Package 'geohabnet'

October 11, 2023

Title Analysis of Cropland Connectivity

Version 1.0.0

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Description Geographical spatial analysis of cropland connectivity.
Allows users to visualize risk index plots for a given set of crops.
The functions are developed as an extension to analysis from Xing et al (2021) <doi:10.1093/biosci/biaa067>.
The primary function is sean() and is indicative of how sensitive analysis is to parameters using kernel models.
Package currently supports crops sourced from Monfreda, C., N. Ramankutty, and J. A. Foley (2008) <doi:10.1029/2007gb002947> ``Farming the planet: 2. Geographic distribution of crop areas, yields, physiological types, and net primary production in the year 2000, Global Biogeochem. Cycles, 22, GB1022" and
International Food Policy Research Institute (2019) <doi:10.7910/DVN/PRFF8V> ``Global Spatially-Disaggregated Crop Production Statistics Data for 2010 Version 2.0, Harvard Dataverse, V4".
This analysis produces 3 maps - mean, variance, and difference for the crop risk index. It applies distance functions and graph operations on network to calculate risk index.
There are multiple ways in which functions can be used -
generate final outcome and then the intermediate outcomes for more sophisticated use cases.
Refer to vignettes.
sean() will set some global variables which can be accessed using \code{the$} prefix. These values are propagated to other functions for performing operations such as distance matrix calculation.
\code{'parameters.yaml'} stores the parameters and values and can be accessed using get_parameters(). Refer it's usage.
The objective of this package to support risk analysis using cropland connectivity on 10 parameters -
host crops, density threshold, aggregation and distance method, resolution, geographic extent, link threshold, kernel models, network metrics and maps.
These parameters serves as an input and are used different phases of analysis workflow.

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Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.2.3

Imports config (>= 0.3.1),
geodata (>= 0.5.8),
geosphere (>= 1.5.18),

igraph (>= 1.4.2),
terra (>= 1.7.29),
easycsv (>= 1.0.8),
yaml (>= 2.3.7),
stats (>= 4.2.3),
stringr (>= 1.5.0),
memoise (>= 2.0.1),
graphics (>= 4.2.3),
rlang (>= 1.1.1),
viridisLite (>= 0.4.2),
beepr (>= 1.3),
rnaturalearth (>= 0.3.3)

Suggests devtools,
knitr,
lintr (>= 3.0.2),
mockthat (>= 0.2.8),
pkgdown (>= 2.0.7.9000),
rmarkdown,
testthat (>= 3.1.7)

URL https://garrettlab.github.io/CroplandConnectivity/,
https://github.com/GarrettLab/CroplandConnectivity/tree/main/geohabnet/,
https://www.garrettlab.com/

BugReports https://github.com/GarrettLab/CroplandConnectivity/issues

VignetteBuilder knitr

R topics documented:

`ccri_diff` *Calculate difference map*

Description

This function produces a map of difference b/w mean and sum indexes in rank of cropland harvested area fraction.

Usage

```
ccri_diff(rast, x, y, global, geoscale, res = reso(), outdir = tempdir())
```

Arguments

`rast`	A raster object for mean index raster difference
`x`	A raster object for cropland harvest
`y`	A raster object for cropland harvest
`global`	logical. `TRUE` if global analysis is required, `FALSE` otherwise. When `TRUE`, `geoscale` is ignored. Default is `TRUE`.
`geoscale`	vector. geographical scale
`res`	numeric. map resolution.
`outdir`	Character. Output directory for saving raster in TIFF format. Default is `tempdir()`.

`ccri_mean` *Calculate mean of raster objects*

Description

Wrapper for `terra::mean()`. Calculates mean of list of rasters.

Usage

```
ccri_mean(
  indexes,
  global = TRUE,
  geoscale = NULL,
  plt = TRUE,
  outdir = tempdir()
)
```

Arguments

indexes	list of rasters. See details.
global	logical. TRUE if global analysis is required, FALSE otherwise. When TRUE, geoscale is ignored. Default is TRUE.
geoscale	vector. geographical scale
plt	TRUE if need to plot mean map, FALSE otherwise and geoscale.
outdir	Character. Output directory for saving raster in TIFF format. Default is tempdir().

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ccri_variance *Calculate variance of CCRI*

Description

This function produces a map of variance of CCRI based on input parameters

Usage

```
ccri_variance(
  indexes,
  rast,
  global,
  geoscale,
  res = reso(),
  outdir = tempdir()
)
```

Arguments

indexes	list of rasters. See details.
rast	A raster object. It will be used in calculating variance.
global	logical. TRUE if global analysis is required, FALSE otherwise. When TRUE, geoscale is ignored. Default is TRUE.
geoscale	vector. geographical scale
res	numeric. map resolution.
outdir	Character. Output directory for saving raster in TIFF format. Default is tempdir().

`connectivity` *Calculate and plot maps*

Description

Calculate mean, variance and difference. The result is produced in form of maps plotted with predefined settings. Currently, the settings for plot cannot be customized. Default value is `TRUE` for all logical arguments

Usage

```
connectivity(
  indexes,
  global = TRUE,
  geoscale,
  res = reso(),
  pmean = TRUE,
  pvar = TRUE,
  pdiff = TRUE,
  outdir = tempdir()
)
```

Arguments

`indexes`	list of rasters. See details.
`global`	logical. `TRUE` if global analysis is required, `FALSE` otherwise. When `TRUE`, `geoscale` is ignored. Default is `TRUE`.
`geoscale`	vector. geographical scale
`res`	numeric. map resolution.
`pmean`	`TRUE` if map of mean should be plotted, `FALSE` otherwise.
`pvar`	`TRUE` if variance map should be plotted, `FALSE` otherwise.
`pdiff`	`TRUE` if difference map should be plotted, `FALSE` otherwise.
`outdir`	Character. Output directory for saving raster in TIFF format. Default is `tempdir()`.

Details

`indexes` are actually risk resulting from operations on crop's raster and parameters provided in either `parameters.yaml` or `sean()`.

It will save all the opted plots using - `pmean`, `pvar` and `pdiff`. File will be saved in provided value of `outdir` or `tempdir()`.If `interactive()` is `TRUE`, then plots can be seen in active plot window. E.g. Rstudio

References

Yanru Xing, John F Hernandez Nopsa, Kelsey F Andersen, Jorge L Andrade-Piedra, Fenton D Beed, Guy Blomme, Mónica Carvajal-Yepes, Danny L Coyne, Wilmer J Cuellar, Gregory A Forbes, Jan F Kreuze, Jürgen Kroschel, P Lava Kumar, James P Legg, Monica Parker, Elmar Schulte-Geldermann, Kalpana Sharma, Karen A Garrett, *Global Cropland Connectivity: A Risk Factor*

for Invasion and Saturation by Emerging Pathogens and Pests, BioScience, Volume 70, Issue 9, September 2020, Pages 744–758, doi:10.1093/biosci/biaa067

Hijmans R (2023). *terra: Spatial Data Analysis*. R package version 1.7-46, \@urlhttps://CRAN.R-project.org/package=terra

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`cropharvest_rast` *Get raster object for crop*

Description

Get cropland information in a form of raster object from data source for crop

Usage

```
cropharvest_rast(crop_name, data_source)
```

Arguments

`crop_name`	Name of the crop
`data_source`	Data source for cropland information

Value

Raster object

Examples

```
## Not run:
cropharvest_rast("avocado", "monfreda")

## End(Not run)
```

`crops_rast` *Get sum of rasters for individual crops*

Description

Takes crop names and returns raster object which is sum of raster of individual crops. Currently, only supports crops listed in `geodata::monfredaCrops()`, `geodata::spamCrops()` If crop is present in multiple sources, then their mean is calculated.

Usage

```
crops_rast(crop_names)
```

Arguments

`crop_names`	A named list of source along with crop names

Value

SpatRaster. Raster object which is sum of all the individual crop rasters

Examples

```
## Not run:
crops_rast(list(monfreda = c("wheat", "barley"), mapspam = c("wheat", "potato")))

## End(Not run)
```

---

dist_methods *Distance methods supported*

Description

Contains supported strategies to calculate distance between two points. Use of one the methods in `sean()` or `sensitivity_analysis()`.

Usage

```
dist_methods()
```

Value

vector

Examples

```
dist_methods()
```

---

geoscale_param *Get geographical scales from the parameters*

Description

This function returns a list of geographical scales set in global and custom extent in `parameters.yaml`. If `global` is `TRUE`, the `CustomExt` is ignored.

Usage

```
geoscale_param()
```

Value

A list of geographical scales

get_parameters *Get Parameters*

Description

Retrieves the parameters and copies the parameter file to the specified output path.

Usage

```
get_parameters(out_path = tempdir(), iwindow = FALSE)
```

Arguments

out_path	character. The output path where the parameter file will be copied. Default is temporary directory tempdir()
iwindow	logical. If TRUE, prompts the user to select the output directory using a file chooser window. Default is FALSE

Details

Using configuration file is an alternative to sean()

Value

character. The path to the copied parameter file.

See Also

set_parameters()

Examples

```
get_parameters()
get_parameters(out = tempdir())
```

get_param_metrics *Get metrics from parameters*

Description

Get metrics and parameters stored in parameters.yaml.

Usage

```
get_param_metrics(params = load_parameters())
```

Arguments

params	R object of load_parameters(). Default is load_parameters().

Value

List of metrics - parameters and values. See usage.

Examples

```
# Get metrics from parameters
get_param_metrics()
get_param_metrics(load_parameters())
```

`get_rasters` *Get rasters object from parameters*

Description

Takes named list of hosts as an input. See host object in `get_parameters()` or `load_parameters()`. This is also a wrapper of `crops_rast()`. Function creates 2 raster object - one is a sum of all the crops specified under sources and other using the provided raster file. See `tiff_torast()`

Usage

```
get_rasters(hosts)
```

Arguments

`hosts`	List of hosts and values. It is synonym to Hosts object in parameters

Value

List of SpatRaster.

See Also

`load_parameters()`, `get_parameters()`, `tiff_torast()`, `cropharvest_rast()`

Examples

```
# Get default rasters
## Not run:
get_rasters(list(mapspam = c("wheat"), monfreda = c("avocado"), file = "some_raster.tif"))

## End(Not run)
```

get_supported_sources *Get supported sources of crops*

Description

When provided, `cropharvest_rast()` will look for cropland data in this specific source.

Usage

```
get_supported_sources()
```

Value

return vector of supported sources. Also used as a lookup to find get raster object.

Examples

```
# Get currently supported sources
get_supported_sources()
```

global_scales *Global geographical extent*

Description

See geographical extents used in global analysis. Returns eastern and western hemisphere extents. Each extent is in the form of c(Xmin, Xmax, Ymin, Ymax).

Usage

```
global_scales()
```

See Also

`set_global_scales()`

load_parameters *Load Parameters from YAML File*

Description

This function loads parameters from a YAML file and stores them in an object.

Usage

```
load_parameters(filepath = .get_helper_filepath(.kparameters_file_type))
```

Arguments

filepath	Path to the YAML file containing the parameters. By default, it takes the value of ".kparameters_file_type" which is set to "parameters.yaml".

Value

object with parameters and values

Examples

```
# Load parameters from default file
load_parameters()
```

model_powerlaw *Calculate risk index using inbuilt models.*

Description

- model_powerlaw() calculates risk index using power law.
- model_neg_exp() calculates risk index using negative exponential.

Usage

```
model_powerlaw(
  beta,
  link_threshold,
  distance_matrix = the$distance_matrix,
  thresholded_crop_values,
  adj_mat = NULL,
  crop_raster,
  crop_cells_above_threshold,
 metrics = the$parameters_config$`CCRI parameters`$NetworkMetrics$InversePowerLaw
)

model_neg_exp(
  gamma_val,
  link_threshold,
```

```
  distance_matrix = the$distance_matrix,
  thresholded_crop_values,
  adj_mat = NULL,
  crop_raster,
  crop_cells_above_threshold,
 metrics = the$parameters_config$`CCRI parameters`$NetworkMetrics$InversePowerLaw
)
```

Arguments

beta	A list of beta values. DispersalParameterBeta in parameters.yaml.
link_threshold	A threshold value for link.
distance_matrix	distance matrix, generated during sean().
thresholded_crop_values	crop values above threshold.
adj_mat	Adjacency matrix(optional) representing un-directed graph network. If this is provided, then gamma_val, distance_matrix, link_threshold and thresholded_crop_values are ignored. These ignored parameters are used to generate adjacency matrix internally. This is the only way to use custom adjacency matrix.
crop_raster	A raster object for cropland harvest.
crop_cells_above_threshold	crop cells above threshold. Only contains cells and not the the values.
metrics	A list 2 vectors - metrics and weights.
gamma_val	A list of beta values. DispersalParameterGamma in parameters.yaml.

Details

Network metrics should be passed as a list of vectors e.g. `list(metrics = c("betweeness"), weights = c(100))`. Default values are fetched from `parameters.yaml` and arguments uses the same structure.

Value

risk index

---

nn_sum *Calculation on network matrix.*

Description

These are basically an abstraction of functions under the igraph package. The functions included in this abstraction are:

- `nn_sum()`: Calculates the sum of nearest neighbors `igraph::graph.knn()`.
- `node_strength()`: Calculates the sum of edge weights of adjacent nodes `igraph::graph.strength()`.
- `betweeness()`: Calculates the vertex and edge betweenness based on the number of geodesics `igraph::betweenness()`.
- `ev()`: Calculates the eigenvector centrality of positions within the network `igraph::evcent()`.

- `closeness()`: measures how many steps is required to access every other vertex from a given vertex `igraph::closeness()`.
- `degree()`: number of adjacent edges `igraph::degree()`.
- `pagerank()`: page rank score for vertices `igraph::page_rank()`.

Usage

```
nn_sum(crop_dm, we)

node_strength(crop_dm, we)

betweeness(crop_dm, we)

ev(crop_dm, we)

degree(crop_dm, we)

closeness(crop_dm, we)

pagerank(crop_dm, we)
```

Arguments

`crop_dm`	Distance matrix. In the internal workflow, the distance matrix comes is a result of operations within `sean()` and risk functions.
`we`	Weight in percentage.

Value

Matrix with the mean value based on the assigned weight.

See Also

Other metrics: `supported_metrics()`

`reset_params` *Reset parameters.yaml*

Description

Resets the values in the `parameters.yaml` file to the default initial values.

Usage

```
reset_params()
```

Examples

```
## Not run:
reset_params()

## End(Not run)
```

reso *Get resolution value*

Description

Resolution stored in `parameter.yaml`. If not present it will result default value.

Usage

```
reso()
```

See Also

`set_reso()`

sa_onrasters *Run senstivity analysis*

Description

Same as `sensitivity_analysis()` but it takes raster object and other parameters as an input.

Usage

```
sa_onrasters(
  rast,
  global = TRUE,
  geoscale,
  link_thresholds,
  host_density_thresholds,
  agg_methods = c("sum", "mean"),
  dist_method = "geodesic",
  res = reso(),
  maps = TRUE,
  outdir = tempdir()
)
```

Arguments

`rast`	Raster object which will be used in analysis.
`global`	Logical. `TRUE` if global analysis, `FALSE` otherwise. Default is `TRUE`
`geoscale`	Vector. Geographical coordinates in the form of c(Xmin, Xmax, Ymin, Ymax)
`link_thresholds`	vector. link threshold values
`host_density_thresholds`	vector. host density threshold values
`agg_methods`	vector. Aggregation methods
`dist_method`	character. One of the values from `dist_methods()`
`res`	numeric. resolution at which operations will run. Default is `reso()`
`maps`	logical. `TRUE` if maps are to be plotted, `FALSE` otherwise
`outdir`	Character. Output directory for saving raster in TIFF format. Default is `tempdir()`.

Details

When `global = TRUE`, `geo_scale` is ignored. Instead uses scales from `global_scales()`.

Value

A list of calculated CCRI indices after operations. An index is generated for each combination of paramters. One combination is equivalent to `sean()` function.

References

Yanru Xing, John F Hernandez Nopsa, Kelsey F Andersen, Jorge L Andrade-Piedra, Fenton D Beed, Guy Blomme, Mónica Carvajal-Yepes, Danny L Coyne, Wilmer J Cuellar, Gregory A Forbes, Jan F Kreuze, Jürgen Kroschel, P Lava Kumar, James P Legg, Monica Parker, Elmar Schulte-Geldermann, Kalpana Sharma, Karen A Garrett, *Global Cropland Connectivity: A Risk Factor for Invasion and Saturation by Emerging Pathogens and Pests*, BioScience, Volume 70, Issue 9, September 2020, Pages 744–758, doi:10.1093/biosci/biaa067

Hijmans R (2023). *terra: Spatial Data Analysis*. R package version 1.7-46, \@urlhttps://CRAN.R-project.org/package=terra

See Also

Use `get_rasters()` to obtain raster object.

Examples

```
rr <- get_rasters(list(monfreda = c("avocado")))
res1 <- sa_onrasters(rr[[1]],
            global = FALSE,
            geoscale = c(-115, -75, 5, 32),
            c(0.0001, 0.00004),
            c(0.0001, 0.00005),
            c("sum", "mean"),
            res = 24)
res2 <- sa_onrasters(rr[[1]],
            global = TRUE,
            link_thresholds = c(0.000001),
            host_density_thresholds = c(0.00015),
            agg_methods = c("sum"),
            res = 24)
```

`sean` *Calculate sensitivity analysis on cropland harvested area fraction*

Description

This function calculates sensitivity analysis on cropland harvested area fraction based on provided parameters. Some parameters are only accessible from `paramters.yaml` and uses value from here

Usage

```
sean(
  rast,
  global = TRUE,
  geoscale,
  agg_methods = c("sum", "mean"),
  dist_method = "geodesic",
  link_threshold = 0,
  host_density_threshold = 0,
  res = reso(),
  maps = TRUE,
  outdir = tempdir()
)
```

Arguments

rast	Raster object which will be used in analysis.
global	Logical. TRUE if global analysis, FALSE otherwise. Default is TRUE
geoscale	Vector. Geographical coordinates in the form of c(Xmin, Xmax, Ymin, Ymax)
agg_methods	vector. Aggregation methods
dist_method	character. One of the values from dist_methods()
link_threshold	numeric. A threshold value for link
host_density_threshold	A host density threshold value
res	numeric. resolution at which operations will run. Default is reso()
maps	logical. TRUE if maps are to be plotted, FALSE otherwise
outdir	Character. Output directory for saving raster in TIFF format. Default is tempdir().

Details

When global = TRUE, geoscale is ignored and global_scales() is used

Value

A list of calculated CCRI values using negative exponential

References

Yanru Xing, John F Hernandez Nopsa, Kelsey F Andersen, Jorge L Andrade-Piedra, Fenton D Beed, Guy Blomme, Mónica Carvajal-Yepes, Danny L Coyne, Wilmer J Cuellar, Gregory A Forbes, Jan F Kreuze, Jürgen Kroschel, P Lava Kumar, James P Legg, Monica Parker, Elmar Schulte-Geldermann, Kalpana Sharma, Karen A Garrett, *Global Cropland Connectivity: A Risk Factor for Invasion and Saturation by Emerging Pathogens and Pests*, BioScience, Volume 70, Issue 9, September 2020, Pages 744–758, doi:10.1093/biosci/biaa067

Hijmans R (2023). *terra: Spatial Data Analysis*. R package version 1.7-46, \@urlhttps://CRAN.R-project.org/package=terra

See Also

Uses connectivity()

Examples

```
avocado <- cropharvest_rast("avocado", "monfreda")
risk_indexes <- sean(avocado)
```

`search_crop` *Search for crop*

Description

It returns the dataset sources in which crop data is available. It's a wrapper around `geodata::spamCrops()` and `geodata::monfredaCrops()`

Usage

```
search_crop(name)
```

Arguments

`name`	name of crop

See Also

`get_supported_sources()`

Examples

```
search_crop("coffee")
search_crop("wheat")

search_crop("jackfruit")
```

`sensitivity_analysis` *Calculate sensitivity analysis on parameters*

Description

This function runs sensitivity analysis on parameters based on parameters provided through `set_parameters()`. It can be used as an entry point for CCRI. By default, it runs analysis on global sclaes`global_scales()`. After analysis is complete, it will suppress maps for outcomes if `maps = FALSE` or `interactive()` is `FALSE`.

Usage

```
sensitivity_analysis(maps = TRUE, alert = TRUE)
```

Arguments

maps	logical. TRUE if maps are to be plotted, FALSE otherwise
alert	logical. TRUE if beep sound is to be played, FALSE otherwise

Value

logical. TRUE if analysis is completed, FALSE otherwise. Errors are not handled.

References

Yanru Xing, John F Hernandez Nopsa, Kelsey F Andersen, Jorge L Andrade-Piedra, Fenton D Beed, Guy Blomme, Mónica Carvajal-Yepes, Danny L Coyne, Wilmer J Cuellar, Gregory A Forbes, Jan F Kreuze, Jürgen Kroschel, P Lava Kumar, James P Legg, Monica Parker, Elmar Schulte-Geldermann, Kalpana Sharma, Karen A Garrett, *Global Cropland Connectivity: A Risk Factor for Invasion and Saturation by Emerging Pathogens and Pests*, BioScience, Volume 70, Issue 9, September 2020, Pages 744–758, doi:10.1093/biosci/biaa067

Hijmans R (2023). *terra: Spatial Data Analysis*. R package version 1.7-46, \@urlhttps://CRAN.R-project.org/package=terra

See Also

sa_onrasters() sean() global_scales() get_parameters() set_parameters() connectivity()

Examples

```
# Run analysis on specified parameters.yaml
sensitivity_analysis()
sensitivity_analysis(FALSE, FALSE)
sensitivity_analysis(TRUE, FALSE)
```

---

set_global_scales *Set global geographical extent*

Description

Set the geographical extents used in global analysis. Each extent should be in the form of c(Xmin, Xmax, Ymin, Ymax)

Usage

```
set_global_scales(value)
```

Arguments

value	list. Named list of eastern and western hemisphere extents. See usage.

See Also

global_scales() terra::ext()

Examples

```
## Not run:
set_global_scales(list(east = c(-24, 180, -58, 60), west = c(-140, -34, -58, 60)))

## End(Not run)
```

`set_parameters` *Set Parameters*

Description

This function allows you to set the parameters by replacing the existing parameters file with a new one. Use `get_parameters()` to modify the parameter values.

Usage

```
set_parameters(new_params, iwindow = FALSE)
```

Arguments

`new_params`	The path to the new parameters file.
`iwindow`	Logical indicating whether to prompt the user to select the new parameters file using a file selection window. Defaults to FALSE.

Value

None

Examples

```
param_fp <- get_parameters()
set_parameters(param_fp)
```

`set_reso` *Set resolution value*

Description

Set `resolution` to be used in analysis. It doesn't modify the `parameters.yaml` but instead a currently loaded instance of it. Must be greater than 0 and less than or equal to 48.

Usage

```
set_reso(value)
```

Arguments

`value`	numeric. Resolution value.

Examples

```
## Not run:
set_reso(24)

## End(Not run)
```

`sp_rast` *raster for mapspam crop.*

Description

get raster for crop in mapspam dataset

Usage

```
sp_rast(crp)
```

Arguments

`crp`	character. name of a crop. Case-insensitive.

Details

See `geodata::spamCrops()` for supported crops.

Value

spatRaster

References

International Food Policy Research Institute, 2020. Spatially-Disaggregated Crop Production Statistics Data in Africa South of the Sahara for 2017. <doi: 10.7910/DVN/FSSKBW>, Harvard Dataverse, V2

See Also

`geodata::spamCrops()` `search_crop()`

Examples

```
## Not run:
sp_rast("rice")

## End(Not run)
```

supported_metrics *Returns metrics currently supported in the analysis.*

Description

Returns metrics currently supported in the analysis.

Usage

```
supported_metrics()
```

Value

vector of supported metrics.

See Also

Other metrics: `nn_sum()`

Examples

```
supported_metrics()
```

tiff_torast *Get raster object from tif file*

Description

This is a wrapper of `terra::rast()` and generates a raster object if provided with a TIF file.

Usage

```
tiff_torast(path_to_tif)
```

Arguments

path_to_tif	TIF file

Value

SpatRaster.

Examples

```
## Not run:
# Generate raster for usage
fp <- .get_helper_filepath("avocado")
tiff_torast(fp)
tiff_torast("avocado_HarvestedAreaFraction.tif")


## End(Not run)
```