

Package ‘gDefrag’

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Author Frederico Mestre, Fernando Ascensao, A. Marcia Barbosa

Maintainer Frederico Mestre <mestre.frederico@gmail.com>

Description Provides a set of tools to help the de-fragmentation process. It works by prioritizing the different sections of linear infrastructures (e.g. roads, power-lines) to increase the available amount of a given resource.

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Description

Provides a set of tools to help the de-fragmentation process. It works by prioritizing the different sections of linear infrastructures (e.g. roads, power-lines) to increase the available amount of a given resource.

Details

Our approach is based on graph-based tools, and relies only on the spatial configuration of the infrastructures and the layer of the resource to be evaluated (e.g. habitat area or quality). The package interprets the landscape as a graph-like structure, taking advantage of the structural simplicity of this approach. The main assumption is that connecting a larger amount of a given resource is benefit for the species persistence, following the concept of 'accessible habitat' by Eigenbrod et al. (2008). It then prioritizes the sections that have higher impact in overall connectivity according to three methods, including the Integral Index of Connectivity (IIC) developed by Pascual-Hortal and Saura (2006). It can be used to assist in the de-fragmentation process caused by any linear infrastructure type, including roads, railways, power lines, pipelines, water channels, and fences. Hereafter we will refer to these simply as 'roads'.

Author(s)

Frederico Mestre, Fernando Ascensao, A. Marcia Barbosa

Maintainer: Frederico Mestre <mestre.frederico@gmail.com>

References

- Eigenbrod, F., Hecnar, S.J., Fahrig, L. (2008). Accessible habitat: an improved measure of the effects of habitat loss and roads on wildlife populations. *Landscape Ecology*, 23(2), 159-168.
- Grilo, C., Ascensao, F., Santos-Reis, M., Bissonette, J.A. (2011). Do well-connected landscapes promote road-related mortality?. *European Journal of Wildlife Research*, 57(4), 707-716.
- Pascual-Hortal, L., Saura, S. (2006). Comparison and development of new graph-based landscape connectivity indices: towards the prioritization of habitat patches and corridors for conservation. *Landscape Ecology*, 21(7), 959-967.
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edge.creation	<i>Creating graph edges</i>
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Description

Generates the graph edges, connecting nodes that are the landscape polygons resulting from the landscape clip by the road network.

Usage

```
edge.creation(nodes, land_polyg, min_length = 0,  
min_pol_area = 0, plot = TRUE, shape = FALSE,  
shape_name_edges = "shape_edges")
```

Arguments

nodes	Output of the function node.creation .
land_polyg	SpatialPolygonsDataFrame loaded with readOGR function of the package rgdal (polygon shapefile).
min_length	Minimum length of the road segment to consider creating an edge representing it.
min_pol_area	Minimum area for a given polygon to be considered a node.
plot	Plot after running? (TRUE/FALSE).
shape	Create a shapefile with the edges (TRUE/FALSE).
shape_name_edges	Name of the edges shapefile to be created (without prioritization).

Details

If shape = TRUE, this function produces one shapefile with edges without prioritization metrics.

Value

The function returns a SpatialLinesDataFrame with the edges. The data table of this data frame has the following fields:

- node_A - Node A Id.
- node_B - Node B Id.
- distance - Euclidean distance between nodes A and B (edge length).
- x_node_A - Node A longitude.
- y_node_A - Node A latitude.
- x_node_B - Node B longitude.
- y_node_A - Node A latitude.

- raster_value_A - Average value of the underlying raster in the polygon represented by node A.
- raster_value_B - Average value of the underlying raster in the polygon represented by node B.
- road_ID - Id of the road represented by the edge.
- road_length - Length of the road represented by the edge.

Author(s)

Frederico Mestre, Fernando Ascensao and A. Marcia Barbosa

See Also

[node.creation](#)

Examples

```
data(road_P)

out1 <- node.creation(land_polyg=road_P, value_col = "forest_sum", scale_nodes = 10,
col_nodes = "pink", cex_labels = 0.7)

out2 <- edge.creation(nodes = out1, land_polyg = road_P, min_length = 0,
min_pol_area = 0, plot = FALSE)
```

gDefrag.full

Wrapper function running the full process

Description

This function runs the full process of creating the nodes, creating the edges and establishing priority edges to be used to enhance connectivity in the landscape for the focal species.

Usage

```
gDefrag.full(land_polyg, method, value_col = NULL,
min_length = 0, min_pol_area = 0, shape = FALSE,
shape_name_nodes = "shape_all_nodes", shape_name_edges = "shape_edges",
shape_name_out = "priorities_shape",
shape_name_nodes_edges = "nodes_with_edges",
scale_nodes = 10, col_nodes = "deepskyblue4", col_labels = "white",
cex_labels = 1, main = "Graph")
```

Arguments

land_polyg	SpatialPolygonsDataFrame loaded with readOGR function of the package rgdal (polygon shapefile). To be internally passed to as <code>node.creation</code> and as <code>edge.creation</code> .
method	Method to be used in the prioritization of the connectivity corridors. Detailed description in the 'details' section. To be internally passed to as <code>prioritize</code> .
value_col	Identification of the column in the 'land_polyg' object. To be internally passed to as <code>prioritize</code> .
min_length	Minimum length of the road segment to consider creating an edge representing it. To be internally passed to as <code>edge.creation</code> .
min_pol_area	Minimum area of the polygon to consider creating a node representing it. To be internally passed to as <code>node.creation</code> .
shape	Produce node and edges shapefiles as output? (TRUE/FALSE) To be internally passed to as <code>node.creation</code> and as <code>edge.creation</code> .
shape_name_nodes	Name of the nodes shapefile to be created (all nodes).
shape_name_edges	Name of the edges shapefile to be created (without prioritization).
shape_name_out	Name of the edges prioritization output shapefile to be created.
shape_name_nodes_edges	Instead of 'shape_name_nodes', that depicts all the nodes, this shape has only nodes considered in the prioritization procedure (those with an area greater than the argument 'min_pol_area' in the function <code>edge.creation</code>).
scale_nodes	Node size scaling factor for correct viewing in the plot. To be internally passed to as <code>plotgraph</code> .
col_nodes	Node colour. To be internally passed to as <code>plotgraph</code> .
col_labels	Labels colour. To be internally passed to as <code>plotgraph</code> .
cex_labels	Text labels font size. To be internally passed to as <code>plotgraph</code> .
main	Main plot title. To be internally passed to as <code>plotgraph</code> .

Details

If `shape = TRUE`, this function produces four shapefiles: all nodes, nodes with area greater than 'min_pol_area', edges without prioritization metrics, edges with prioritization metrics. The prioritization of the connectivity corridors can be made with one of four methods (argument 'method'):

- 'value' - Give priority to edges connecting nodes with higher attribute value (e.g. suitability for the focal species);
- 'traffic' - Give priority to edges representing roads with more traffic;
- 'IIC' - Give priority to edges with higher overall impact in the connectivity. This was implemented by using the Integral Index of Connectivity (IIC) developed by Pascual-Hortal and Saura (2006). The connectivity relevance of each edge was evaluated using the approach described in Pascual-Hortal and Saura (2006): $dIIC = ((I1 - I2) / I1) * 100$, where I1 is the IIC before edge removal and I2 is the IIC after.
- 'between' - Give priority to edges with higher betweenness using the function `edge.betweenness.estimate` from the `igraph` package.
- 'AWM' - Give priority to edges connecting nodes with more suitable area.

Value

This function produces a 'list' object. This list has two objects, one with the information of the nodes and the other with the information on the edges.

Author(s)

Frederico Mestre, Fernando Ascensao and A. Marcia Barbosa

See Also

[node.creation](#), [edge.creation](#), [plotgraph](#), [prioritize](#)

Examples

```
data(road_P)

out1 <- gDefrag.full(land_polyg = road_P, method = "value",
  value_col = "forest_sum", main = "value-based graph",
  shape = FALSE)
```

node.creation	<i>Creating graph nodes</i>
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Description

Generates the nodes that are the landscape polygons resulting from the landscape clip by infrastructure network.

Usage

```
node.creation(land_polyg, value_col, plot = TRUE,
  scale_nodes = 1, col_nodes = "deepskyblue4", cex_labels = 1,
  shape = FALSE, shape_name_nodes = "shape_nodes")
```

Arguments

land_polyg	SpatialPolygonsDataFrame loaded with readOGR function of the package rgdal (polygon shapefile).
value_col	Which column in the SpatialPolygonsDataFrame has the relevant information on the underlying raster. This information must be retrieved previously using a GIS, if the user wishes to use the method 'value' for prioritizing edges.
plot	Plot the output (TRUE/FALSE).
scale_nodes	Scaling factor to represent the nodes graphically.
col_nodes	Nodes colour.

cex_labels Font size of the labels.
shape Create a shapefile with the edges (TRUE/FALSE).
shape_name_nodes Name of the nodes shapefile to be created (all nodes).

Details

If shape = TRUE, this function produces one shapefile with all the nodes.

Value

The function returns a SpatialPointsDataFrame with the nodes. The data table of this data frame has the following fields:

- X - Node latitude.
- Y - Node longitude.
- pol_value - Underlying raster value summed for each polygon.
- pol_area - Polygon area.

Author(s)

Frederico Mestre, Fernando Ascensao and A. Marcia Barbosa

See Also

[edge.creation](#)

Examples

```
data(road_P)

out1 <- node.creation(land_polyg = road_P, value_col = "forest_sum",
  scale_nodes = 10, col_nodes = "pink", cex_labels = 1)
```

plotgraph

Plotting the spatial graph

Description

Function to plot the spatial graph

Usage

```
plotgraph(nodes, edges, land_polyg, scale_nodes = 10, col_nodes = "darkblue",
  col_labels = "darkblue", cex_labels = 1, main = "Graph")
```

Arguments

nodes	Output of node.creation .
edges	Output of edge.creation .
land_polyg	SpatialPolygonsDataFrame with the polygons defined by roads and the study site limits.
scale_nodes	Node size scaling factor for correct viewing in the plot.
col_nodes	Node colour.
col_labels	Labels colour.
cex_labels	Text labels font size.
main	Main plot title.

Details

Produces a plot of the study site, nodes and edges with the prioritization order.

Value

This function plots the spatial graph.

Author(s)

Frederico Mestre, Fernando Ascensao and A. Marcia Barbosa

See Also

[node.creation](#), [edge.creation](#), [prioritize](#), [road_P](#)

Examples

```
data(road_P)

#Obtaining nodes
out1 <- node.creation(land_polyg = road_P, value_col = "forest_sum",
  scale_nodes = 10, col_nodes = "pink", cex_labels = 1)

#Obtaining edges
out2 <- edge.creation(nodes = out1, land_polyg = road_P,
  min_length = 0, min_pol_area = 0)

#Prioritize
out3 <- prioritize(nodes = out1, edges = out2, method = "value")

#Plotting results
plotgraph(nodes = out1, edges = out3, land_polyg = road_P, main = "Habitat value")
```

prioritize *Prioritizing connectivity corridors*

Description

Function to establish a priority on the edges (connections) to be created in order to de-fragment a landscape.

Usage

```
prioritize(nodes, edges, method, shape=FALSE, shape_name_out = "priorities_shape",
           shape_name_nodes_edges = "nodes_with_edges")
```

Arguments

nodes	Output of node.creation .
edges	Output of edge.creation .
method	Method to be used in the prioritization of the connectivity corridors. Detailed description in the 'details' section.
shape	Create a shapefiles? (TRUE/FALSE).
shape_name_out	Name of the shapefile to be created.
shape_name_nodes_edges	Name of the nodes shapefile to be created. This shape has only nodes considered in the prioritization procedure (those with an area greater than the argument 'min_pol_area' in the function edge.creation).

Details

If `shape = TRUE`, this function produces two shapefiles: nodes with area greater than 'min_pol_area' (in [edge.creation](#)), edges with prioritization metrics. The prioritization of the connectivity corridors can be made with one of these methods (argument 'method'):

- 'value' - Give priority to edges connecting nodes with higher attribute value (e.g. suitability for the focal species);
- 'IIC' - Give priority to edges with higher overall impact in the connectivity. This was implemented by using the Integral Index of Connectivity (IIC) developed by Pascual-Hortal and Saura (2006). The connectivity relevance of each edge was evaluated using the approach described in Pascual-Hortal and Saura (2006): $dIIC = ((I1 - I2) / I1) * 100$, where I1 is the IIC before edge removal and I2 is the IIC after.
- 'between' - Give priority to edges with higher betweenness using the function `edge.betweenness.estimate` from the `igraph` package.
- 'AWM' - Area Weighted Metric - The user should provide the value of the variable to be weighted by area (e.g. habitat area) in the 'value_col' argument of the function [node.creation](#). The expression for the Area Weighted Metric is: $AWM = (value_A * area_B) + (value_B * area_A)$, where: 'value_A' and 'area_B' are the node A argument and the habitat in polygon B, respectively and 'value_B' and 'area_A' are the node B argument and the habitat in polygon A, respectively.

Value

The output is a `SpatialLinesDataFrame` with an additional field, 'priorization'. This field provides the desired prioritization value.

Note

The computation of IIC is made resorting to the node attribute in place of the 'habitat patch areas' (ai and aj) as originally devised by Pascual-Hortal & Saura (2006).

Author(s)

Frederico Mestre, Fernando Ascensao and A. Marcia Barbosa

References

Pascual-Hortal, L., Saura, S. (2006). Comparison and development of new graph-based landscape connectivity indices: towards the prioritization of habitat patches and corridors for conservation. *Landscape Ecology*, 21(7), 959-967.

See Also

[node.creation](#), [edge.creation](#)

Examples

```
data(road_P)

##### To the value, betweenness and connectivity metrics #####

#Obtaining nodes
out1 <- node.creation(land_polyg = road_P, value_col = "forest_sum",
scale_nodes = 10, col_nodes = "pink", cex_labels = 1)

#Obtaining edges
out2 <- edge.creation(nodes = out1, land_polyg = road_P,
min_length = 0, min_pol_area = 0)

#Prioritize
out3 <- prioritize(nodes = out1, edges = out2,
method = "value")#Raster value
out4 <- prioritize(nodes = out1, edges = out2,
method = "between")#Betweenness
out5 <- prioritize(nodes = out1, edges = out2,
method = "IIC") #Connectivity

##### To the area weighted metric #####

#Obtaining nodes
out6 <- node.creation(land_polyg = road_P, value_col = "proportion",
scale_nodes = 10, col_nodes = "pink", cex_labels = 1)
```

```
#Obtaining edges
out7 <- edge.creation(nodes = out6, land_polyg = road_P,
min_length = 0, min_pol_area = 0)

#Prioritize
out8 <- prioritize(nodes = out6, edges = out7,
method = "AWM")#Habitat proportion
```

road_P

Road polygons dataset with habitat area

Description

Sample dataset. SpatialPolygonsDataFrame with the polygons defined by roads and the study site limits. The field characterizing the raster is the habitat area in the polygon.

Usage

```
data("road_P")
```

Format

An object of the class 'SpatialPolygonsDataFrame' [package "sp"] with six features. The argument table should have a column with the values of an underlying raster (e.g. averaged or summed).

Details

This dataset has the spatial information on the polygons in the sample study site. The polygons are defined by the roads and the limits of the sample study site (Odemira Municipality, Portugal). The field "forest_amount" is to be used with method = 'hab_prop' in the function `prioritize`, a new metric that accounts for habitat proportion in each polygon. However, it can also be used with method = 'value' or 'IIC'. The field "forest_sum" can not be used with the method = "hab_prop" of the function `prioritize`, since this method uses only the ratio of suitable habitat. It is intended for use with the methods 'IIC' and 'value'.

See Also

[node.creation](#), [edge.creation](#), [prioritize](#)

Examples

```
data(road_P)
str(road_P)
```

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