for all demog and envt indicators
usastats_query() # or statestats_query('us') # can say us or US or USA or usa etc.
usastats_query(PCTILES = 'mean')
usastats_means() # same but nicer looking format in console
usastats_means(dig=4)
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_d)]
usastats[!(usastats$PCTILE < 50), c("PCTILE", names_e)]
## in 1 state, see mean and key percentiles for all demog and envt indicators

```

```

statestats_query('MD')
## in 1 state, see mean and key percentiles for just demog indicators
statestats_queryd('MD')
## 1 indicator in 1 state, see a few key percentiles and mean
statestats_query('MD','proximity.tsdf')
## mean of 1 indicator for each state
statestats_query(varnames = 'proximity.tsdf')
## using full blockgroup dataset, not lookup tables of percentiles,
blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_d, names_e)]
## see all total counts (not just US means),
## demographics including subgroups,
## but not environmental indicators.
t(round(EJAMbatch.summarizer::ustotals2(bg = EJAM::blockgroupstats),2))
t(blockgroupstats[, lapply(.SD, function(x) mean(x, na.rm=T)), .SDcols= c(names_e, names_d)])

## End(Not run)

```

workbook_output_styled

workbook_output_styled Format EJAM tabular outputs for saving as Excel spreadsheet

Description

workbook_output_styled Format EJAM tabular outputs for saving as Excel spreadsheet

Usage

```

workbook_output_styled(
  overall,
  eachsite,
  graycolnums = NULL,
  narrowcolnums = NULL,
  graycolor = "gray",
  narrow6 = 6,
  ...
)

```

Arguments

overall	table to save in one tab, from EJAM analysis of indicators overall (one row)
eachsite	table to save in another tab, from EJAM analysis site by site (one row per site)
graycolnums	which columns to deemphasize
narrowcolnums	which columns to make narrow
graycolor	color used to deemphasize some columns
narrow6	how narrow
...	other params passed along to openxlsx::writeData()

Details

NEED TO MERGE THIS WITH EJAMEjscreenapi's prep_for_excel()

Value

a workbook, ready to be saved in spreadsheet format, with tabs like "Overall" and "Each Site"

Examples

```
## Not run:
wb <- workbook_output_styled(datasetResults())$results_overall, datasetResults())$results_bysite)
openxlsx::saveWorkbook(wb, "results.xlsx", overwrite = TRUE)

## End(Not run)
```

write_pctiles_lookup	<i>create lookup table of percentiles 0 to 100 and mean for each indicator by State or USA total</i>
----------------------	--

Description

create lookup table of percentiles 0 to 100 and mean for each indicator by State or USA total

Usage

```
write_pctiles_lookup(
  x,
  zone.vector = NULL,
  zoneOverallName = "USA",
  wts = NULL
)
```

Arguments

x	data.frame with numeric data. Each column will be examined to calculate mean, sd, and percentiles, for each zone
zone.vector	optional names of states or regions, for example. same length as wts, or rows in mydf
zoneOverallName	optional. Default is USA.
wts	not used in EJScreen percentiles anymore

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