

Package ‘rsMove’

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Description Tools to analyze animal tracking data with remote sensing.

LazyData TRUE

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backSample	<i>backSample</i>
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Description

Background sample selection.

Usage

```
backSample(xy = xy, rid = rid, method = method, img = img, nb = NULL)
```

Arguments

<code>xy</code>	Object of class <i>SpatialPoints</i> of <i>SpatialPointsDataFrame</i> .
<code>rid</code>	Vector of region identifiers for each sample.
<code>method</code>	One of <i>random</i> or <i>pca</i> . Default is <i>random</i> .
<code>img</code>	Object of class <i>RasterLayer</i> , <i>RasterStack</i> or <i>RasterBrick</i> .
<code>nb</code>	Number of random background samples.

Details

The function selects a set of n background samples where n is determined by *nb*. If *nb* is not provided all background pixels are considered. If the method is *random* these samples are returned in the form of a *SpatialPoints* object. If *pca* is used, a PCA analysis is performed over all available samples (provided and random). For each unique region ID (*rid*) the median and the Median Absolute Deviation (MAD) is estimated for a given Principal Component (PC). Background samples where the difference between their variance and the variance of the region samples exceeds the MAD are selected. Only the samples that are selected by all unique regions are kept. This process is repeated for all PC's with eigenvalues greater than 1 (Kaiser rule). The final set of samples corresponds to the unique records selected by the different PC's. These samples are returned as a *SpatialPointsDataFrame* containing the variables extracted from *img*.

Value

A *SpatialPoints* or a *SpatialPointsDataFrame*.

See Also

[labelSample](#) [hotMove](#) [dataQuery](#)

Examples

```
{
  require(raster)

  # read raster data
  file <- list.files(system.file('extdata', '', package="rsMove"), 'tc.*tif', full.names=TRUE)
  rsStk <- stack(file)

  # read movement data
  moveData <- read.csv(system.file('extdata', 'konstanz_20130805-20130811.csv', package="rsMove"))
  moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(rsStk))

  # find sample regions
  label <- labelSample(xy=moveData, rad=500, npx=2, pxr=30)

  # select background samples
  ind <- which(label>0) # selected samples
  bSamples <- backSample(xy=moveData[ind,], rid=label[ind], img=rsStk, method='random')
}
```

dataQuery

dataQuery

Description

Query environmental data for coordinate pairs.

Usage

```
dataQuery(xy = xy, st = NULL, img = img, rt = NULL, tb = NULL,
          bs = NULL, fun = NULL)
```

Arguments

xy	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
st	Object of class <i>Date</i> with <i>xy</i> observation dates.
img	Object of class <i>RasterLayer</i> , <i>RasterStack</i> or <i>RasterBrick</i> .
rt	Object of class <i>Date</i> with <i>img</i> observation dates.

tb	Two element vector with temporal search buffer, expressed in days.
bs	Buffer size (unit depends on the raster projection).
fun	Passes an external function.

Details

Returns environmental variables from a raster object for a given set of x and y coordinates. A buffer size (*bs*) and a user defined function (*fun*) can be specified to sample within an area. The default is to estimate a weighted mean. If raster acquisition times are provided (*rt*) and the date of sampling (*st*). In this case, the function will treat the raster data as a time series and search for clear pixel in time within the constraints of a temporal buffer defined by *tb*. *tb* passes two values which represent the size of the buffer in two directions: before and after the target date. This allows for backward and forward sampling.

Value

A n object of class *vector* or *data.frame*.

See Also

[sampleMove](#) [backSample](#)

Examples

```
{
  require(raster)

  # read raster data
  file <- list.files(system.file('extdata', '', package="rsMove"), 'tc.*tif', full.names=TRUE)
  rsStk <- stack(file)
  rsStk <- stack(rsStk, rsStk, rsStk) # dummy files for the example

  # read movement data
  moveData <- read.csv(system.file('extdata', 'konstanz_20130805-20130811.csv', package="rsMove"))
  moveData <- SpatialPointsDataFrame(moveData[1:10,1:2], moveData[1:10,], proj4string=crs(rsStk))

  # raster dates
  r.date <- seq.Date(as.Date("2013-08-01"), as.Date("2013-08-09"), 1)

  # sample dates
  o.date <- as.Date(moveData@data$date)

  # retrieve remote sensing data for samples
  rsQuery <- dataQuery(xy=moveData, st=o.date, img=rsStk, rt=r.date, tb=c(30,30))
}
```

getEnv	<i>getEnv</i>
--------	---------------

Description

Interface to download ecologically relevant data.

Usage

```
getEnv(d.path = NULL, d.source = NULL, t.var = NULL, ref = NULL,
       p.raster = FALSE, p.res = NULL, pad = 10)
```

Arguments

<code>d.path</code>	Output data path for downloaded data.
<code>d.source</code>	Data source. One of "EarthEnv", "GFC", "GSW", "CCI" or "HSM".
<code>t.var</code>	Target variable.
<code>ref</code>	Projected spatial object from which an extent can be derived.
<code>p.raster</code>	Logical. Should the output be reprojected?
<code>p.res</code>	Pixel resolution (used if <code>p.raster</code> is TRUE).
<code>pad</code>	Number of cells used to pad the output raster when resampling.

Details

Downloads data from earthenv.org. To check which variables can be downloaded, run the function without specifying `t.var` and specifying `d.source`. This will return a data frame listing the existing variables for a given data source. Here, the user can refer to the column `"code"` to retrieve the keywords that can be passed to the function through `t.var`. The keywords recognized by `d.source` are:

- `"EarthEnv"` - EarthEnv project.
- `"GFC"` - Maryland University Global Forest Change.
- `"GSW"` - JRC Global Surface Water.
- `"CCI"` - ESA CCI Global land cover.
- `"HSM"` - JRC Human Settlement map.
- `"NEO"` - NASA Earth Observations.

If `t.var` contains `"DEM90"` from "EarthEv" or any variable from `"GFC"` or `"GSW"`, The function might require more time as it will mosaic the tiles with which the reference spatial object (`ref`) overlap with. In any circumstance, the output will be cropped to the extent of `ref` and, if prompted, use it as a reference to reproject the output.

Value

One or multiple raster objects.

References

<http://www.earthenv.org/> <https://earthenginepartners.appspot.com/science-2013-global-forest>
<https://global-surface-water.appspot.com/> <http://ghsl.jrc.ec.europa.eu/> <http://maps.elie.ucl.ac.be/CCI/viewer/> <https://neo.sci.gsfc.nasa.gov/>

See Also

[dataQuery proSat](#)

Examples

```
{
  # return list of variables
  ee.var <- getEnv(d.source="EarthEnv")
  gfc.var <- getEnv(d.source="GFC")
  gsw.var <- getEnv(d.source="GSW")
  cci.var <- getEnv(d.source="CCI")
  hsm.var <- getEnv(d.source="HSM")
  neo.var <- getEnv(d.source="NEO")
}
```

hotMove

hotMove

Description

Identifies hotspots of samples using a pixel based approach.

Usage

```
hotMove(xy = xy, pxr = pxr, shp = FALSE)
```

Arguments

<code>xy</code>	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
<code>pxr</code>	Grid resolution. Unit depends on <code>xy</code> projection.
<code>shp</code>	Logical. Should the function provide polygons? Default is FALSE.

Details

The function builds a matrix for a given resolution and a spatial extent derived from a set of samples (`xy`). First, these samples are converted into unique pixel coordinates. Then, the spatial connectivity of these pixels is evaluated using a 8-neighbor connected component labelling algorithm. The function returns a vector with the region ID per sample (*\$indices*). If `shp` is TRUE the function also returns a shapefile for each region defined by the convex hull of the samples within them (*\$polygons*).

Value

A list object.

See Also

[sampleMove hotMoveStats](#)

Examples

```
{  
  
  require(raster)  
  
  # reference data  
  sprj <- crs("+proj=longlat +ellps=WGS84 +no_defs")  
  moveData <- read.csv(system.file('extdata', 'latlon_example.csv', package="rsMove"))  
  moveData <- SpatialPointsDataFrame(moveData[,2:3], moveData, proj4string=sprj)  
  
  # extract regions  
  hm <- hotMove(xy=moveData, pxr=0.1, shp=TRUE)  
  
  # plot shapefile (color by region)  
  plot(hm$polygons, col=hm$indices)  
  
  # add new information to original shapefile  
  moveData@data$indices <- hm$indices  
  
}
```

hotMoveStats

hotMoveStats

Description

Provides statistics for the output of `hotMove`.

Usage

```
hotMoveStats(rid = NULL, o.time = NULL, tUnit = NULL, aid = NULL,  
             method = "deg")
```

Arguments

<code>rid</code>	List object as provided by <code>hotMove()</code> .
<code>o.time</code>	Object of class <i>Date</i> , <i>POSIXlt</i> or <i>POSIXct</i> .
<code>tUnit</code>	Time unit for stats. Default is <i>days</i> . See difftime for additional keywords.
<code>aid</code>	Optional. Unique identifiers.
<code>method</code>	Method used to estimate polygon area.

Details

This functions analysis the attributes of sample regions define by hotMove(). Alternatively, the user can keep *rid* as NULL. This case, all information will be assumed as part of one region. For each sample region, the function returns the amount of samples (*ms*). If a vector of unique identifiers is provided (*aid*) the number of unique identifiers observed within each region is also reported. If temporal information is provided (*time*) the function identifies unique temporal segment corresponding to periods of consecutive days with observations. For each segment, the function reports on the amount of segments (*nts*) as well as the minimum (*mnt*), maximum (*mxt*) and mean (*avt*) of the time segments and the total amount of time that they amount to (*tts*). For each region, the function will also report on the start and end of each temporal segment (*\$temporal.segments*) and will provide the sample indices for associated to each segment (*\$segment.indices*). If *rid* contains polygons for each region, the function also reports on the area of convex polygons. In this case, the user can use the *method* keyword to specify how the polygons should be handled. If the polygons are in lat-lon, method *deg* can be used to re-project each polygon to its corresponding UTZ zone before retrieving the area. This is the default, if the polygons are in a cartesian coordinate system use *m*.

Value

A data frame.

See Also

[hotMove](#)

Examples

```
{
require(raster)

# reference data
sprj <- CRS("+proj=longlat +ellps=WGS84 +no_defs")
moveData <- read.csv(system.file('extdata', 'latlon_example.csv', package="rsMove"))
moveData <- SpatialPointsDataFrame(moveData[,2:3], moveData, proj4string=sprj)

# extract regions
hm <- hotMove(xy=moveData, pxr=0.1, shp=TRUE)

# plot shapefile (color by region)
plot(hm$polygons, col=hm$indices)

# add new information to original shapefile
moveData@data <- cbind(moveData@data, hm$indices)

# derive statistics
hm.region.stats <- hotMoveStats(rid=hm, o.time=as.Date(moveData@data$timestamp))
hm.time.stats <- hotMoveStats(o.time=as.Date(moveData@data$timestamp))
}
```

imgInt	<i>imgInt</i>
--------	---------------

Description

Temporal linear interpolation of raster data.

Usage

```
imgInt(img = NULL, rd = rd, td = td, bs = NULL, xy = NULL,
       edata = NULL)
```

Arguments

img	Object of class <i>RasterStack</i> or <i>RasterBrick</i> .
rd	Raster dates. Object of class <i>Date</i> .
td	Target dates. Object of class <i>Date</i> .
bs	wo element vector with temporal search buffer (expressed in days).
xy	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
edata	Object of class <i>data frame</i> or <i>matrix</i> with remote sensing data.

Details

Performs a pixel-wise linear interpolation over a raster for a given set of dates (*td*). A temporal buffer (*bs*) is required to limit the search for reference data points (*rd*). *bs* is defined by a two element vector which limits the search of images in the past and future. If *xy* is provided the function only considers the pixels that overlap with these sample points. Otherwise, a *RasterBrick* is provided. However, if *edata* is provided, *xy* and *img* are ignored, and a data frame is provided.

Value

A *RasterBrick* or a *data frame*.

See Also

@seealso [dataQuery](#) [timeDir](#) [spaceDir](#) [moveSeg](#)

Examples

```
{
  require(raster)

  #' # read raster data
  file <- list.files(system.file('extdata', '', package="rsMove"), 'tc.*tif', full.names=TRUE)
  rsStk <- stack(file)
  rsStk <- stack(rsStk, rsStk, rsStk) # dummy files for the example
```

```

# read movement data
moveData <- read.csv(system.file('extdata', 'konstanz_20130805-20130811.csv', package="rsMove"))
moveData <- SpatialPointsDataFrame(moveData[1:10,1:2], moveData[1:10,], proj4string=crs(rsStk))

# raster dates
rd = seq.Date(as.Date("2012-01-01"), as.Date("2012-12-31"), 45)

# target dates
td = as.Date("2012-04-01")

# interpolate raster data to target dates
i.img <- imgInt(img=rsStk, rd=rd, td=td, bs=c(60,60), xy=moveData)
}

```

labelSample

labelSample

Description

Region labeling of samples based on their spatial connectivity.

Usage

```
labelSample(xy = xy, rad = rad, npt = NULL, npx = NULL, pxr = r)
```

Arguments

<code>xy</code>	Object of class <i>SpatialPoints</i> of <i>SpatialPointsDataFrame</i> .
<code>rad</code>	Minimum radius. Unit depends on the projection of the data.
<code>npt</code>	Minimum pixel count per pixel.
<code>npx</code>	Minimum number of pixels.
<code>pxr</code>	Pixel resolution as a valid raster layer.

Details

First, the samples are converted to pixel coordinates and then one of two occur: 1) if *npt* is set, the function removes pixels with a pixel count smaller than the one specified; 2) If *npx* is set, the connectivity between neighboring samples is evaluated and regions with a count small than the specified value are filtered. Only one option may be set at a time. Then, the remaining pixels are dilated and the samples are again labeled accounting for regions that are not connected but are near to each other. Regions within a given distance of each other (defined by *rad*) are aggregated. The grid used for this analysis is built from the spatial extent of *xy* and a given pixel resolution (*pxr*). If *pxr* is a raster, this will be used to define the dimensions of the grid. Doing so can be of use when the user has pre-select environmental predictors that will be used for modeling. Note that the finer the resolution the more independent regions are likely to be returned. The output is a vector with ID's assigning each sample to its region. Samples filtered by *npt* or *npx* will be returned as zeros.

Value

A vector.

See Also

[sampleMove](#) [hotMove](#)

Examples

```
{  
  
  require(raster)  
  
  # read raster data  
  r <- raster(system.file('extdata', 'tcb_1.tif', package="rsMove"))  
  
  # read movement data  
  moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))  
  moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(r))  
  
  # derive region labels  
  labels <- labelSample(xy=moveData, rad=90, npx=2, pxr=30)  
  
}
```

modelApply

modelApply

Description

Apply a model or an ensemble of models to raster data.

Usage

```
modelApply(model, img, fun = NULL)
```

Arguments

model	List object as provided by <i>moveModel()</i> .
img	Object of class <i>RasterLayer</i> , <i>RasterStack</i> or <i>RasterBrick</i> .
fun	Function that specifies how the model should be applied.

Details

The function uses the output of *moveModel()*. If this contains a list of models from multiple runs, the function creates a stack of predictions and summarizes it on a pixel-by-pixel basis using a weighted mean. The weights are defined by the average performance for *presence* and *background* samples in each iteration.

Value

A *Raster*.

See Also

[segRaster](#) [moveModel](#)

Examples

```
## Not run:

require(raster)

# read remote sensing data
file <- list.files(system.file('extdata', '', package="rsMove"), 'tc.*tif', full.names=TRUE)
rsStk <- stack(file)

# read movement data
moveData <- read.csv(system.file('extdata', 'konstanz_20130805-20130811.csv', package="rsMove"))
moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(rsStk))

# retrieve remote sensing data for samples
rsQuery <- dataQuery(xy=moveData, img=rsStk, remove.dup=TRUE)

# identify unique sample regions
label <- labelSample(xy=rsQuery, rad=90, npx=1, pxr=rsStk)

# select background samples
ind <- which(label>0) # selected samples
bSamples <- backSample(xy=moveData[ind,], rid=label[ind], img=rsStk, nb=4000, method='pca')

# derive model predictions
fun <- function(x,y) {train(x, y, method="rf", trControl=trainControl(method='oob'))}
p.model <- moveModel(p.data=rsQuery@data, a.data=bSamples@data, label=label, fun=fun, nruns=1)

# derive prediction from model ensemble
fun <- function(x,y) {predict(x, y, type='prob')[[1]]$`1`}
prob <- modelApply(p.model, rsStk, fun=fun)

# see output
plot(prob)

## End(Not run)
```

Description

Provides historical information on cloud cover.

Usage

```
moveCloud(xy = xy, o.time = o.time, d.path = NULL, b.size = NULL,  
          remove.file = TRUE, p.res = T)
```

Arguments

<code>xy</code>	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
<code>o.time</code>	Object of class <i>Date</i> .
<code>d.path</code>	Output data path for downloaded data.
<code>b.size</code>	Two element vector with temporal buffer size (expressed in days).
<code>remove.file</code>	Logical. Should the files be deleted after usage?
<code>p.res</code>	Should the output be plotted on screen? Default is TRUE.

Details

This function makes uses daily cloud fraction data from NASA's NEO service. For each observation date (*o.time*), the function downloads the correspondent image and extracts the percent of cloud cover for the samples acquired at the target date. If *d.path* is specified, the function will look within the provided directory for the required files. If so, they won't be downloaded. If *d.buffer* is specified, for each date, the function will consider images before and after within a temporal buffer. *d.buffer* requires two elements which specify the buffer size before and after the target date. These new images will be used to report on the closest time step with the lowest possible cloud cover. The final report provides information on:

- *day.cover*: cloud cover for the observation date
- *p.day.before*: date before the observation date with the lowest cloud cover
- *p.cover.before*: cloud cover for *p.day.before*
- *p.day.after*: date after the observation date with the lowest cloud cover
- *p.cover.after*: cloud cover for *p.day.after*

The output will also contain a two plots with information on the distance between the observation dates and the closest date with the lowest cloud cover. The plots show the amount of samples that are covered in each of the target dates for the best dates before (*\$plot.before*) and after (*\$plot.after*) the observation dates.

Value

A list.

References

<https://cneos.jpl.nasa.gov/>

See Also

[sMoveRes](#) [tMoveRes](#)

Examples

```
## Not run:

require(raster)

# read raster data
r <- raster(system.file('extdata', 'tcb_1.tif', package="rsMove"))

# read movement data
moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))
moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(r))

# test function for 5, 10 20 and 30 m
od <- as.Date(moveData@data$date)
c.cover <- moveCloud(xy=moveData, o.time=od, d.path=".", b.size=30)

## End(Not run)
```

moveModel

moveModel

Description

Spatially stratified predictive modeling.

Usage

```
moveModel(p.data = p.data, a.data = a.data, label = NULL, fun = fun,
          nruns = 1)
```

Arguments

p.data	Object of class <i>data.frame</i> with environmental variables for presence samples.
a.data	Object of class <i>data.frame</i> with environmental variables for background samples.
label	Region labels. If missing, "p.data" is assumed as one region.
fun	A function with the modeling algorithm to use.
nruns	Number of runs. Default is 1.

Details

For n iterations, where n is the number of unique sample regions, the function uses one sample region for validation while the remaining ones are used for training. The background samples are split randomly at each iteration. The final accuracy, provided as a F1-score for both presence and background samples, is derived from the total of true and false positives ($f1$). Additionally, for each run, the function returns a model ($model$) which is trained using all the samples. This output can be passed to `modelApply()`. By default, the function uses a Random Forest classifier. However, the a user specified function can be passed using *fun*.

Value

A list.

See Also

[sampleMove](#) [labelSample](#) [backSample](#) [modelApply](#) [train](#)

Examples

```
## Not run:

require(rgdal)
require(raster)
require(sp)

# read remote sensing data
file <- list.files(system.file('extdata', '', package="rsMove"), 'tc.*tif', full.names=TRUE)
rsStk <- stack(file)

# read movement data
moveData <- read.csv(system.file('extdata', 'konstanz_20130805-20130811.csv', package="rsMove"))
moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(rsStk))

# retrieve remote sensing data for samples
rsQuery <- dataQuery(xy=moveData, img=rsStk, remove.dup=TRUE)

# identify unique sample regions
label <- labelSample(xy=rsQuery, rad=3000, pxr=rsStk)

# select background samples
ind <- which(label>0) # selected samples
bSamples <- backSample(xy=moveData[ind,], rid=label[ind], img=rsStk, method='pca')

# derive model predictions
fun <- function(x,y) {train(x, y, method="rf", trControl=trainControl(method='oob'))}
out <- moveModel(p.data=rsQuery@data, a.data=bSamples@data, label=label, fun=fun, nruns=1)

## End(Not run)
```

 moveReduce

moveReduce

Description

Remote sensing based point segmentation.

Usage

```
moveReduce(xy = xy, ot = ot, img = img)
```

Arguments

<code>xy</code>	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
<code>ot</code>	Object of class <i>Date</i> , <i>POSIXlt</i> or <i>POSIXct</i> with <code>xy</code> observation dates.
<code>img</code>	Object of class <i>RasterLayer</i> , <i>RasterStack</i> or <i>RasterBrick</i> .

Details

SReduces a set of input samples (`xy`) based on their assignment to unique pixels within a reference raster (`img`). The function looks at consecutive points ordered by time (`ot`) and aggregates samples if they remain within the same pixel. If the same pixel is revisited on a later time, that observation is kept as a separate occurrence. For each temporal segment, the function returns mean x and y coordinates, the start and end timestamps, the mean timestamp and the elapsed time.

Value

A *list*.

See Also

[sampleMove](#) [moveSeg](#)

Examples

```
{
  require(raster)

  # read raster data
  r <- raster(system.file('extdata', 'tcb_1.tif', package="rsMove"))

  # read movement data
  moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))
  moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(r))

  # observation time
  o.time <- strptime(paste0(moveData@data$date, ' ', moveData@data$time), format="%Y/%m/%d %H:%M:%S")
}
```



```
# reduce amount of samples
move.reduce <- moveReduce(xy=moveData, ot=o.time, img=r)

}
```

moveSeg

moveSeg

Description

Remote sensing based point segmentation

Usage

```
moveSeg(xy = xy, edata = edata, type = "cont", ot = NULL,
        b.size = NULL, threshold = NULL, s.fun = NULL)
```

Arguments

<code>xy</code>	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
<code>edata</code>	Object of class <i>RasterLayer</i> or <i>data.frame</i> .
<code>type</code>	Raster data type. One of <i>cont</i> (continues) or <i>cat</i> (for categorical).
<code>ot</code>	Object of class <i>Date</i> , <i>POSIXlt</i> or <i>POSIXct</i> with <i>xy</i> observation dates.
<code>b.size</code>	Buffer size expressed in the map units.
<code>threshold</code>	Change threshold.
<code>s.fun</code>	Output summary function. Default is mean.

Details

Segmentation of a point shapefile based on the spatial variability of a raster dataset. When the *method* is set to *'cont'*, the raster data is assumed to be continuous. Then, the function determines the percentual change between each pair of two consecutive coordinate pairs. If this change is above a predefined *threshold*, a new pointer is added and the previous sequence of samples is labeled as a unique segment. If *method* is set as *'cat'*, the function assumes the raster data is categorical ignoring the *theshold* and *s.fun* keywords. In this case, a new segment is identified if the any change is observed between two consecutife points. The output consists of a list containing a *SpatialPointsDataFrame* (*\$points*) reporting on the segment ID (*sid*) associated to each sample and a data frame (*\$report*) with the amount of points in each region and the value returned by *s.fun*. If *ot* is provided, the function also provides the elapsed time within each segment. If *fun* is set by the user, the provided function will be used to summarize the raster values at each segment. Also, if *edata* is a *RasterStack* or a *RasterBrick*, *r.fun* is used to reduce the multi-layered object to a single layer. he user can either use a Principal Component Analysis (PCA) analysis by setting *r.fun* to *pca* or provide a function. If *pca* is selected, the first Principal Component (PC) *threshold* will be set automatically to the standard deviation of the first PC. This is the default. If *method* is *cat* the function will assume the data is categorical and wil define segments whenever a change occurs.

Value

A list.

See Also

[dataQuery imgInt](#)

Examples

```
{
  require(raster)

  # read raster data
  r <- raster(system.file('extdata', 'tcb_1.tif', package="rsMove"))

  # read movement data
  moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))
  moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(r))

  # observation time
  o.time <- strptime(paste0(moveData@data$date, ' ', moveData@data$time), format="%Y/%m/%d %H:%M:%S")

  # perform directional sampling
  seg <- moveSeg(xy=moveData, ot=o.time, edata=r, type="cont", threshold=0.1)
}
```

plotMove

plotMove

Description

Plots per pixel time and value.

Usage

```
plotMove(x = x, y = y, o.time = NULL, value = NULL, type = NULL)
```

Arguments

x	Vector of x coordinates.
y	Vector of y coordinates.
o.time	Vector with time length.
value	Vector with environmental data.edata Object of class <i>RasterLayer</i> or <i>data.frame</i> .
type	One of 'cont' or 'cat'. Defines the type of <i>value</i> .

Details

This function plots a provided set of x and y coordinates adding information on the elapsed time at each coordinate (*e.time*) and/or a given environmental variable (*value*). If only *time* or *value* are set, the function builds a scatterplot where the size of the points is defined by the input variables. If both are provided, the size of the points is defined by the elapsed time and the raster value is used to build a color scheme for the points. When *value* is provided, the keyword *type* is required. It will influence how the plots are built. The breaks a limits for the point size and colors is define automatically.

Value

A *ggplot* object.

See Also

[dataQuery](#) [moveReduce](#)

Examples

```
{  
  
  require(raster)  
  
  # read raster data  
  r <- raster(system.file('extdata', 'tcb_1.tif', package="rsMove"))  
  
  # read movement data  
  moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))  
  moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(r))  
  
  # observation time  
  o.time <- strptime(paste0(moveData@data$date, ' ', moveData@data$time), format="%Y/%m/%d %H:%M:%S")  
  
  # reduce amount of samples  
  move.reduce <- moveReduce(xy=moveData, ot=o.time, img=r)  
  
  # query data  
  ov <- dataQuery(xy=move.reduce$points, img=r)  
  
  # plot output  
  x <- move.reduce$points@coords[,1]  
  y <- move.reduce$points@coords[,2]  
  et <- move.reduce$points@data$elapsed.time  
  op <- plotMove(x=x, y=y, o.time=et, value=ov[,1], type="cont")  
  
}
```

poly2sample

poly2sample

Description

Convert spatial polygons into point samples.

Usage

```
poly2sample(pol = pol, re = NULL, mpc = NULL, pr = NULL)
```

Arguments

<code>pol</code>	Object of class <i>SpatialPolygons</i> or <i>SpatialPolygonDataFrame</i> .
<code>re</code>	Object of class <i>Extent</i> or a raster object from which an extent can be derived.
<code>mpc</code>	Minimum pixel cover (0-100). Default is 100.
<code>pr</code>	Pixel resolution.

Details

Determines coordinates of pixels within a given extent. If *re* is missing the target extent corresponds to the extent of the polygon layer. In this case, *pr* is required. If *re* is an *Extent* object or *re* is missing *rp* is required. *mpc* can be used to filter pixels with low purity, i.e. pixels where the percentage of area cover by a polygon is below the defined threshold. The output provides the coordinates (*x* and *y*) the pixel percent cover (*cover*) for each selected pixel.

Value

A *SpatialPointsDataFrame*.

See Also

[dataQuery](#) [imgInt](#)

Examples

```
{  
  
  require(raster)  
  
  # load example probability image  
  file <- system.file('extdata', 'konstanz_probabilities.tif', package="rsMove")  
  img <- raster(file)  
  
  # load area of interest  
  file <- system.file('extdata', 'konstanz_roi.shp', package="rsMove")  
  roi <- shapefile(file)
```

```

# segment probabilities
samples <- poly2sample(pol=roi, re=img)

}

```

proSat

proSat

Description

Interface to download and process satellite data.

Usage

```

proSat(t.var = NULL, xy = NULL, o.time = NULL, d.path = NULL,
       p.raster = FALSE, p.res = NULL, user.cred = NULL, pad = 10)

```

Arguments

<code>t.var</code>	Target variable.
<code>xy</code>	Object from which an extent can be derived.
<code>o.time</code>	Object of class <i>Date</i> .
<code>d.path</code>	Output data path for downloaded data.
<code>p.raster</code>	Logical. Should the output be re-projected?
<code>p.res</code>	Target pixel resolution (if <code>p.raster</code> is TRUE).
<code>user.cred</code>	Two element character vector containing username and password.
<code>pad</code>	Padding expressed in number of pixels. Used when re-projecting and resampling.

Details

Downloads and pre-processes pre-selected satellite datasets That provide ecologically meaningful variables. `xy` is required to define a reference extent for which the data will be downloaded and cropped. If `p.raster` is TRUE, the function will also re-project the output to the projection of `xy`. Note that `xy` does not required reprojection a priori. The function will find the appropriate projection to use when deriving a reference extent. If `p.raster` is TRUE `p.res` is also required to determine the output pixel resolution. The outputs are presented in GTiff format and are cropped and masked by default. Note that the download files might not correspond to the dates supplied through `o.time`. The function will consider the temporal resolution of the target dataset and compare the possible and the requested dates. If a requested date is not found, the function will instead provide the closest image time. As a consequence, the number of downloaded files might be lesser than the number of dates. The function will inform the user on this by providing a list object which contains the downloaded dates (*\$date*) as well as the path for the correspondent image (*\$path*).

Value

One or multiple raster objects.

Note

To consult the list of provided variables, run the function without specifying any input. This will provide information on the variables, variable codes (used in *t.var*), temporal and spatial resolution and the sensor of origin. Some variables might require login credentials. Check the table to know which credentials to use and assign them through *user.cred*.

See Also

[getEnv imgInt dataQuery](#)

Examples

```
{
  # return list of variables
  modis.var <- proSat()
}
```

rsComposite

rsComposite

Description

Compositing of remote sensing data based on GPS tracking dates.

Usage

```
rsComposite(img, rd, ot, cm = 1, type = "norm", d.buffer = NULL)
```

Arguments

img	Object of class <i>RasterSpack</i> or <i>RasterBrick</i> .
rd	Object of class <i>Date</i> with <i>img</i> observation dates.
ot	Object of class <i>Date</i> with reference dates.
cm	Number of deviations from the target date. Default is 1.
type	One of "norm" or "pheno" or "pheno2".
d.buffer	Search buffer (expressed in days).

Details

The function uses a multi-layer raster object to build a composite. It looks at a given set of dates (e.g. GPS tracking dates) and estimates a reference date to build the composite for defined by the median of *ot*. The median is then used to estimate Median Absolute Deviation (MAD) which specifies the size of the buffer set around the target date within which bands will be considered. Here, *cm* is used as a multiplier to enlarge the temporal buffer. Alternatively, a user define temporal buffer is allowed by using the keyword *d.buffer*. If *ot* contains only one element, the function will use it as a reference date. In this case, if *d.buffer* is NULL the function will set it to 30 by default. The way how the function handles temporal information depends on the *type* keyword. If set to *norm*, the function will search for the nearest possible dates within the temporal buffer. However, if *pheno* is set, then the day of the year will be given priority. Thus, if multi-year raster data is provided, older data with a DOY closer to the target that will be used when possible. The output provides: #'

- *value* - composite of target images
- *dates* - per pixel date code
- *count* - pixel count of *dates*
- *na.count* - count of NA values
- *target* - target date
- *mad* - temporal buffer

If *pheno2* is used, for each pixel, the function will estimate a weighted mean of the clear pixels within the temporal buffer. The weights represent the inverse time difference between the target and the available dates give higher weights to small differences.

Value

A list.

See Also

[imgInt dataQuery proSat](#)

Examples

```
## Not run:

require(raster)

# read raster data
file <- list.files(system.file('extdata', '', package="rsMove"), 'tc.*tif', full.names=TRUE)
rsStk <- stack(file)
rsStk <- stack(rsStk, rsStk, rsStk) # dummy files for the example

# raster dates
rd = seq.Date(as.Date("2013-01-01"), as.Date("2013-12-31"), 45)

# target date
ot = as.Date("2013-06-01")
```

```
# build composite
r.comp <- rsComposite(rsStk, rd, ot, d.buffer=90)

## End(Not run)
```

sampleMove

sampleMove

Description

Sampling of possible stops along a movement track.

Usage

```
sampleMove(xy = xy, ot = ot, error = error, method = "m",
           tUnit = NULL)
```

Arguments

xy	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
ot	Object of class <i>Date</i> , <i>POSIXlt</i> or <i>POSIXct</i> with the same length as <i>xy</i> .
error	Distance (in meters).
method	How should the distance be estimated? One of 'm' or 'deg'. Default is 'm'.
tUnit	Time unit to estimate elapsed time. See difftime for keywords. Default is <i>mins</i> .

Details

This function offers a simple approach to sample from locati where an animal showed little or no movement based on GPS tracking data. It looks at the distance among consecutive samples (*error*) and estimates mean coordinates for the temporal segments where the animal moved less than the defined distance from the first location of the segment. The user should selected *method* in accordance with the projection system associated to the data. If 'm' it estimates the euclidian distance. If 'deg' it uses the haversine formula. The output reports on the mean sample coordinates for the sample locations ('x' and 'y'), the start, end and total time spent per sample ('time' expressed in minutes) and the total number of observations per sample ('count').

Value

A *SpatialPointsDataFrame*.

See Also

[labelSample](#) [backSample](#) [dataQuery](#)

Examples

```

{

  require(raster)

  # reference data
  sprj <- crs("+proj=longlat +ellps=WGS84 +no_defs")
  moveData <- read.csv(system.file('extdata', 'latlon_example.csv', package="rsMove"))
  moveData <- SpatialPointsDataFrame(moveData[,2:3], moveData, proj4string=sprj)

  # sampling without reference grid
  ot = strptime(moveData$timestamp, "%Y-%m-%d %H:%M:%S")
  output <- sampleMove(xy=moveData, ot=ot, error=7, method='deg')

  # compare original vs new samples
  plot(moveData, col="black", pch=16)
  points(output$x, output$y, col="red", pch=15)

}

```

 satTime

satTime

Description

Finds available satellite dates for tracking data.

Usage

```
satTime(o.time = o.time, t.var = t.var)
```

Arguments

<code>o.time</code>	Object of class <i>Date</i> .
<code>t.var</code>	Target variable.

Details

This function compares a set of input dates (*o.time*) to the possible dates for which satellite data can be downloaded. This analysis is performed for the set of variables provided through the function `proSat()`. The function provides a report with the closest dates available for each of the samples and a faceted plot for each unique year showing the observed dates, the dates covered by the satellite data and the dates covered by both.

Value

One or multiple plots.

See Also

[moveCloud proSat](#)

Examples

```
{
  # return list of variables
  var.ls <- satTime(o.time=as.Date("2013-08-04"), t.var="ndvi")
}
```

segRaster

segRaster

Description

Connencte-region based raster segmentation.

Usage

```
segRaster(prob, pt = 0.1, mp = 0.5)
```

Arguments

prob	Object of class <i>RasterLayer</i> .
pt	Difference threshold. Default is 0.05.
mp	Minimum value. Default is 0.5.

Details

The function segments an input layer using a connected component region labeling approach. For each pixel, the function estimates the difference between it and its imediate 8 neighbors. The pixels where the difference is below the defined threshold (*ct*) are aggregated into a single region. The user can define a minimum pixel value using *mp* which will limit the range of pixels under evaluation. The result contains a raster with unique values for each segment region region (*\$segment*) as well as a data frame (*\$stats*) with statistics for each region. The data frame report on the minimum (*min*), maximum (*max*), mean (*mean*) and standard deviation (*sd*) of the pixels contained in each region.

Value

A list object.

See Also

[moveModel modelApply](#)

Examples

```
{  
  
  require(raster)  
  
  # load example probability image  
  file <- system.file('extdata', 'konstanz_probabilities.tif', package="rsMove")  
  probImg <- raster(file)  
  
  # segment probabilities  
  rs <- segRaster(probImg)  
  
}
```

sMoveRes

sMoveRes

Description

Evaluates how a change in raster resolution impactst the availability of data points.

Usage

```
sMoveRes(xy = xy, pxr = pxr)
```

Arguments

<code>xy</code>	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
<code>pxr</code>	vector of target resolutions.

Details

Given a vector of pixel resolutions (*pxr*), the function determines the number of unique pixels and unique pixel groups. Additionally, for each pixel, the function returns the corresponding pixel indices per resolution showing which samples would be grouped. The function returns a data frame (*\$stats*) and a plot (*\$plot*) with the statistics per resolution as well as a data frame with the pixel indices per resolution (*\$indices*).

Value

A *list*.

See Also

[tMoveRes specVar](#)

Examples

```

{

  require(raster)

  # read movement data
  moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))
  moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData)

  # test function for 5, 10 20 and 30 m
  a.res <- sMoveRes(xy=moveData, pxr=c(5, 10, 20, 30))

}

```

spaceDir

spaceDir

Description

Spatial directional raster analysis.

Usage

```
spaceDir(xy = xy, ot = NULL, img = img, dir = dir, type = type,
         dm = "m", b.size = NULL, fun = NULL, npx = 2)
```

Arguments

xy	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
ot	Object of class <i>Date</i> , <i>POSIXlt</i> or <i>POSIXct</i> with <i>xy</i> observation dates.
img	Object of class <i>RasterLayer</i> .
dir	One of <i>fwd</i> , <i>bwd</i> or <i>both</i> . Default is <i>both</i> .
type	One of 'cont' or 'cat'. Defines which type of variable is in use.
dm	One of 'm' or 'deg' specifying the projection unit. Default is 'm'.
b.size	Buffer size expressed in the map units.
fun	Summary function.
npx	Minimum number of pixels used in <i>fun</i> . Default is 2.

Details

This function evaluates how do environmental conditions change in space along a movement track. First, it looks at consecutive observations to define spatial segments. All the pixels between the two endpoints of the segment are considered in this process. The user can use the argument *dir* to prompt the function to focus on previous (*bwd*) or following (*fwd*) observations or to look in both directions (*both*) when defining the segments. The output consists of a list containing two

shapefiles, one *SpatialPointsDataFrame* (*\$endpoints*) with the endpoints of each segment and a *SpatialLinesDataFrame* (*\$segments*) representing each segment. The data frames provided with these shapefiles reports on:

- *x* - mean x coordinates
- *y* - mean y coordinates
- *timestamp* - mean observation time
- *pixel.time* - elapsed time within a pixel for a given segment
- *travel.distance* - distance between a segment endpoints
- *travel.time* - elapsed time between a segment endpoints

The additional information added to the data frame depends on the type of data. If *type* is set to *'cont'*, the function will assume the input raster is a continuous variable and estimate a statistical metric for each segment. If none is provided through the keyword *fun*, the slope will be returned by default and assigned to the column *'stat'* within the output data frame. If *type* is set to *'cat'*, the function will assume the data is categorical. As a result, the keyword *fun* is ignored. Instead, the function will return the proportion of pixels occupied by each class for each segment. In addition, the dominant class (*'main'*) and the shannon index (*'shannon'*) will be returned for each segment within the output data frame. In order to include the area surrounding the sample points in this analysis, *d.buffer* can be used. This keyword dilates the interpolated sample points by a given distance transforming the initial line into a polygon. @note If *d.buffer* is used, the default *fun* (the slope) is not adequate given that spatial order of the samples is not preserved. This should be considered when using these keywords simultaneously,

Value

A list.

See Also

[timeDir](#) [dataQuery](#) [imgInt](#)

Examples

```
{
  require(raster)

  # read raster data
  r <- raster(system.file('extdata', 'tcb_1.tif', package="rsMove"))

  # read movement data
  moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))
  moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(r))

  # observation time
  ot <- strptime(paste0(moveData@data$date, ' ', moveData@data$time), format="%Y/%m/%d %H:%M:%S")

  # perform directional sampling
  of <- function(x) {lm(x~c(1:length(x)))$coefficients[2]}
```

```
s.sample <- spaceDir(xy=moveData, ot=ot, img=r, dir="bwd", type='cont', fun=of)
}
```

specVar *specVar*

Description

Quantifies how changes in the resolution of a raster affects the perception of spectral complexity.

Usage

```
specVar(img = img, xy = NULL, pxr = pxr)
```

Arguments

<code>img</code>	Object of class <i>RasterLayer</i> .
<code>xy</code>	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
<code>pxr</code>	vector of target resolutions.

Details

Given a raster object, the function determines how reducing the resolution of the raster impacts our ability to perceive the complexity of the landscape. The function aggregates *img* to each of the given pixel resolutions (*pxr*) and estimates the Mean Absolute Error (MAE) for each pixel with the aggregated layer where the differences are estimated from the pixel within the original raster that overlap with the target pixel. If a point shapefile is provided (*xy*), the function will only report on the values that overlap with the points. The output of the function consists of:

- *mae* - MAE, either for all pixels or for the points within *xy*
- *pixel.optimal* - raster reporting on the resolution with the lowest MAE for each pixel
- *pixel.optimal.stats* - Proportion of samples per resolution derived from *pixel.optimal*
- *plot* - boxplots of the variability of the MAE per resolution

If *xy* is set, the output will contain a vector with the optimal resolution per sample (*\$sample.optimal*).

Value

A *list*.

See Also

[tMoveRes](#) [sMoveRes](#)

Examples

```
## Not run:

require(raster)

# read raster data
r <- raster(system.file('extdata', 'tcb_1.tif', package="rsMove"))

# read movement data
moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))
moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(r))

# apply function
s.var <- specVar(img=r, xy=moveData, pxr=60)

## End(Not run)
```

timeDir

timeDir

Description

Temporal directional raster analysis.

Usage

```
timeDir(xy = NULL, ot = ot, img = NULL, edata = NULL, rt = rt,
        mws = NULL, dir = NULL, fun = NULL)
```

Arguments

xy	Object of class "SpatialPoints" or "SpatialPointsDataFrame".
ot	Object of class <i>Date</i> , <i>POSIXlt</i> or <i>POSIXct</i> with <i>xy</i> observation dates.
img	Object of class <i>RasterStack</i> or <i>RasterBrick</i> .
edata	Object of class <i>data frame</i> .
rt	Object of class <i>Date</i> , <i>POSIXlt</i> or <i>POSIXct</i> with <i>img</i> observation dates.
mws	Moving window size (expressed in days).
dir	One of <i>fwd</i> , <i>bwd</i> or <i>both</i> . Default is <i>both</i> .
fun	Summary function.

Details

This function evaluates how do environmental conditions change in time along a movement track. First, it compares the observation times (*ot*) of *xy* against the acquisition times (*rt*) of *img* to search for relevant information within a pre-defined temporal window (*mws*). The user can chose to only consider time steps before (*bwd*) or after (*fwd*) the target observation time or look at both directions (*both*). If the latest is chosen, the function applies *mws* equally to both directions. After selecting adequate temporal information, a statistical metric (*fun*) is used to summarize the selected time steps. By default, the slope will be used. The slope is estimated from a linear regression estimated between the acquisition times of *img* and their corresponding values. When providing a new function, set *x* for time and *y* for the raster values. The output reports on:

- *x* - mean x coordinates
- *y* - mean y coordinates
- *timestamp* - mean observation time
- *pixel.time* - elapsed time within a pixel for a given segment
- *stat*: statistical metric

If *edata* is provided, *img* will be ignores as *edata* will contain the environmental data with each column representing a different variable. Otherwise, this data will be retrieved from *img*. Also, if *edata* is provided, *xy* is not required. However, it can be provided to built a spatial plot with the results of the analysis.

Value

A vector.

See Also

[spaceDir](#) [dataQuery](#) [imgInt](#)

Examples

```
{
  require(raster)

  # read raster data
  file <- list.files(system.file('extdata', '', package="rsMove"), 'tc.*tif', full.names=TRUE)
  rsStk <- stack(file)
  rsStk <- stack(rsStk, rsStk, rsStk) # dummy files for the example

  # read movement data
  moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))
  moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(rsStk))

  # raster dates
  rd <- seq.Date(as.Date("2013-08-01"), as.Date("2013-08-09"), 1)

  # sample dates
  ot <- strptime(paste0(moveData@data$date, ' ', moveData@data$time), format="%Y/%m/%d %H:%M:%S")
}
```



```
# perform directional sampling
of <- function(x,y) {lm(y~x)$coefficients[2]}
time.env <- timeDir(xy=moveData, ot=ot, img=rsStk, rt=rd, mws=10, dir="bwd", fun=of)
}
```

tMoveRes

tMoveRes

Description

Provides historical information on cloud cover.

Usage

```
tMoveRes(xy = xy, o.time = o.time, t.res = t.res, s.res = s.res)
```

Arguments

<code>xy</code>	Object of class <i>SpatialPoints</i> or <i>SpatialPointsDataFrame</i> .
<code>o.time</code>	Object of class <i>Date</i> , <i>POSIXlt</i> or <i>POSIXct</i> with <i>xy</i> observation dates.
<code>t.res</code>	Temporal resolution.
<code>s.res</code>	Spatial resolution.

Details

Given a vector of temporal resolutions (*t.res*), the function determines the number of unique pixels and unique pixel groups after their temporal aggregation. The function returns the corresponding pixel indices per resolution showing which samples would be grouped (*\$indices*). The function returns a data frame (*\$stats*) and a plot (*\$plot*) with the statistics per temporal resolution.

Value

A *list*.

See Also

[sMoveRes specVar](#)

Examples

```
## Not run:

require(raster)

# read raster data
r <- raster(system.file('extdata', 'tcb_1.tif', package="rsMove"))

# read movement data
moveData <- read.csv(system.file('extdata', 'konstanz_20130804.csv', package="rsMove"))
moveData <- SpatialPointsDataFrame(moveData[,1:2], moveData, proj4string=crs(r))

# test function for 5, 10 20 and 30 m
a.res <- tMoveRes(xy=moveData, dpath='.')

## End(Not run)
```

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