# Package 'moveNT'

# April 21, 2017

<b>Fitle</b> An R package for the analysis of movement data using network theory					
<b>Version</b> 0.0.0.9000					
<b>Description</b> This package provides a series of functions to analyse movement data using network theory.					
<b>Depends</b> R (>= 3.3.2)					
License GPL (>=3)					
Encoding UTF-8					
LazyData true					
RoxygenNote 6.0.1.900	0				
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adj2stack	Calculation of network metrics				

### Description

Transform an adjancency matrix to a series of network metrics at the node-level (weight, degree, betweenness, transitivity, eccentricity) and graph level (diameter, transitivity, density, and modularity)

## Usage

```
adj2stack(adjmov, grph = T)
```

2 clustnet

#### **Arguments**

adjmov Adjacency matrix, need to be an object produced by function traj2adj

grph Whether node level metrics are to be plotted

#### Value

A raster stack object

#### **Examples**

```
traj1<-sim_mov(type="0U", npatches=3, grph=T)
stck<-adj2stack(traj2adj(traj1, res=100), grph=T)</pre>
```

clustnet

Normal mixture model for clustering of node level metrics

#### **Description**

Apply a normal mixture model to a node-level metric

#### Usage

```
clustnet(stack, id = 2, nclust = 2, grph = T)
```

#### **Arguments**

stack An object produce by the function adj2stack

id Metric to be used (2=Weight, 3=Degree, 4=Betweenness, 5=Transitivity, 6=Ec-

centricity)

grph Whether resulting classification should be plotted

#### Value

A list object containing a Mclust object and a raster object

```
traj1<-sim_mov(type="0U", npatches=3, grph=T)
stck<-adj2stack(traj2adj(traj1, res=100), grph=T)
cl<-clustnet(stck, id=2, nclust=2, grph=T)
summary(cl[[1]])</pre>
```

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graphmet

Summarize graph-level metrics

#### Description

Summarize graph-level metrics from an object generated by adj2stack

#### Usage

```
graphmet(grid)
```

#### **Arguments**

grid An object generated by the function adj2stack

id Metric to be used (2=Weight, 3=Degree, 4=Betweenness, 5=Transitivity, 6=Ec-

centricity)

#### Value

A vector

#### **Examples**

```
traj1<-sim_mov(type="OU", npatches=3, grph=T)
stck<-adj2stack(traj2adj(traj1, res=quant(traj1)), grph=T)
graphmet(stck)</pre>
```

quant

Sample quantile of distance for ltraj object

#### **Description**

Wrapper function that extract the sample quantile of distance

#### Usage

```
quant(x, p = 0.5)
```

### Arguments

x A ltraj object

p Probability, default=0.5 (median)

#### Value

A vector of length p

```
traj1<-sim_mov(type="0U", npatches=3, grph=T)
stck<-adj2stack(traj2adj(traj1, res=quant(traj1)), grph=T)</pre>
```

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SIM	mov

Simulation of patch-based movement trajectory

#### Description

Simulate a movement trajectory with user defined number of patches and interpatch movement

#### Usage

```
sim_mov(type = c("2states", "OU"), npatches = 5, ratio = 5,
  nswitch = 150, ncore = 200, spacecore = 200,
  seq_visit = sample(1:npatches, nswitch, replace = T), stepDist = "gamma",
  angleDist = "vm", stepPar = c(0.5, 3, 1, 5), anglePar = c(pi, 0, 0.5,
  2), s = diag(40, 2), grph = F)
```

#### **Arguments**

type	whether movement within patches should be based on a 2states process (from package moveHMM) or a Bivariate Ornstein-Uhlenbeck process (OU) (from package adehabitatLT)
npatches	Number of patches, default=5
ratio	Ratio (in percent) of locations associated to interpatch movement, default=5
nswitch	Number of switch/depart from patches, default=150
ncore	Number of locations within a patch per visit, default=200
spacecore	Minimum distance between center of patches, default=200
seq_visit	Specify the sequence of visit among patches, default is random sequence
stepDist	Distribution for step length if 2states specified in type, see simData of moveHMM package
angleDist	Distribution for turn angle if 2states specified in type, see simData of moveHMM package
stepPar	Parameters for step length distribution if 2states specified in type, see simData of moveHMM package
anglePar	Parameters for turn angle distribution if 2states specified in type, see simData of moveHMM package
S	Parameters for the OU process, see simm.mou of adehabitatLT package
grph	Whether a graph of the trajectory should be produced, default=F

#### Value

A ltraj (adehabitatLT) object

```
traj1<-sim_mov(type="OU", npatches=3, grph=T)
traj2<-sim_mov(type="2states", npatches=2, grph=T)</pre>
```

traj2adj 5

t	raj2adj	Generation of adjacency matrix from movement data

#### Description

Transform an Itraj object to an adjacency matrix using a user-specified grid size

#### Usage

```
traj2adj(mov, res = 100, grid = NULL)
```

#### **Arguments**

mov Movement trajectory, need to be a ltraj object

res Grid size

grid User specified grid (a raster), needs to have a larger extent than the movement

trajectory

#### Value

A list of object containing the adjacency matrix, the grid use, and patch/corridor identification (only useful if sim\_mov was used)

#### **Examples**

```
traj1<-sim_mov(type="0U", npatches=3, grph=T)
adj<-traj2adj(traj1, res=100)</pre>
```

val

Extract occupied cells in a raster object

#### **Description**

Extract only occupied cells in a raster object,

#### Usage

```
val(grid, id)
```

#### **Arguments**

grid An object generated by the function adj2stack

id Metric to be used (2=Weight, 3=Degree, 4=Betweenness, 5=Transitivity, 6=Ec-

centricity)

#### Value

A vector

6 val

```
traj1<-sim_mov(type="0U", npatches=3, grph=T)
stck<-adj2stack(traj2adj(traj1, res=quant(traj1)), grph=T)
mean(val(stck, 2))</pre>
```

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