Package ‘trackdem’

February 27, 2020

Type Package
Title Particle Tracking and Demography
Version 0.5.2
Date 2020-02-27
Author Marjolein Bruijning, Marco D. Visser, Caspar A. Hallmann, Eelke Jongejans
Maintainer Marjolein Bruijning <mbruijning@princeton.edu>
Description Obtain population density and body size structure, using video material or image sequences as input. Functions assist in the creation of image sequences from videos, background detection and subtraction, particle identification and tracking. An artificial neural network can be trained for noise filtering. The goal is to supply accurate estimates of population size, structure and/or individual behavior, for use in evolutionary and ecological studies.
License GPL-2
URL https://github.com/marjoleinbruijning/trackdem
BugReports https://github.com/marjoleinbruijning/trackdem/issues
Encoding UTF-8
Depends
Imports png, neuralnet, raster, Rcpp, MASS, grDevices, graphics, stats, shiny
LinkingTo Rcpp, RcppArmadillo,
RoxygenNote 6.1.1
SystemRequirements Python (>=2.7), Libav, ExifTool
Suggests knitr, rmarkdown, testthat
VignetteBuilder knitr
NeedsCompilation yes
Repository CRAN
Date/Publication 2020-02-27 22:10:02 UTC
createBackground

Description
createBackground detects the still background, containing all motionless pixels (non particles). Three different methods to detect the background can be used.

Usage
createBackground(colorimages, method = "mean")

Arguments
colorimages Array of class 'TrDm' containing all images, obtained by loadImages.
method Use method='mean' to calculate the mean value for each pixel and color. Use method='powerroot' to deflate dark values (note, this can only be used for dark particles on a light background). Use method='filter' to replace pixels in which movement has occurred with the mode of neighboring values. Note that method='filter' is computationally more intensive.

Value
Array of class 'TrDm' and 'colorimage' containing detected background.
Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Examples

```r
## Not run:
dir.create("images")
## Create image sequence
traj <- simulTrajec(path="images",
                   nframes=30, nIndividuals=20, domain="square",
                   h=0.01, rho=0.9,
                   sizes=runif(20, 0.004, 0.006))
## Load images
dir <- "images"
allFullImages <- loadImages (dirPictures=dir, nImages=1:30)
stillBack <- createBackground(allFullImages, method="mean")
plot(stillBack)
## End(Not run)
```

`createImageSeq` creates an image sequence (.png) using video files as input. All movies within a directory will be converted into an image sequence. For each movie, a new directory is created containing the recorded date and name of the movie.

Usage

```r
createImageSeq(moviepath = "Movies", imagepath = "ImageSequences",
               x = 1920, y = 1080, fps = 15, nsec = 2, start = NULL,
               stop = NULL, ext = "MTS", libavpath = "avconv",
               exiftoolpath = "exiftool", pythonpath = "python", verbose = FALSE,
               logfile = FALSE)
```

Arguments

- **moviepath**: Path to existing directory containing the video files. By default, 'Movies' is used.
- **imagepath**: Path to location of a directory in which image sequences should be saved. By default, 'ImageSequences' is used (and created if not existing).
- **x**: Number of pixels in horizontal direction; default is 1920 (HD).
- **y**: Number of pixels in vertical direction; default is 1080 (HD).
- **fps**: Frames per second, default is 15.
findThreshold

**findThreshold**

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>nsec</td>
<td>Duration of movie that is exported, default is 2 seconds. When movie length is greater than nsec, the nsec seconds in the exact middle of the movie are exported.</td>
</tr>
<tr>
<td>start</td>
<td>Start time (in seconds) from where the video is converted (optional). By default, the nsec middle second of the video are used. If a a start time is specified and no stop time, nsec seconds starting from start are converted.</td>
</tr>
<tr>
<td>stop</td>
<td>End time (in seconds) from where the video is converted (optional). By default, the nsec middle second of the video are used. When an end time but no start time are specified, conversion starts at nsec seconds before end.</td>
</tr>
<tr>
<td>ext</td>
<td>The extension of the video. Default is 'MTS'. All formats supported by libav are accepted. To convert videos with different extensions, use for example c('MTS','mp4').</td>
</tr>
<tr>
<td>libavpath</td>
<td>Path to location where the executable file for libav can be found (named 'avconv.exe'), in case it is not found automatically, e.g. 'C:/Users/libav/usr/bin/avconv.exe'.</td>
</tr>
<tr>
<td>exiftoolpath</td>
<td>Path to location where the executable file for ExifTool can be found, in case it is not found automatically. For instance, use 'exiftool(-k).exe', if located in the working directory.</td>
</tr>
<tr>
<td>pythonpath</td>
<td>Path to location where the executable file for Python 2.7 can be found, in case it is not found automatically. For instance, use 'C:/Python27/python.exe'.</td>
</tr>
<tr>
<td>verbose</td>
<td>Logical. By default FALSE. Set to TRUE will print additional information.</td>
</tr>
<tr>
<td>logfile</td>
<td>Logical. By default FALSE. Set to TRUE will create a log file in the working directory.</td>
</tr>
</tbody>
</table>

**Author(s)**

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

**Examples**

```r
## Not run:
createImageSeq(moviepath="Movies",imagepath="ImageSequences", nsec=3,ext="AVI")
## End(Not run)
```

---

**Description**

This function can help to find a threshold value to distinguish noise from particles of interest.

**Usage**

```r
findThreshold(images, frame = 1, colorimages = NULL)
```
identifyParticles

Arguments

images Array containing images containing all moving particles, as obtained from `subtractBackground`.
frame Number specifying which frame to use. Default is frame 1.
colorimages Array containing original color images. By default, the original color images are obtained from the global environment.

Value

Returns the number that is interactively chosen by the user. Use this threshold value in `identifyParticles`.

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Examples

```r
## Not run:
dir.create("images")
dir <- dir.create("images")
traj <- simulTrajec(path="images",
                   nframes=30,nIndividuals=20,domain="square",
                   h=0.01,rho=0.9,
                   sizes=runif(20,0.004,0.006))
dir <- "images"
allFullImages <- loadImages(dir,dirPictures=dir,nImages=1:30)
stillBack <- createBackground(allFullImages,method="mean")
allImages <- subtractBackground(stillBack)
thr <- findThreshold(allImages,frame=10)
## End(Not run)
```

identifyParticles Identify moving particles

Description

`identifyParticles` identifies moving particles using the subtracted images obtained from `subtractBackground`. Function uses Connected Component Labeling and obtains particle statistics based on code developed for the orphaned package SDMTools (written by Jeremy VanDerWal).

Usage

```r
identifyParticles(sbg, threshold = -0.1, pixelRange = NULL,
                  qthreshold = NULL, select = "dark", colorimages = NULL,
                  autoThres = FALSE, perFrame = FALSE, frames = NULL)
```
identifyParticles

Arguments

**sbg** Array containing images containing all moving particles, as obtained from `subtractBackground`.

**threshold** Thresholds for including particles. A numeric vector containing three values; one for each color. Otherwise, supply one value which is to be used for all three colors. For a chosen quantile for each frame, use `qthreshold`. Default is `threshold=-0.1`, which works for dark particles on a light background. Alternatively, set `autoThres` below for an automatic threshold.

**pixelRange** Default is `NULL`. Numeric vector with minimum and maximum particle size (area), used as a first filter to identify particles. Use if particle of interest are of a known size range (in pixels).

**qthreshold** Default is `NULL`. Supply a value, to do thresholding based on quantile. Quantile is calculated for each frame separately.

**select** Select dark particles (`'dark'`), light particles (`'light'`), or both (`'both'`), compared to background. Default is `'dark'`.

**colorimages** Array containing original color images. By default, the original color images are obtained from global environment.

**autoThres** Logical. `TRUE` to get an automated threshold for each color layer. Default is `FALSE`.

**perFrame** Logical. If `autoThres=TRUE`, set at `TRUE` to calculate a threshold for each frame separately. Default is `FALSE`. Note that is can be computationally intensive to calculate a threshold for each frame.

**frames** When `autoThres=TRUE` and `allFrames=FALSE`, supply a numeric vector specifying over which frames the automated threshold should be calculated on (e.g. `c(1,3,5,7,9,11)` for all odd frames from 1 to 11).

Value

Returns a dataframe of class `'TrDm'` and `'particles'`, containing particle statistics with identified particles for each frame.

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Examples

```r
## Not run:
dir.create("images")
## Create image sequence
traj <- simulTrajec(path="images",
nframes=30,nIndividuals=20,domain="square",
h=0.01,rho=0.9,
sizes=runif(20,0.004,0.006))
## Load images
dir <- "images"
allFullImages <- loadImages (dirPictures=dir,nImages=1:30)
stillBack <- createBackground(allFullImages,method="mean")
```
allImages <- subtractBackground(stillBack)
partIden <- identifyParticles(allImages, threshold=-0.1,
       pixelRange=c(3,400))
plot(partIden)
summary(partIden)

## End(Not run)

loadImages

Description

loadImages loads png images as three dimensional arrays. The objects created through the function can be used for image analysis.

Usage

loadImages(dirPictures, filenames = NULL, nImages = 1:30,
          xranges = NULL, yranges = NULL)

Arguments

dirPictures  The path of the folder where the images can be found.
filenames    Default is NULL, or all files. If all files should NOT be loaded, here specify which files to use, as a character string.
nImages      Numeric vector specifying which images in the directory should be loaded; default is 1:30.
xranges      By default the full image is loaded; specify to subset the number of columns.
yranges      By default the full image is loaded; specify to subset the number of rows.

Value

Array of class 'TrDm' and 'colorimages' containing all loaded images.

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Examples

## Not run:
dir.create("images")
## Create image sequence
traj <- simulTrajec(path="images",
    nframes=30,nIndividuals=20,domain="square",
    h=0.01,rho=0.9,
    sizes=runif(20,0.004,0.006))
## Load images

dir <- "images"
allFullImages <- loadImages (dirPictures=dir,nImages=1:30)
plot(allFullImages)

## End(Not run)

---

**manuallySelect**

*Manually identify true and false positives with a GUI.*

**Description**

`manuallySelect` opens a graphic user interface to create training data for a neural net by manually selecting true and false positives (i.e. correctly identified particles and noise, respectively).

**Usage**

```r
manuallySelect(particles, colorimages = NULL, frames = NULL)
```

**Arguments**

- `particles` A data frame of class 'TrDm' with particle statistics for each frame, obtained by `identifyParticles`.
- `colorimages` An array with the original full color images, in order to plot on the original images. If `NULL`, the original color images are used, obtained from the global environment.
- `frames` A vector defining the frame(s) that should be used. Default is `NULL`; in that case the frame with the maximum number of identified particles is used.

**Value**

List containing three elements: true positives, false positives, and the evaluated frame.

**Author(s)**

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

**Examples**

```r
## Not run:
dir.create("images")
## Create image sequence
traj <- simulTrajec(path="images",
                     nframes=30,nIndividuals=20,domain="square",
                     h=0.01,rho=0.9,movingNoise=TRUE,
                     parsMoving = list(density=20, duration=10, size=1,
                                       speed = 10, colRange = c(0,1)),
                     sizes=runif(20,0.004,0.006))
```
mergeTracks

## Load images

dir <- "images"
allFullImages <- loadImages (dirPictures=dir,nImages=1:30)
stillBack <- createBackground(allFullImages,method="mean")
allImages <- subtractBackground(stillBack)
partIden <- identifyParticles(allImages,threshold=-0.1,
    pixelRange=c(3,400))

# select the nframes with the most identified particles
nframes <- 3
frames <- order(tapply(partIden$patchID,partIden$frame,length),
    decreasing=TRUE)[1:nframes]
mId <- manuallySelect(particles=partIden,frame=frames)

## End(Not run)

mergeTracks

Merge track records

Description

mergeTracks attempts to merge to two track objects as obtained by trackParticles.

Usage

mergeTracks(records1, records2, L = NULL, weight = NULL,
    logsizes = FALSE)

Arguments

records1 Object of class 'tracked', obtained using trackParticles.
records2 Object of class 'tracked', obtained using trackParticles that should be linked
to records1.
L Numeric. Maximum cost for linking a particle to another particle. When the
cost is larger, particles will be not be linked (resulting in the begin or end of a
segment). If NULL, the same L as used to create records2 is used.
weight Vector containing 3 weights to calculate costs. Depending on the study system
user may want to value certain elements over others. Weights are ordered as fol-
lows; first number gives the weight for differences in x and y coordinates; second
number gives the weight for particle size differences. Note that the difference
between the predicted location and the observed location is not taken into ac-
count in this function. If NULL, the same weights as used to create records2 is
used.
logsizes Logical. Default is FALSE. Set to TRUE to take the natural logarithm of body
sizes, when calculating the cost of linking two particles.

Value

A list of class 'TrDm' and 'records'. Use 'summary' and 'plot'.
Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Examples

```r
## Not run:
## Create image sequence
dir.create("images")
traj <- simulTrajec(path="images",
  nframes=60,nIndividuals=20,domain="square",
  h=0.01,rho=0.9,sizes=runif(20,0.004,0.006))
## Analyse first part
dir <- "images"
allFullImages1 <- loadImages (dirPictures=dir,nImages=1:30)
stillBack1 <- createBackground(allFullImages1)
allImages1 <- subtractBackground(bg=stillBack1)
partIden1 <- identifyParticles(sbg=allImages1,
  pixelRange=c(1,500),
  threshold=-0.1)
records1 <- trackParticles(partIden1,L=20,R=2)
## Analyse second part
allFullImages2 <- loadImages (dirPictures=dir,nImages=31:60)
stillBack2 <- createBackground(allFullImages2)
allImages2 <- subtractBackground(bg=stillBack2)
partIden2 <- identifyParticles(sbg=allImages2,
  pixelRange=c(1,500),
  threshold=-0.1)
records2 <- trackParticles(partIden2,L=20,R=2)
## Merge tracks
records <- mergeTracks(records1,records2)
plot(records,colorimages=allFullImages1,type="trajectories",incThres=10)
## End(Not run)
```

plot.TrDm

Methods for class 'TrDm'.

Description

plot methods for class 'TrDm'.

Usage

```r
## S3 method for class 'TrDm'
plot(x, frame = 1, type = NULL, incThres = NULL,
  colorimages = NULL, cl = 1, path = NULL, name = "animation",
  libavpath = NULL, ...)
```
print.summaryTrDm

Arguments

- **x**: An object of class 'TrDm'.
- **frame**: Choose which frame to be plotted. By default, frame=1.
- **type**: Only for 'tracked' objects. By default, both trajectories and size distribution are plotted. Choose 'trajectories' to plot only trajectories on the original color image. Choose 'sizes' to only plot the particle size distribution. Choose 'animation' to create an .mp4 animation. Here, images are temporarily saved in path. Set name of file with argument name.
- **incThres**: Minimum length of tracked segments for particles to be included. By default an automated threshold is calculated. Only for 'tracked' objects.
- **colorimages**: Original color images. By default, original color images are obtained from the global environment.
- **cl**: When plotting a subtracted background image, choose which color layer is plotted. By default, cl=1.
- **path**: When creating an animation, choose directory in which images are saved temporarily, and where the animation should be saved.
- **name**: of animation; by default 'animation'.
- **libavpath**: Path to location where the executable file for libav can be found (named 'avconv.exe'), in case it is not found automatically, e.g. 'C:/Users/libav/usr/bin/avconv.exe'. Only required when creating an animation.
- ... further arguments passed to plotRGB

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

print.summaryTrDm  print methods for class 'TrDm'.

Description

print methods for class 'TrDm'.

Usage

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

Arguments

- **x**: Object of class 'summaryTrDm'.
- ... Further arguments passed to or from other methods.

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser
print.TrDm

Methods for class 'TrDm'.

Description

print methods for class 'TrDm'.

Usage

## S3 method for class 'TrDm'
print(x, ...)

Arguments

x 
Object of class 'TrDm'.

... 
Further arguments passed to or from other methods.

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

runBatch

Batch analysis

Description

runBatch analyzes all image sequences in a specified directory. Use this function when settings have been optimized previously on a single or selection of movies/image sequences.

Usage

runBatch(path, settings = NULL, dirnames = NULL, nImages = 1:30, pixelRange = NULL, threshold = -0.1, qthreshold = NULL, select = "dark", nn = NULL, incThres = NULL, plotOutput = FALSE, plotType = "trajectories", L = 20, R = 2, weight = c(1, 1, 1), autoThres = FALSE, perFrame = FALSE, methodBg = "mean", frames = NULL, saveAll = FALSE)

Arguments

path 
A character vector of path name that contains all directories with image sequences.

settings 
Object of class 'tracked' containing all optimized settings in attributes, as obtained from trackParticles. Alternatively, settings can be specified using arguments described below.
runBatch

If not all image sequences should be analyzed, specify which files to use as a character string.

See loadImages

See identifyParticles

See identifyParticles

See identifyParticles

See identifyParticles

Name of artificial neural net if apply it to images. Default is NULL, resulting in no neural net being applied.

Minimum number of frames that a particle must be present. By default, automated estimate is used.

Default is FALSE. Set TRUE to plot results.

Default is 'trajectories'. Other options are 'sizes' and 'animation'.

See trackParticles

See trackParticles

See trackParticles

See createBackground

See identifyParticles

Logical. Set TRUE to save for each image sequence the full object obtained from loadImages. Default is FALSE.

Dataframe with estimated population size for each image sequence.

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

loadImages, createBackground, subtractBackground, identifyParticles, trackParticles.

## Not run:
## Simulate 3 image sequences
wd <- getwd()
folders <- paste0(rep("images",3),1:3)
populations <- c(15,25,50)
dir.create("./batchTest")
setwd("./batchTest")
for(i in 1:length(folders)){

```
simulTrajec:

Simulate trajectories and save as png files.

Description

simulTrajec simulates movement trajectories within a bounded space, movements are set with speed \( h \) and may be correlated in direction \( \rho \). Function simulates movement of particles in a video sequence of certain number of frames \( n\text{frames} \) in length. Images are saved as png files.

Usage

```r
simulTrajec(nframes = 20, nIndividuals = 10, h = 0.02, rho = 0,
            domain = "square", correctBoundary = TRUE,
            sizes = stats::runif(nIndividuals) * 0.012 + 0.01,
            staticNoise = FALSE, movingNoise = FALSE, name = "trajectory",
            path = NULL, parsMoving = list(density = 10, duration = 10, size = 1,
            speed = 10, colRange = c(0, 1)), parsStatic = list(density = 10, blur =
            TRUE, blurCoef = 0.025, sizes = NULL, col = "red"), width = 480,
            height = NULL)
```

Arguments

- **nframes**: Number of time frames (steps).
- **nIndividuals**: Number of individual trajectories.
- **h**: Displacement speed in pixels.
- **rho**: Correlation parameter for angle of displacement.
- **domain**: One of "square" or "circle", imposing a [0-1,0-1] rectangle domain, or a circular domain of radius 1, respectively. correct boundary ensure individual trajectories do not cross the domain
- **correctBoundary**: Logical. TRUE to make sure that individuals cannot leave the image.
subtractBackground

**sizes** Vector of sizes for each simulated particle of length nIndividuals.

**staticNoise** Logical. If TRUE, static noise is added.

**movingNoise** Logical. If TRUE, moving noise is added.

**name** Stem of the filename.

**path** to location where the created images should be saved. By default the working directory is used.

**parsMoving** List of parameters used to generate moving noise these include the density of noise particles (density), their duration (in n frames; duration), their size (size=1), their speed (speed=10) and the range or colors they are randomly drawn from (colRange=c(0,1)).

**parsStatic** List of parameters used to generate static noise. These include the density (per image) of noise particles (density), whether spots look blurry (blur=TRUE or blur=FALSE), a blurring coefficient (blurCoef=0.025), and the size of the spots (with sizes).

**width** of created png image. By default 480.

**height** of create png image. If NULL, width is used.

**Author(s)**

Caspar A. Hallmann, Marjolein Bruijning & Marco D. Visser

**Examples**

```r
## Not run:
dir.create("images")
## Create image sequence and save as png's in the working directory.
traj <- simulTrajec(path="images",
                   nframes=30, nIndividuals=20, domain="square",
                   h=0.01, rho=0.9,
                   sizes=runif(20, 0.004, 0.006))

## End(Not run)
```

---

**subtractBackground**  
Background subtraction

**Description**

subtractBackground subtracts each image from a previously created still background. The objects created through the function contain all changing pixels (i.e. movement).

**Usage**

```r
subtractBackground(bg, colorimages = NULL)
```
summary.TrDm

Arguments

bg          Array containing still background, as returned from createBackground.
colorimages Array containing all frames, obtained by loadImages. Default is NULL, in this case the original images are used from the global environment.

Value

Returns array of class 'TrDm' and 'sbg' with same size as images, subtracted from background.

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Examples

## Not run:
dir.create("images")
## Create image sequence
traj <- simulTrajec(path="images",
                   nframes=30, nIndividuals=20, domain="square",
                   h=0.01, rho=0.9,
                   sizes=runif(20, 0.004, 0.006))
## Load images
dir <- "images"
allFullImages <- loadImages(dirPictures=dir, nImages=1:30)
stillBack <- createBackground(allFullImages, method="mean")
allImages <- subtractBackground(stillBack)
plot(allImages)
## End(Not run)

summary.TrDm

summary methods for class 'TrDm'.

Description

summary methods for class 'TrDm'.

Usage

## S3 method for class 'TrDm'
summary(object, incThres = NULL,
        funSize = stats::median, ...)
testNN

Arguments

object       an object of class 'TrDm'.
incThres     Minimum length of tracked segments for particles to be included. By default an
             automated threshold is calculated. Only for 'tracked' objects.
funSize      Statistic to be calculated to obtain particle sizes. By default median is used.
             ... further arguments passed to or from other methods.

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Description

Fits multiple neural networks to a dataset; data set has been randomly assigned to each of three
categories: train, validate and test. A final neural net is selected based on a fit statistic (either
precision, recall or the F1-score). All neural networks are trained to the training dataset. Neural
network may vary in the number of hidden layers. Classification thresholds are selected based on
the validation data, and then the final neural network is selected based on the test data.

Usage

testNN(dat, stat = "F", maxH = 5, repetitions = 3, prop = c(8, 1,
        1), predictors = NULL, pca = TRUE, thr = 0.95, ...)

Arguments

dat          a previously constructed dataset obtained from manuallySelect.
stat          Fit statistic. May be "precision", "recall", or "F" for the harmonic mean of
             precision and recall.
maxH          maximum number of hidden layers to test note that more layers will require
             more time to fit.
repetitions   the number of repetitions for the neural network’s training.
prop          the proportion or ratio for each class c(training, validation, test).
predictors   Optional. A set of custom predictors for the neural network. Default uses all
             columns in dat.
pca          Logical. TRUE by default. Should the set of predictors be compressed to the
             most informative? In short, should a principal component analysis be conducted
to select axis that explain at least a fraction thr (see below) of the variance in
the full set of predictors?
thr           Threshold for pca (above).
...          additional parameters, passed to neuralnet.
Details

The neural networks may be selected based on precision, recall or a F1-score (default). In binary classification, precision is the number of correct positive results divided by the number of all positive predictions. Recall is the number of correct positive results divided by the number of positive results that could have been returned if the algorithm was perfect. A F1 score (F-score/ F-measure) is a statistical measure of accuracy. F1 scores considers both the precision and the recall. A F1 score may be seen as a weighted average (harmonic mean) of the precision and recall. Precision, recall and F1 scores are at best 1 and at worst 0.

Value

Returns trained artificial neural net.

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Examples

```r
## Not run:
dir.create("images")
## Create image sequence
traj <- simulTrajec(path="images",
nframes=30,nIndividuals=20,domain="square",
h=0.01,rho=0.9,movingNoise=TRUE,
parsMoving = list(density=20, duration=10, size=1,
 speed = 10, colRange = c(0,1)),
sizes=runif(20,0.004,0.006))
## Load images
dir <- "images"
allFullImages <- loadImages (dirPictures=dir,nImages=1:30)
stillBack <- createBackground(allFullImages,method="mean")
allImages <- subtractBackground(stillBack)
partIden <- identifyParticles(allImages,threshold=-0.1,
pixelRange=c(3,400))
nframes <- 3
frames <- order(tapply(partIden$patchID,partIden$frame,length),
decreasing=TRUE)[1:nframes]
mid <- manuallySelect(particles=partIden,frame=frames)
finalNN <- testNN(dat=mid,repetitions=10,maxH=4,prop=c(6,2,2))
summary(finalNN)
## End(Not run)
```

trackdem - Particle Tracking and Demography
trackParticles

**Author(s)**
Marjolein Bruijning, Caspar A. Hallmann, Marco D. Visser, Eelke Jongejans

---

**trackParticles**

*Track particles*

**Description**

trackParticles reconstructs trajectories by linking particles.

**Usage**

```r
trackParticles(particles, L = 50, R = 2, weight = c(1, 1, 1),
              costconstant = FALSE, logsizes = FALSE)
```

**Arguments**

- `particles` Object of class 'particles', obtained using `identifyParticles`.
- `L` Numeric. Maximum cost for linking a particle to another particle. When the cost is larger, particles will be not be linked (resulting in the begin or end of a segment). Default set at 50.
- `R` Integer. Link to how many subsequent frames? Default set at 2.
- `weight` Vector containing 3 weights to calculate costs. Depending on the study system, users may want to value certain elements over others. For instance, when individuals can vary in size over frames (which happens when objects move away or towards a camera) the "size" weight may be decreased. Weights are ordered as follows; first number gives the weight for differences in x and y coordinates; second number gives the weight for particle size differences; third number gives the difference between the predicted location and the observed location. The latter is calculated using the location of the identified particle in the previous frame.
- `costconstant` Logical. Default is FALSE. This increases the maximum cost L proportional to the number of images in between two frames (when R > 1). Set to TRUE keeps maximum cost L constant for all 1:R frames.
- `logsizes` Logical. Default is FALSE. Set to TRUE to take the natural logarithm of body sizes, when calculating the cost of linking two particles.

**Value**

A list of class 'TrDm' and 'records'. Use 'summary' and 'plot'.

**Author(s)**
Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser
## Examples

```
## Not run:
dir.create("images")
## Create image sequence
traj <- simulTrajec(path="images", 
nframes=30,nIndividuals=20,domain="square", 
h=0.01,rho=0.9, 
sizes=runif(20,0.004,0.006))
## Load images
dir <- "images"
allFullImages <- loadImages (dirPictures=dir,nImages=1:30)
stillBack <- createBackground(allFullImages,method="mean")
allImages <- subtractBackground(stillBack)
partIden <- identifyParticles(allImages,threshold=-0.1, 
    pixelRange=c(3,400))
records <- trackParticles(particles,L=40,R=2)
summary(records)
plot(records,type="trajectories")
## End(Not run)
```

### update.particles

**Update identified particles.**

Apply trained artificial neural network to particleStat object.

#### Usage

```
## S3 method for class 'particles'
update(object, neuralnet, pca = TRUE, 
    colorimages = NULL, sbg = NULL, ...)
```

#### Arguments

- **object**: Object of class `nnTrackdemObject`.
- **neuralnet**: Trained neural net obtained from `testNN`.
- **pca**: Logical. By default `TRUE`, indicating that a principal component analysis is performed on the predictors.
- **colorimages**: An array with the original full color images, in order to plot on the original images, obtained by `loadImages`. By default the original color images are used.
- **sbg**: Images subtracted from background, as obtained by `subtractBackground`. By default, the original subtracted images are used.
- **...**: further arguments passed to or from other methods.
Value

Data frame class 'particles', containing updated particle statistics (excluding particles that have been filtered out by the neural net).

Author(s)

Marjolein Bruijning, Caspar A. Hallmann & Marco D. Visser

Examples

```r
## Not run:
dir.create("images")
## Create image sequence
traj <- simulTrajec(path="images",
  nframes=30, nIndividuals=20, domain='square',
  h=0.01, rho=0.9, movingNoise=TRUE,
  parsMoving = list(density=20, duration=10, size=1,
    speed = 10, colRange = c(0,1)),
  sizes=runif(20, 0.004, 0.006))
## Load images
dir <- "images"
allFullImages <- loadImages(dirPictures=dir, nImages=1:30)
stillBack <- createBackground(allFullImages, method="mean")
allImages <- subtractBackground(stillBack)
partIden <- identifyParticles(allImages, threshold=-0.1,
  pixelRange=c(3,400))
nframes <- 3
frames <- order(tapply(partIden$patchID, partIden$frame, length),
  decreasing=TRUE)[1:nframes]
mId <- manuallySelect(particles=partIden, frame=frames)
finalNN <- testNN(dat=mId, repetitions=10, maxH=4, prop=c(6,2,2))
partIdenNN <- update(particles=partIden, neuralnet=finalNN)
## End(Not run)
```
Index

createBackground, 2, 13, 16
createImageSeq, 3

findThreshold, 4

identifyParticles, 5, 8, 13, 19

loadImages, 2, 7, 13, 16, 20

manuallySelect, 8
mergeTracks, 9

plot.TrDm, 10
plotRGB, 11
print.summaryTrDm, 11
print.TrDm, 12

runBatch, 12

simulTrajec, 14
subtractBackground, 5, 6, 13, 15, 20
summary.TrDm, 16

testNN, 17, 20
trackdem, 18
trackdem-package (trackdem), 18
trackParticles, 9, 12, 13, 19

update.particles, 20