Package ‘disaggregation’

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Type Package

Title Disaggregation Modelling

Version 0.1.3

Description Fits disaggregation regression models using 'TMB' ('Template Model Builder'). When the response data are aggregated to polygon level but the predictor variables are at a higher resolution, these models can be useful. Regression models with spatial random fields. A useful reference for disaggregation modelling is Lucas et al. (2019) <doi:10.1101/548719>.

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LazyData true

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Imports maptools, raster, foreach, sp, parallel, doParallel, rgeos, splancs, rgdal, Matrix, stats, TMB, dplyr, ggplot2, cowplot, sparseMVN, utils

Additional_repositories https://inla.r-inla-download.org/R/stable

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as.disag_data

Function to fit the disaggregation model

Description

Function to fit the disaggregation model

Usage

as.disag_data(
  polygon_shapefile,
  shapefile_names,
  covariate_rasters,
  polygon_data,
  covariate_data,
  aggregation_pixels,
  coordsForFit,
  coordsForPrediction,
  startendindex,
  mesh = NULL
)
Arguments

- **polygon_shapefile**
  - SpatialPolygonsDataFrame containing the response data
- **shapefile_names**
  - List of 2: polygon id variable name and response variable name from polygon_shapefile
- **covariate_rasters**
  - RasterStack of covariates
- **polygon_data**
  - data.frame with two columns: polygon id and response
- **covariate_data**
  - data.frame with cell id, polygon id and covariate columns
- **aggregation_pixels**
  - vector with value of aggregation raster at each pixel
- **coordsForFit**
  - coordinates of the covariate data points within the polygons in polygon_shapefile
- **coordsForPrediction**
  - coordinates of the covariate data points in the whole raster extent
- **startendindex**
  - matrix containing the start and end index for each polygon
- **mesh**
  - inla.mesh object to use in the fit

Value

A list is returned of class `disag_data`. The functions `summary`, `print` and `plot` can be used on `disag_data`. The list of class `disag_data` contains:

- **polygon_shapefile**
  - The SpatialPolygonsDataFrame used as an input.
- **covariate_rasters**
  - The RasterStack used as an input.
- **polygon_data**
  - A data frame with columns of `area_id`, `response` and `N` (sample size: all NAs unless using binomial data). Each row represents a polygon.
- **covariate_data**
  - A data frame with columns of `area_id`, `cell_id` and one for each covariate in `covariate_rasters`. Each row represents a pixel in a polygon.
- **aggregation_pixels**
  - An array with the value of the aggregation raster for each pixel in the same order as the rows of `covariate_data`.
- **coordsForFit**
  - A matrix with two columns of x, y coordinates of pixels within the polygons. Used to make the spatial field.
- **coordsForPrediction**
  - A matrix with two columns of x, y coordinates of pixels in the whole Raster. Used to make predictions.
- **startendindex**
  - A matrix with two columns containing the start and end index of the pixels within each polygon.
- **mesh**
  - A INLA mesh to be used for the spatial field of the disaggregation model.
Description

`build_mesh` function takes a SpatialPolygons object and mesh arguments to build an appropriate mesh for the spatial field.

Usage

```r
build_mesh(shapes, mesh.args = NULL)
```

Arguments

- **shapes**: shapefile covering the region under investigation.
- **mesh.args**: list of parameters that control the mesh structure. `convex`, `concave` and `resolution`, to control the boundary of the inner mesh, and `max.edge`, `cut` and `offset`, to control the mesh itself, with the parameters having the same meaning as in the INLA functions `inla.convex.hull` and `inla.mesh.2d`.

Details

The mesh is created by finding a tight boundary around the polygon data, and creating a fine mesh within the boundary and a coarser mesh outside. This speeds up computation time by only having a very fine mesh within the area of interest and having a small region outside with a coarser mesh to avoid edge effects.

Six mesh parameters can be specified as arguments: `convex`, `concave` and `resolution`, to control the boundary of the inner mesh, and `max.edge`, `cut` and `offset`, to control the mesh itself, with the names meaning the same as used by INLA functions `inla.convex.hull` and `inla.mesh.2d`.

Defaults are: `pars <- list(convex = -0.01, concave = -0.5, resolution = 300, max.edge = c(3.0, 8), cut = 0.4, offset = c(1, 15))`.

Value

An inla.mesh object

Examples

```r
## Not run:
polygons <- list()
for(i in 1:100) {
  row <- ceiling(i/10)
  col <- ifelse(i %% 10 != 0, i %% 10, 10)
  xmin = 2*(col - 1); xmax = 2*col; ymin = 2*(row - 1); ymax = 2*row
  polygons[[i]] <- rbind(c(xmin, ymax), c(xmax,ymax), c(xmax, ymin), c(xmin,ymin))
}
```
polys <- do.call(raster::spPolygons, polygons)
response_df <- data.frame(area_id = 1:100, response = runif(100, min = 0, max = 10))
spdf <- sp::SpatialPolygonsDataFrame(polys, response_df)

my_mesh <- build_mesh(spdf)

## End(Not run)

---

**disaggregation-deprecated**

*Deprecated functions in disaggregation*

### Description

These functions still work but will be removed (defunct) in the next version.

### Details

- **fit_model**: This function is deprecated, and will be removed in the next version of this package.

---

**dummy**  

*Roxygen commands*

### Description

Roxygen commands

### Usage

dummy()
fit_model

Fit the disaggregation model

Description

fit_model function takes a disag_data object created by prepare_data and performs a Bayesian disaggregation fit.

Usage

fit_model(
data,  
priors = NULL, 
family = "gaussian", 
link = "identity", 
iterations = 100, 
field = TRUE, 
iid = TRUE,  
hess_control_parscale = NULL, 
hess_control_ndeps = 1e-04,  
silent = TRUE
)

disag_model(
data,  
priors = NULL, 
family = "gaussian", 
link = "identity", 
iterations = 100, 
field = TRUE, 
iid = TRUE,  
hess_control_parscale = NULL, 
hess_control_ndeps = 1e-04,  
silent = TRUE
)

Arguments

data disag_data object returned by prepare_data function that contains all the necessary objects for the model fitting

priors list of prior values

family likelihood function: gaussian, binomial or poisson

link link function: logit, log or identity

iterations number of iterations to run the optimisation for

field logical. Flag the spatial field on or off
iid logical. Flag the iid effect on or off

hess_control_parscale
Argument to scale parameters during the calculation of the Hessian. Must be the same length as the number of parameters. See optimHess for details.

hess_control_ndeps
Argument to control step sizes during the calculation of the Hessian. Either length 1 (same step size applied to all parameters) or the same length as the number of parameters. Default is 1e-3, try setting a smaller value if you get NaNs in the standard error of the parameters. See optimHess for details.

silent logical. Suppress verbose output.

Details

The model definition
The disaggregation model make predictions at the pixel level:

$$\text{link}(\text{pred}_i) = \beta_0 + \beta X + GP(s_i) + u_i$$

And then aggregates these predictions to the polygon level using the weighted sum (via the aggregation raster, \(agg_i\)):

$$\text{cases}_j = \sum_{i \in j} \text{pred}_i \times \text{agg}_i$$
$$\text{rate}_j = \frac{\sum_{i \in j} \text{pred}_i \times \text{agg}_i}{\sum_{i \in j} \text{agg}_i}$$

The different likelihood correspond to slightly different models (\(y_j\) is the response count data):

- Gaussian: If \(\sigma\) is the dispersion of the pixel data, \(\sigma_j\) is the dispersion of the polygon data, where \(\sigma_j = \sigma \sqrt{\sum agg_i^2 / \sum agg_i}\)
$$dnorm(y_j / \sum agg_i, \text{rate}_j, \sigma_j)$$
- predicts incidence rate.

- Binomial: For a survey in polygon \(j\), \(y_j\) is the number positive and \(N_j\) is the number tested.
$$dbinom(y_j, N_j, \text{rate}_j)$$
- predicts prevalence rate.

- Poisson:
$$dpois(y_j, \text{cases}_j)$$
- predicts incidence count.

Specify priors for the regression parameters, field and iid effect as a single list. Hyperpriors for the field are given as penalised complexity priors you specify \(\rho_{\text{min}}\) and \(\rho_{\text{prob}}\) for the range of the field where \(P(\rho < \rho_{\text{min}}) = \rho_{\text{prob}}\), and \(\sigma_{\text{min}}\) and \(\sigma_{\text{prob}}\) for the variation of the field where \(P(\sigma > \sigma_{\text{min}}) = \sigma_{\text{prob}}\). Also, specify pc priors for the iid effect.
The `family` and `link` arguments are used to specify the likelihood and link function respectively. The likelihood function can be one of `gaussian`, `poisson` or `binomial`. The link function can be one of `logit`, `log` or `identity`. These are specified as strings.

The field and iid effect can be turned on or off via the `field` and `iid` logical flags. Both are default `TRUE`.

The `iterations` argument specifies the maximum number of iterations the model can run for to find an optimal point.

The `silent` argument can be used to publish/supress verbose output. Default `TRUE`.

Value

A list is returned of class `disag_model`. The functions `summary`, `print` and `plot` can be used on `disag_model`. The list of class `disag_model` contains:

- `obj` The TMB model object returned by `MakeADFun`.
- `opt` The optimized model object returned by `nlminb`.
- `sd_out` The TMB object returned by `sdreport`.
- `data` The `disag_data` object used as an input to the model.
- `model_setup` A list of information on the model setup. Likelihood function (`family`), link function (`link`), logical: whether a field was used (`field`) and logical: whether an iid effect was used (`iid`).

Examples

```r
## Not run:
polygons <- list()
for(i in 1:100) {
  row <- ceiling(i/10)
  col <- ifelse(i %% 10 != 0, i %% 10, 10)
  xmin = 2*(col - 1); xmax = 2*col; ymin = 2*(row - 1); ymax = 2*row
  polygons[[i]] <- rbind(c(xmin, ymax), c(xmax,ymax), c(xmax, ymin), c(xmin,ymin))
}
polys <- do.call(raster::spPolygons, polygons)
response_df <- data.frame(area_id = 1:100, response = runif(100, min = 0, max = 10))
spdf <- sp::SpatialPolygonsDataFrame(polys, response_df)

r <- raster::raster(ncol=20, nrow=20)
r <- raster::setExtent(r, raster::extent(spdf))
r[] <- sapply(1:raster::ncell(r), function(x) rnorm(1, ifelse(x %% 20 != 0, x %% 20, 20), 3))
r2 <- raster::raster(ncol=20, nrow=20)
r2[] <- sapply(1:raster::ncell(r), function(x) rnorm(1, ceiling(x/10), 3))
cov_rasters <- raster::stack(r, r2)

c1 <- parallel::makeCluster(2)
doParallel::registerDoParallel(c1)
test_data <- prepare_data(polygon_shapefile = spdf,
                          covariate_rasters = cov_rasters)
```
getCovariateRasters

parallel::stopCluster(cl)
foreach::registerDoSEQ()

result <- fit_model(test_data, iterations = 2)

## End(Not run)

---

**getCovariateRasters**  
*Get a RasterStack of covariates from a folder containing .tif files*

**Description**

Looks in a specified folder for raster files. Returns a RasterStack of the rasters cropped to the extent specified by the `shape` parameter.

**Usage**

```r
getcovariateRasters(directory, file_pattern = "tif", shape)
```

**Arguments**

- `directory`  
  Filepath to the directory containing the rasters.
- `file_pattern`  
  Pattern the filenames must match. Default is all files ending in `.tif`.
- `shape`  
  An object with an extent that the rasters will be cropped to.

**Value**

A RasterStack of the raster files in the directory

**Examples**

```r
## Not run:
getcovariateRasters(  
  '/home/rasters',  
  '.tif',  
  shape)

## End(Not run)
```
getPolygonData

Extract polygon id and response data into a data.frame from a SpatialPolygonsDataFrame

Description

Returns a data.frame with a row for each polygon in the SpatialPolygonsDataFrame and columns: area_id, response and N, containing the id of the polygon, the values of the response for that polygon, and the sample size respectively. If the data is not survey data (the sample size does not exist), this column will contain NAs.

Usage

getPolygonData(
  shape,
  id_var = "area_id",
  response_var = "response",
  sample_size_var = NULL
)

Arguments

shape A SpatialPolygons object containing response data.
id_var Name of column in shape object with the polygon id. Default 'area_id'.
response_var Name of column in shape object with the response data. Default 'response'.
sample_size_var For survey data, name of column in SpatialPolygonsDataFrame object (if it exists) with the sample size data. Default NULL.

Value

A data.frame with a row for each polygon in the SpatialPolygonsDataFrame and columns: area_id, response and N, containing the id of the polygon, the values of the response for that polygon, and the sample size respectively. If the data is not survey data (the sample size does not exist), this column will contain NAs.

Examples

```r
polygons <- list()
for(i in 1:100) {
  row <- ceiling(i/10)
  col <- ifelse(i %% 10 != 0, i %% 10, 10)
  xmin = 2*(col - 1); xmax = 2*col; ymin = 2*(row - 1); ymax = 2*row
  polygons[[i]] <- rbind(c(xmin, ymax), c(xmax,ymax), c(xmax, ymin), c(xmin,ymin))
}
```
### getStartendindex

**Function to match pixels to their corresponding polygon**

**Description**

From the covariate data and polygon data, the function matches the polygon id between the two to find which pixels from the covariate data are contained in each of the polygons.

**Usage**

```r
getStartendindex(covariates, polygon_data, id_var = "area_id")
```

**Arguments**

- `covariates`: data.frame with each covariate as a column and id column.
- `polygon_data`: data.frame with polygon id and response data.
- `id_var`: string with the name of the column in the covariate data.frame containing the polygon id.

**Details**

Takes a data.frame containing the covariate data with a polygon id column and one column for each covariate, and another data.frame containing polygon data with a polygon id, response and sample size column (as returned by `getPolygonData` function).

Returns a matrix with two columns and one row for each polygon. The first column is the index of the first row in covariate data that corresponds to that polygon, the second column is the index of the last row in covariate data that corresponds to that polygon.

**Value**

A matrix with two columns and one row for each polygon. The first column is the index of the first row in covariate data that corresponds to that polygon, the second column is the index of the last row in covariate data that corresponds to that polygon.
Examples

```r
covs <- data.frame(area_id = c(1, 1, 1, 2, 2, 3, 3, 3, 3), response = c(3, 9, 5, 2, 3, 6, 7, 3, 5))
response <- data.frame(area_id = c(1, 2, 3), response = c(4, 7, 2), N = c(NA, NA, NA))
getStartendindex(covs, response, 'area_id')
```

Description

`make_model_object` function takes a `disag_data` object created by `prepare_data` and creates a TMB model object to be used in fitting.

Usage

```r
make_model_object(
  data, 
  priors = NULL, 
  family = "gaussian", 
  link = "identity", 
  field = TRUE, 
  iid = TRUE, 
  silent = TRUE 
)
```

Arguments

- `data`: `disag_data` object returned by `prepare_data` function that contains all the necessary objects for the model fitting
- `priors`: list of prior values
- `family`: likelihood function: `gaussian`, `binomial` or `poisson`
- `link`: link function: `logit`, `log` or `identity`
- `field`: logical. Flag the spatial field on or off
- `iid`: logical. Flag the iid effect on or off
- `silent`: logical. Suppress verbose output.
The model definition

The disaggregation model make predictions at the pixel level:

\[ \text{link}(\text{pred}_i) = \beta_0 + \beta X + GP(s_i) + u_i \]

And then aggregates these predictions to the polygon level using the weighted sum (via the aggregation raster, \(agg_i\)):

\[ \text{cases}_j = \sum_{i \in j} \text{pred}_i \times agg_i \]
\[ \text{rate}_j = \frac{\sum_{i \in j} \text{pred}_i \times agg_i}{\sum_{i \in j} agg_i} \]

The different likelihood correspond to slightly different models (\(y_j\) is the response count data):

- **Gaussian**: If \(\sigma\) is the dispersion of the pixel data, \(\sigma_j\) is the dispersion of the polygon data, where \(\sigma_j = \sigma \sqrt{\sum agg_i / \sum agg_i}\)

  \[ \text{dnorm}(y_j / \sum agg_i, \text{rate}_j, \sigma_j) \]

  - predicts incidence rate.

- **Binomial**: For a survey in polygon \(j\), \(y_j\) is the number positive and \(N_j\) is the number tested.

  \[ \text{dbinom}(y_j, N_j, \text{rate}_j) \]

  - predicts prevalence rate.

- **Poisson**: \(dpois(y_j, \text{cases}_j)\)

  - predicts incidence count.

Specify priors for the regression parameters, field and iid effect as a single list. Hyperpriors for the field are given as penalised complexity priors you specify \(\rho_{\text{min}}\) and \(\rho_{\text{prob}}\) for the range of the field where \(P(\rho < \rho_{\text{min}}) = \rho_{\text{prob}}\), and \(\sigma_{\text{min}}\) and \(\sigma_{\text{prob}}\) for the variation of the field where \(P(\sigma > \sigma_{\text{min}}) = \sigma_{\text{prob}}\). Also, specify pc priors for the iid effect.

The family and link arguments are used to specify the likelihood and link function respectively. The likelihood function can be one of gaussian, poisson or binomial. The link function can be one of logit, log or identity. These are specified as strings.

The field and iid effect can be turned on or off via the field and iid logical flags. Both are default TRUE.

The iterations argument specifies the maximum number of iterations the model can run for to find an optimal point.

The silent argument can be used to publish/supress verbose output. Default TRUE.

Value

The TMB model object returned by MakeADFun.
### Examples

```r
## Not run:
polygons <- list()
for(i in 1:100) {
  row <- ceiling(i/10)
  col <- ifelse(i %% 10 != 0, i %% 10, 10)
  xmin = 2*(col - 1); xmax = 2*col; ymin = 2*(row - 1); ymax = 2*row
  polygons[[i]] <- rbind(c(xmin, ymax), c(xmax,ymax), c(xmax, ymin), c(xmin,ymin))
}

polys <- do.call(raster::spPolygons, polygons)
response_df <- data.frame(area_id = 1:100, response = runif(100, min = 0, max = 10))
spdf <- sp::SpatialPolygonsDataFrame(polys, response_df)

r <- raster::raster(ncol=20, nrow=20)
r <- raster::setExtent(r, raster::extent(spdf))
r[] <- sapply(1:raster::ncell(r), function(x) rnorm(1, ifelse(x %% 20 != 0, x %% 20, 20), 3))
r2 <- raster::raster(ncol=20, nrow=20)
r2 <- raster::setExtent(r2, raster::extent(spdf))
r2[] <- sapply(1:raster::ncell(r), function(x) rnorm(1, ceiling(x/10), 3))
cov_rasters <- raster::stack(r, r2)

c1 <- parallel::makeCluster(2)
doParallel::registerDoParallel(cl)
test_data <- prepare_data(polygon_shapefile = spdf, coariate_rasters = cov_rasters)
parallel::stopCluster(cl)
foreach::registerDoSEQ()

result <- make_model_object(test_data)
## End(Not run)
```

---

**parallelExtract**

*Parallel extraction of raster stack by shape file.*

### Description

Parallelisation is performed across rasters, not shapes. So this function is only useful if you are extracting data from many raster layers. As the overhead for parallel computation in windows is high it only makes sense to parallelise in this way.

### Usage

```r
parallelExtract(raster, shape, fun = mean, id = "OBJECTID", ...)
```
Arguments

- **raster**: A RasterBrick or RasterStack object.
- **shape**: A SpatialPolygons object.
- **fun**: The function used to aggregate the pixel data. If NULL, raw pixel data is returned.
- **id**: Name of column in shape object to be used to bind an ID column to output.
- **...**: Other arguments to raster::extract.

Value

A data.frame with columns of polygon id, cell id (if fun = NULL) and a column for each raster in the stack.

Examples

```r
## Not run:
polygons <- list()
for(i in 1:100) {
  row <- ceiling(i/10)
  col <- ifelse(i %% 10 != 0, i %% 10, 10)
  xmin = 2*(col - 1); xmax = 2*col; ymin = 2*(row - 1); ymax = 2*row
  polygons[[i]] <- rbind(c(xmin, ymax), c(xmax,ymax), c(xmax, ymin), c(xmin,ymin))
}
polys <- do.call(raster::spPolygons, polygons)
response_df <- data.frame(area_id = 1:100, response = runif(100, min = 0, max = 10))
spdf <- sp::SpatialPolygonsDataFrame(polys, response_df)
r <- raster::raster(ncol=20, nrow=20)
r <- raster::setExtent(r, raster::extent(spdf))
r[] <- sapply(1:raster::ncell(r), function(x) rnorm(1, ifelse(x %% 20 != 0, x %% 20, 20), 3))
r2 <- raster::raster(ncol=20, nrow=20)
r2 <- raster::setExtent(r2, raster::extent(spdf))
r2[] <- sapply(1:raster::ncell(r), function(x) rnorm(1, ceiling(x/10), 3))
cov_rasters <- raster::stack(r, r2)

cl <- parallel::makeCluster(2)
doParallel::registerDoParallel(cl)
result <- parallelExtract(cov_rasters, spdf, fun = NULL, id = 'area_id')
parallel::stopCluster(cl)
foreach::registerDoSEQ()

## End(Not run)
```
plot.disag_data  *Plot input data for disaggregation*

**Description**
Plotting function for class *disag_data* (the input data for disaggregation).

**Usage**
```r
## S3 method for class 'disag_data'
plot(x, which = c(1, 2, 3), ...)
```

**Arguments**
- `x`: Object of class *disag_data* to be plotted.
- `which`: If a subset of plots is required, specify a subset of the numbers 1:3.
- `...`: Further arguments to `plot` function.

**Details**
Produces three plots: polygon response data, covariate rasters and INLA mesh.

**Value**
A list of three plots: the polygon plot (ggplot), covariate plot (spplot) and INLA mesh plot (ggplot).

plot.disag_model  *Plot results of fitted model*

**Description**
Plotting function for class *disag_model* (the result of the disaggregation fitting).

**Usage**
```r
## S3 method for class 'disag_model'
plot(x, ...)
```

**Arguments**
- `x`: Object of class *disag_model* to be plotted.
- `...`: Further arguments to `plot` function.

**Details**
Produces two plots: results of the fixed effects and in-sample observed vs predicted plot.
plot.disag_prediction

Value
A list of two ggplot plots: results of the fixed effects and an in-sample observed vs predicted plot

Description
Plot mean and uncertainty predictions from the disaggregation model results

Usage
## S3 method for class 'disag_prediction'
plot(x, ...)

Arguments
x Object of class disag_prediction to be plotted.
... Further arguments to plot function.

Details
Produces raster plots of the mean prediction, and the lower and upper confidence intervals.

Value
A list of plots of rasters from the prediction: mean prediction, lower CI and upper CI.

predict.disag_model

Description
predict.disag_model function takes a disag_model object created by disaggregation::disag_model and predicts mean and uncertainty maps.

Usage
## S3 method for class 'disag_model'
predict(object, newdata = NULL, predict_iid = FALSE, N = 100, CI = 0.95, ...)

predict.disag_model

Description
Predict mean and uncertainty from the disaggregation model result

Usage
## S3 method for class 'disag_model'
predict(object, newdata = NULL, predict_iid = FALSE, N = 100, CI = 0.95, ...)

predict.disag_model
Arguments

- **object**: disag_model object returned by disag_model function.
- **newdata**: If NULL, predictions are made using the data in model_output. If this is a raster stack or brick, predictions will be made over this data.
- **predict_iid**: logical. If TRUE, any polygon iid effect from the model will be used in the prediction. Default FALSE.
- **N**: Number of realisations. Default: 100.
- **CI**: Confidence interval to be calculated from the realisations. Default: 0.95.
- **...**: Further arguments passed to or from other methods.

Details

To predict over a different spatial extent to that used in the model, a RasterStack covering the region to make predictions over is passed to the argument **newdata**. If this is not given predictions are made over the data used in the fit.

The **predict_iid** logical flag should be set to TRUE if the results of the iid effect from the model are to be used in the prediction.

For the uncertainty calculations, the number of the realisations and the size of the confidence interval to be calculated are given by the arguments **N** and **CI** respectively.

Value

An object of class **disag_prediction** which consists of a list of two objects:

- **mean_prediction**: List of:
  - **prediction**: Raster of mean predictions based.
  - **field**: Raster of the field component of the linear predictor.
  - **iid**: Raster of the iid component of the linear predictor.
  - **covariates**: Raster of the covariate component of the linear predictor.

- **uncertainty_prediction**: List of:
  - **realisations**: RasterStack of realisations of predictions. Number of realisations defined by argument **N**.
  - **predictions_ci**: RasterStack of the upper and lower credible intervals. Defined by argument **CI**.

Examples

```r
## Not run:
predict(fit_result)

## End(Not run)
```
predict_model

Function to predict mean from the model result

Description

`predict_model` function takes a `disag_model` object created by `disaggregation::disag_model` and predicts mean maps.

Usage

```r
predict_model(model_output, newdata = NULL, predict_iid = FALSE)
```

Arguments

- `model_output`: `disag_model` object returned by `disag_model` function
- `newdata`: If NULL, predictions are made using the data in `model_output`. If this is a raster stack or brick, predictions will be made over this data. Default NULL.
- `predict_iid`: If TRUE, any polygon iid effect from the model will be used in the prediction. Default FALSE.

Details

Function returns rasters of the mean predictions as well as the covariate and field contributions to the linear predictor.

To predict over a different spatial extent to that used in the model, a RasterStack covering the region to make predictions over is passed to the argument `newdata`. If this is not given predictions are made over the data used in the fit.

The `predict_iid` logical flag should be set to TRUE if the results of the iid effect from the model are to be used in the prediction.

Value

The mean prediction, which is a list of:

- `prediction`: Raster of mean predictions based.
- `field`: Raster of the field component of the linear predictor.
- `iid`: Raster of the iid component of the linear predictor.
- `covariates`: Raster of the covariate component of the linear predictor.

Examples

```r
## Not run:
predict_model(result)
```

## End(Not run)
predict_uncertainty  

Function to predict uncertainty from the model result

Description

predict_uncertainty function takes a disag_model object created by disaggregation::disag_model and predicts upper and lower credible interval maps.

Usage

predict_uncertainty(
  model_output,  
  newdata = NULL,  
  predict_iid = FALSE,  
  N = 100,  
  CI = 0.95
)

Arguments

model_output  
  disag_model object returned by disag_model function.
newdata  
  If NULL, predictions are made using the data in model_output. If this is a raster stack or brick, predictions will be made over this data. Default NULL.
predict_iid  
  If TRUE, any polygon iid effect from the model will be used in the prediction. Default FALSE.
N  
  number of realisations. Default: 100.
CI  
  confidence interval. Default: 0.95.

Details

Function returns a RasterStack of the realisations as well as the upper and lower credible interval rasters.

To predict over a different spatial extent to that used in the model, a RasterStack covering the region to make predictions over is passed to the argument newdata. If this is not given predictions are made over the data used in the fit.

The predict_iid logical flag should be set to TRUE if the results of the iid effect from the model are to be used in the prediction.

The number of the realisations and the size of the confidence interval to be calculated are given by the arguments N and CI respectively.

Value

The uncertainty prediction, which is a list of:

• realisations RasterStack of realisations of predictions. Number of realisations defined by argument N.
• predictions_ci RasterStack of the upper and lower credible intervals. Defined by argument CI.
prepare_data

Examples

```r
## Not run:
predict_uncertainty(result)
## End(Not run)
```

prepare_data

## Not run: predict_uncertainty(result)
## End(Not run)

### Description

The `prepare_data` function is used to extract all the data required for fitting a disaggregation model. Designed to be used in the `disaggregation::fit_model` function.

### Usage

```r
prepare_data(
  polygon_shapefile, 
  covariate_rasters, 
  aggregation_raster = NULL, 
  id_var = "area_id", 
  response_var = "response", 
  sample_size_var = NULL, 
  mesh.args = NULL, 
  na.action = FALSE, 
  makeMesh = TRUE, 
  ncores = 2
)
```

### Arguments

- **polygon_shapefile**: SpatialPolygonDataFrame containing at least two columns: one with the id for the polygons (`id_var`) and one with the response count data (`response_var`); for binomial data, i.e., survey data, it can also contain a sample size column (`sample_size_var`).
- **covariate_rasters**: RasterStack of covariate rasters to be used in the model.
- **aggregation_raster**: Raster to aggregate pixel level predictions to polygon level e.g. population to aggregate prevalence. If this is not supplied a uniform raster will be used.
- **id_var**: Name of column in SpatialPolygonDataFrame object with the polygon id.
- **response_var**: Name of column in SpatialPolygonDataFrame object with the response data.
- **sample_size_var**: For survey data, name of column in SpatialPolygonDataFrame object (if it exists) with the sample size data.
prepare_data

mesh.args  list of parameters that control the mesh structure with the same names as used by INLA.

na.action  logical. If TRUE, NAs in response will be removed, covariate NAs will be given the median value, aggregation NAs will be set to zero. Default FALSE (NAs in response or covariate data within the polygons will give errors).

makeMesh  logical. If TRUE, build INLA mesh, takes some time. Default TRUE.

ncores  Number of cores used to perform covariate extraction.

Details

Takes a SpatialPolygonDataFrame with the response data and a RasterStack of covariates. Extract the values of the covariates (as well as the aggregation raster, if given) at each pixel within the polygons (parallelExtract function). This is done in parallel and n.cores argument is used to set the number of cores to use for covariate extraction. This can be the number of covariates used in the model.

The aggregation raster defines how the pixels within each polygon are aggregated. The disaggregation model performs a weighted sum of the pixel prediction, weighted by the pixel values in the aggregation raster. For disease incidence rate you use the population raster to aggregate pixel incidence rate by summing the number of cases (rate weighted by population). If no aggregation raster is provided a uniform distribution is assumed, i.e. the pixel predictions are aggregated to polygon level by summing the pixel values.

Makes a matrix that contains the start and end pixel index for each polygon. Builds an INLA mesh to use for the spatial field (getStartendindex function).

The mesh.args argument allows you to supply a list of INLA mesh parameters to control the mesh used for the spatial field (build_mesh function).

The na.action flag is automatically off. If there are any NAs in the response or covariate data within the polygons the prepare_data method will error. Ideally the NAs in the data would be dealt with beforehand, however, setting na.action = TRUE will automatically deal with NAs. It removes any polygons that have NAs as a response, sets any aggregation pixels with NA to zero and sets covariate NAs pixels to the median value for the that covariate.

Value

A list is returned of class disag_data. The functions summary, print and plot can be used on disag_data. The list of class disag_data contains:

polygon_shapefile  The SpatialPolygonDataFrame used as an input.

covariate_rasters  The RasterStack used as an input.

disag_data  A data frame with columns of area_id, response and N (sample size: all NAs unless using binomial data). Each row represents a polygon.

covariate_data  A data frame with columns of area_id, cell_id and one for each covariate in covariate_rasters. Each row represents a pixel in a polygon.
**aggregation_pixels**

An array with the value of the aggregation raster for each pixel in the same order as the rows of `covariate_data`.

**coordsForFit**

A matrix with two columns of x, y coordinates of pixels within the polygons. Used to make the spatial field.

**coordsForPrediction**

A matrix with two columns of x, y coordinates of pixels in the whole Raster. Used to make predictions.

**startIndex**

A matrix with two columns containing the start and end index of the pixels within each polygon.

**mesh**

A INLA mesh to be used for the spatial field of the disaggregation model.

---

**Examples**

```r
polygons <- list()
for(i in 1:100) {
  row <- ceiling(i/10)
  col <- ifelse(i %% 10 != 0, i %% 10, 10)
  xmin = 2*(col - 1); xmax = 2*col; ymin = 2*(row - 1); ymax = 2*row
  polygons[[i]] <- rbind(c(xmin, ymax), c(xmax,ymax), c(xmax, ymin), c(xmin,ymin))
}
polys <- do.call(raster::spPolygons, polygons)
response_df <- data.frame(area_id = 1:100, response = runif(100, min = 0, max = 10))
spdf <- sp::SpatialPolygonsDataFrame(polys, response_df)

r <- raster::raster(ncol=20, nrow=20)
r <- raster::setExtent(r, raster::extent(spdf))
r[] <- sapply(1:raster::ncell(r), function(x) rnorm(1, ifelse(x %% 20 != 0, x %% 20, 20), 3))
r2 <- raster::raster(ncol=20, nrow=20)
r2 <- raster::setExtent(r2, raster::extent(spdf))
r2[] <- sapply(1:raster::ncell(r), function(x) rnorm(1, ceiling(x/10), 3))
cov_rasters <- raster::stack(r, r2)
test_data <- prepare_data(polygon_shapefile = spdf,
                          covariate_rasters = cov_rasters)
```

---

**print.disag_data**

`Print function for disaggregation input data`

**Description**

Function that prints the input data from the disaggregation model.
Print function for decomposition fit result.

Function that prints the result of the fit from the decomposition model.

Usage

```r
## S3 method for class 'disag_model'
print(x, ...)

Arguments

x
Object returned from disag_model.

... Further arguments to print function.

Details

Prints the negative log likelihood, model parameters and calculates metrics from in-sample performance.
**print.disag_prediction**

*Print function for disaggregation prediction*

**Description**

Function that prints the prediction from the disaggregation model.

**Usage**

```r
## S3 method for class 'disag_prediction'
print(x, ...)
```

**Arguments**

- `x`: Object returned from predict.disag_model.
- `...`: Further arguments to `print` function.

**Details**

Prints the number of polyons and pixels, the number of pixels in the largest and smallest polygons and summaries of the covariates.

---

**summary.disag_data**

*Summary function for disaggregation input data*

**Description**

Function that summarizes the input data from the disaggregation model.

**Usage**

```r
## S3 method for class 'disag_data'
summary(object, ...)
```

**Arguments**

- `object`: Object returned from prepare_data.
- `...`: Further arguments to `summary` function.

**Details**

Prints the number of polyons and pixels, the number of pixels in the largest and smallest polygons and summaries of the covariates.

**Value**

A list of the number of polyons, the number of covariates and summaries of the covariates.
summary.disag_model  Summary function for disaggregation fit result

Description
Function that summarises the result of the fit from the disaggregation model.

Usage
```r
## S3 method for class 'disag_model'
summary(object, ...)
```

Arguments
- `object`: Object returned from `disag_model`.
- `...`: Further arguments to `summary` function.

Details
Prints the negative log likelihood, model parameters and calculates metrics from in-sample performance.

Value
A list of the model parameters, negative log likelihood and metrics from in-sample performance.

summary.disag_prediction  Summary function for disaggregation prediction

Description
Function that summarizes the prediction from the disaggregation model.

Usage
```r
## S3 method for class 'disag_prediction'
summary(object, ...)
```

Arguments
- `object`: Object returned from `predict.disag_model`.
- `...`: Further arguments to `summary` function.
summary.disag_prediction

Details

Prints the number of polygons and pixels, the number of pixels in the largest and smallest polygons and summaries of the covariates.

Value

A list of the number of polygons, the number of covariates and summaries of the covariates.
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