Vignette moveNT

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Simulating movement strategies - *sim_mov*

The function *sim_mov* generates movement trajectories including patches and movement between patches. Movement within patches can follow an Ornstein-Uhlenbeck process (based on *simm.mou* function from package *adehabitatLT*) or two-states movement model (based on *simmData* function from package *moveHMM*). Movement between patches is following a brownian bridge movement model (based on *simm.bb* function from package *adehabitatLT*). Generated outputs are of the class *ltraj* from package *adehabitatlt*.

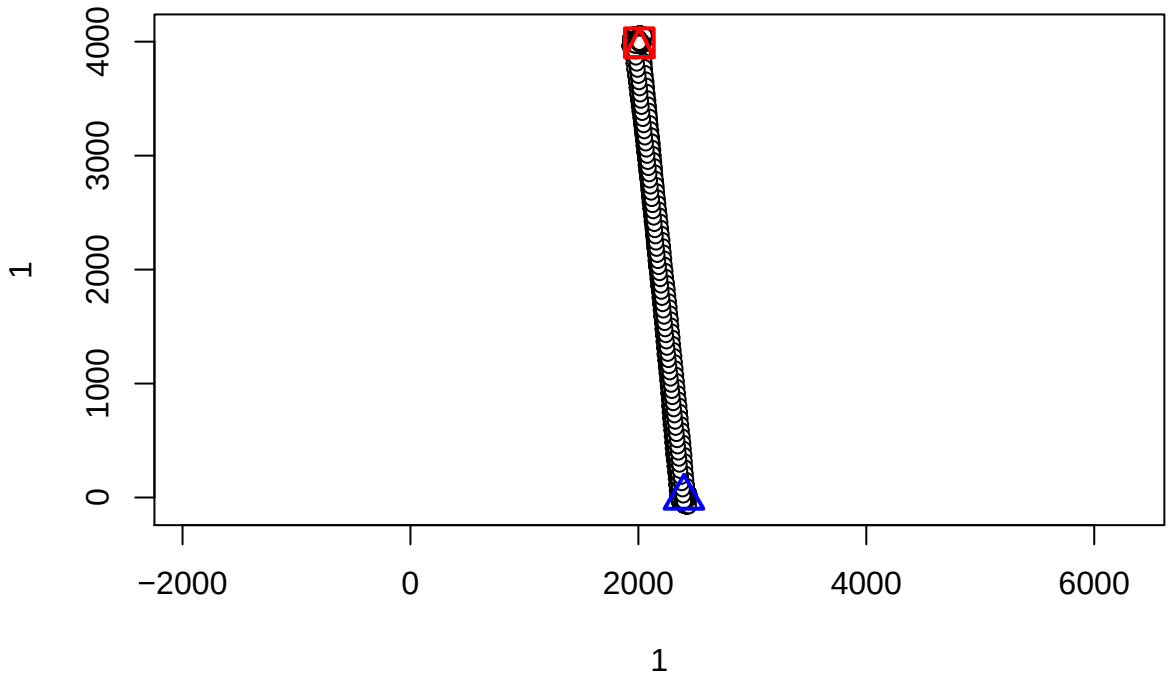
```
# Simulating migration with two-states model
mig<-sim_mov(type="2states", npatches=2, ratio=2, nswitch=25, ncore=150, grph=F)
mig
```

```
##
## ************ List of class ltraj ************
##
## Type of the traject: Type II (time recorded)
## * Time zone: GMT *
## Regular traject. Time lag between two locs: 1 seconds
##
## Characteristics of the bursts:
##   id burst nb.reloc NAs          date.begin            date.end
## 1 id    id     4575   0 1960-01-01 00:00:01 1960-01-01 01:16:15
##
##
##  infolocs provided. The following variables are available:
## [1] "out.Corri"
```

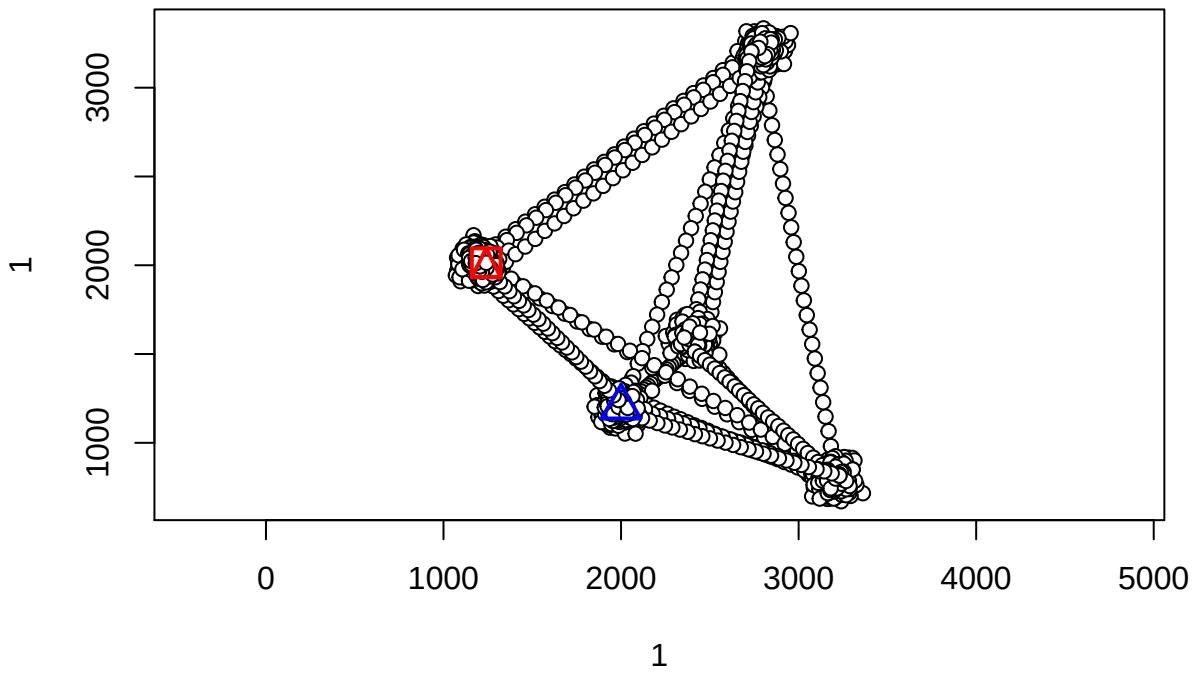
```
head(ld(mig))
```

```
##          x         y                date            dx            dy
## 1 2400.000 0.0000000 1960-01-01 00:00:01 -6.445834e-01  6.951748e-01
## 2 2399.355 0.6951748 1960-01-01 00:00:02  4.959303e-01  5.159112e-01
## 3 2399.851 1.2110860 1960-01-01 00:00:03  9.231926e-01  2.499306e-01
## 4 2400.775 1.4610166 1960-01-01 00:00:04  7.173123e-06 -1.963805e-06
## 5 2400.775 1.4610146 1960-01-01 00:00:05  3.374780e-01  6.426785e-02
## 6 2401.112 1.5252825 1960-01-01 00:00:06 -5.498167e-01  4.466795e+00
##           dist dt       R2n  abs.angle  rel.angle id burst out.Corri
## 1 9.480273e-01  1 0.0000000  2.3184508         NA id    id         2
## 2 7.156194e-01  1 0.8987557  0.8051427 -1.5133081 id    id         2
## 3 9.564255e-01  1 1.4888270  0.2643868 -0.5407560 id    id         2
## 4 7.437084e-06  1 2.7344809 -0.2672248 -0.5316116 id    id         2
## 5 3.435430e-01  1 2.7344862  0.1881824  0.4554072 id    id         2
## 6 4.500506e+00  1 3.5630855  1.6932700  1.5050876 id    id         2
```

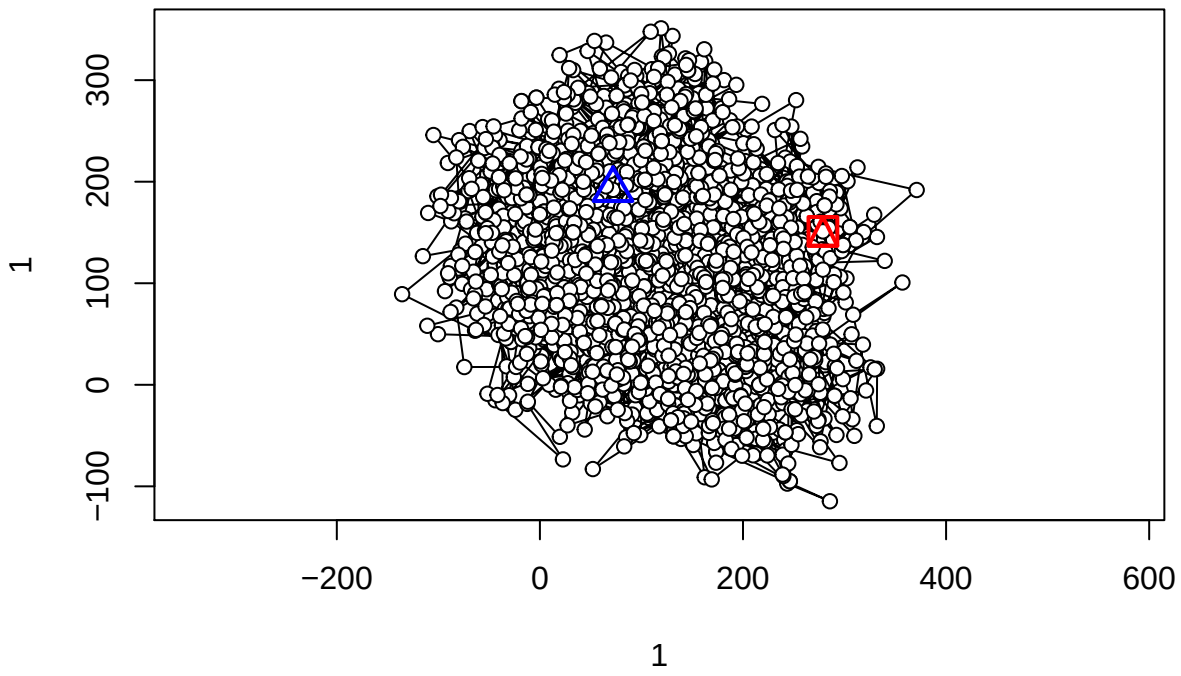
```
plot(mig)
```




```
# Simulating multi-patches movement with Ornstein-Uhlenbeck process
patches<-sim_mov(nswitch=25, ncore=150, ratio=5, type="OU", npatches=5, grph=T)
```




```
# Simulating sedentary movement
seden<-sim_mov(type="OU", npatches=10, spacecore=12, ratio=3, nswitch=150, ncore=20, grph=T)
```



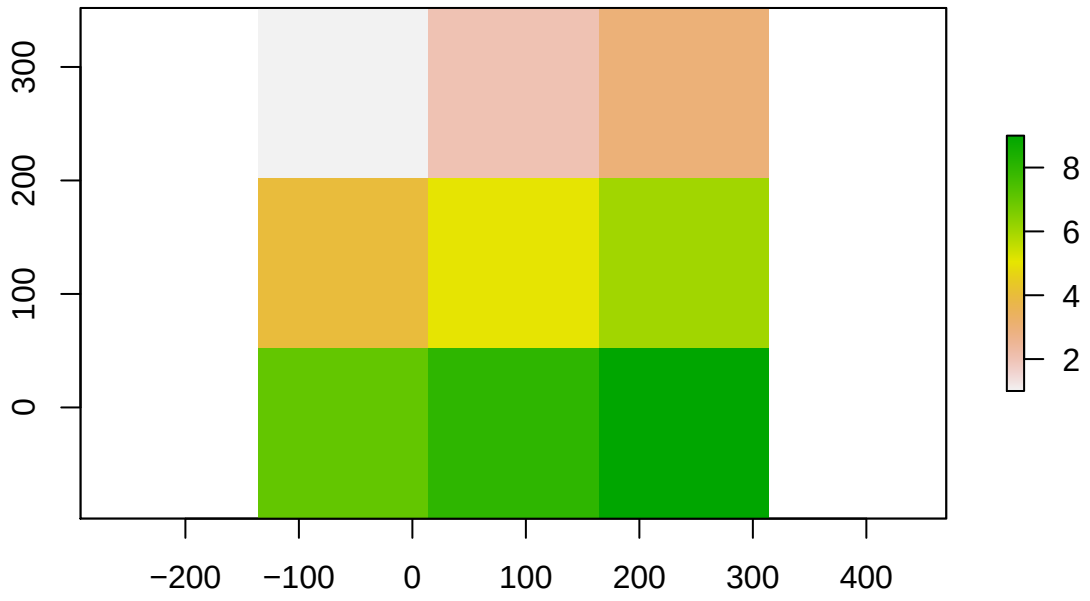
Converting movement to adjacency matrix - *traj2adj*

The function *traj2adj* converts a trajectory object of class *ltraj* to an adjacency matrix. This is done by overlapping a grid over the relocation data and tallying the number of transitions among each pixel. Users need to specify the grid size, which can be based on distance travelled. The function *quant* is a wrapper that allows to sample a quantile of step length distribution from a *ltraj* object. Output produced by *traj2adj* is a list containing the adjacency matrix, the grid used (raster format), and a raster indicating pixel numbers that are occupied. These rasters are used by other functions such as *adj2stack* and *clustnet*.

```
# Using sedentary movement and user specific grid-size
adj_seden<-traj2adj(seden, res=150) #Pixel size of 150m
adj_seden[[1]] # Adjency matrix
```

```
##       [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9]
##  [1,]   21   13    0   14   14    0    0    0    0
##  [2,]   12  279   17    8   84    8    0    0    0
##  [3,]    0   19   19    0    5   23    0    0    0
##  [4,]   18   11    0  201  128    0   10    7    0
##  [5,]   11   81    4  120  973   73   15  134   17
##  [6,]    0    5   27    0   60  312    0   20   51
##  [7,]    0    0    0   14   10    0   13   11    0
##  [8,]    0    0    0   18  131   13   10  261   58
##  [9,]    0    0    0    0   22   47    0   59  247
```

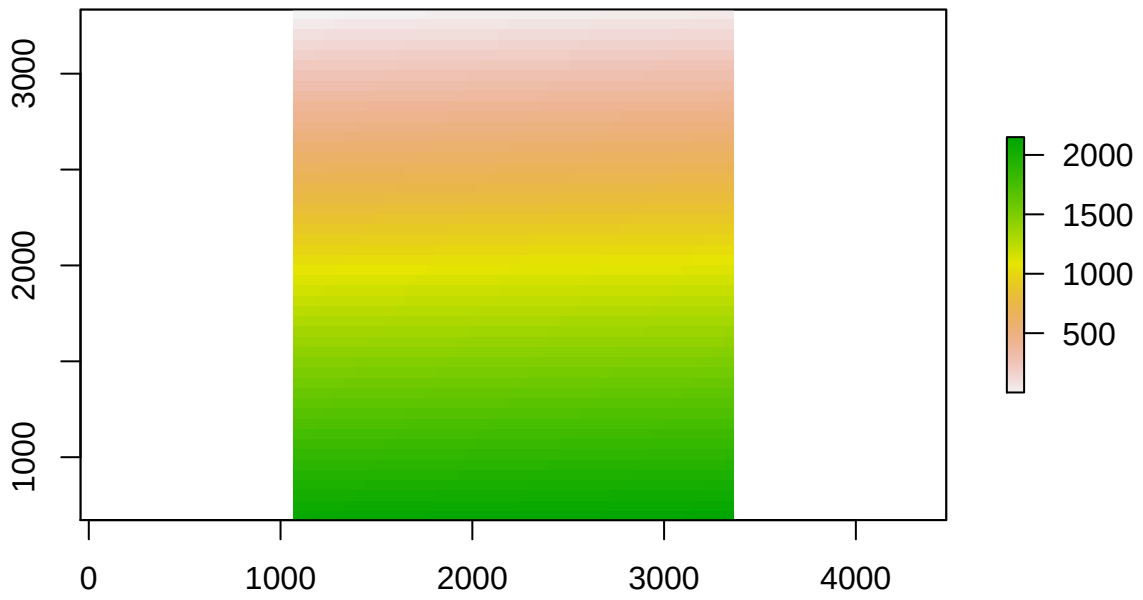
```r
plot(adj_seden[[2]]) #Plot grid used
```



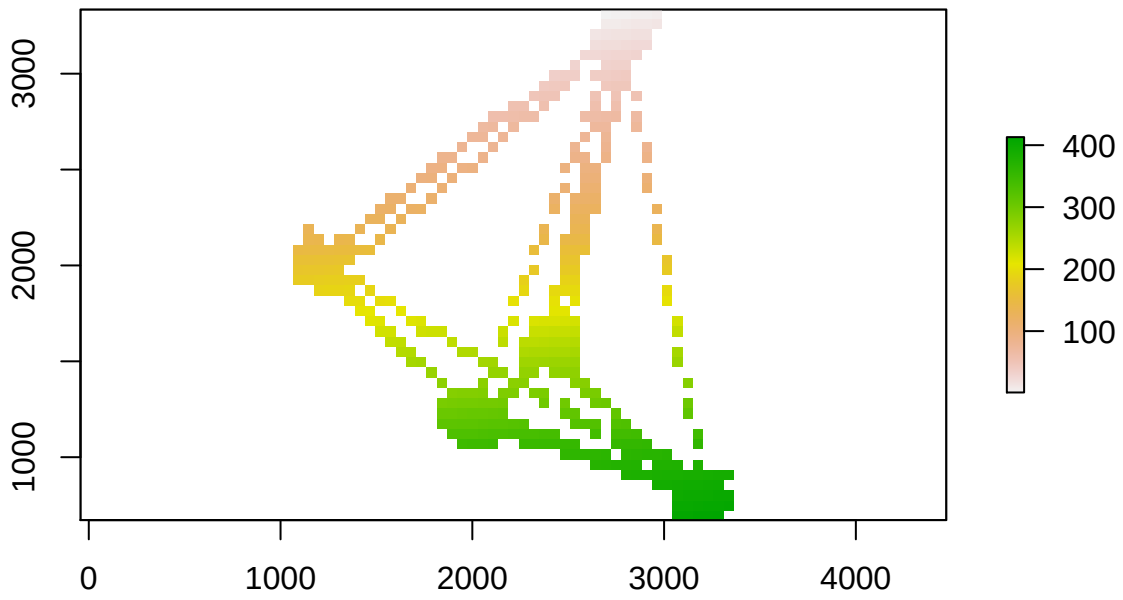

```r
# Using multi-patches movement and median distance travelled
adj_patches<-traj2adj(patches, res=quant(patches, p=0.5)) #Grid size based on median
dim(adj_patches[[1]]) # Size of the adjacency matrix
```

```
## [1] 413 413
```

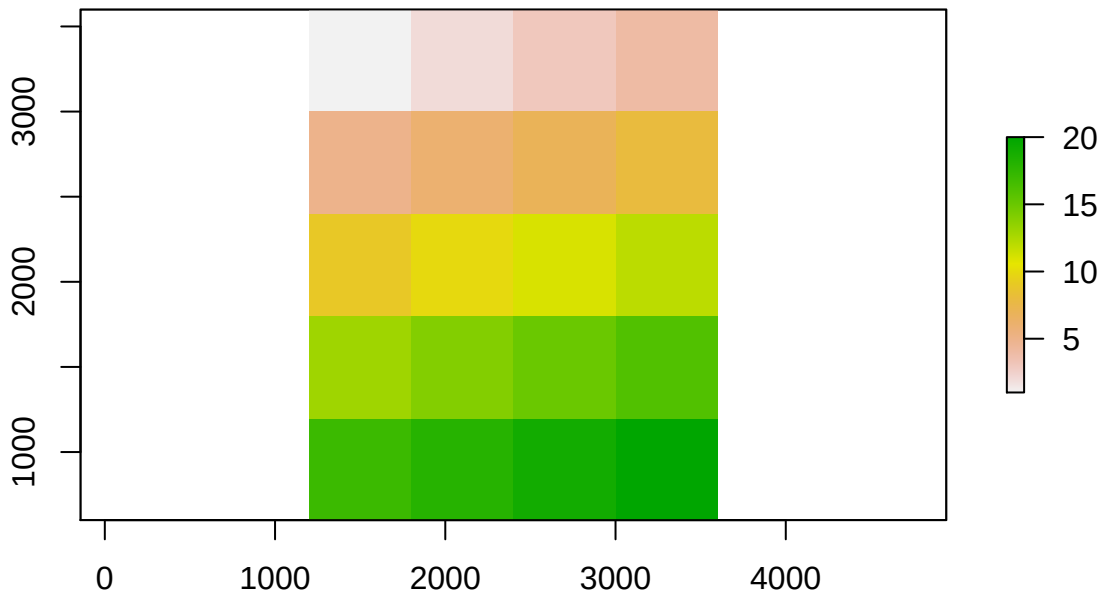
```r
plot(adj_patches[[2]]) #Plot grid used
```




```
plot(adj_patches[[3]]) #Plot occupied pixels
```



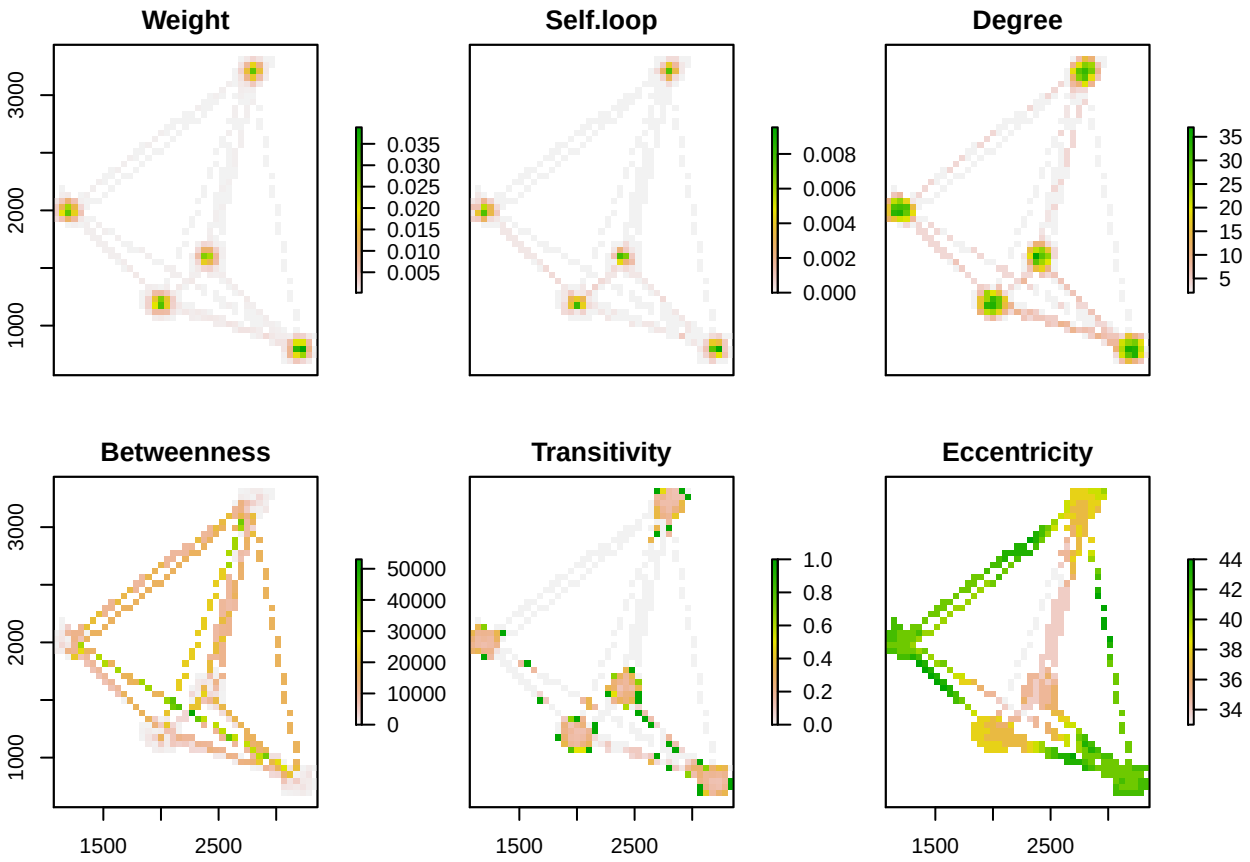

```
# Using user defined grid
ras<-raster(nrows=10, ncols=10, xmn=0, ymn=0, xmx=6000, ymx=6000)
adj_patches2<-traj2adj(patches, res=quant(patches, p=0.5), grid=ras) #Grid size based on median
plot(adj_patches2[[2]]) #Crop version of the grid created
```



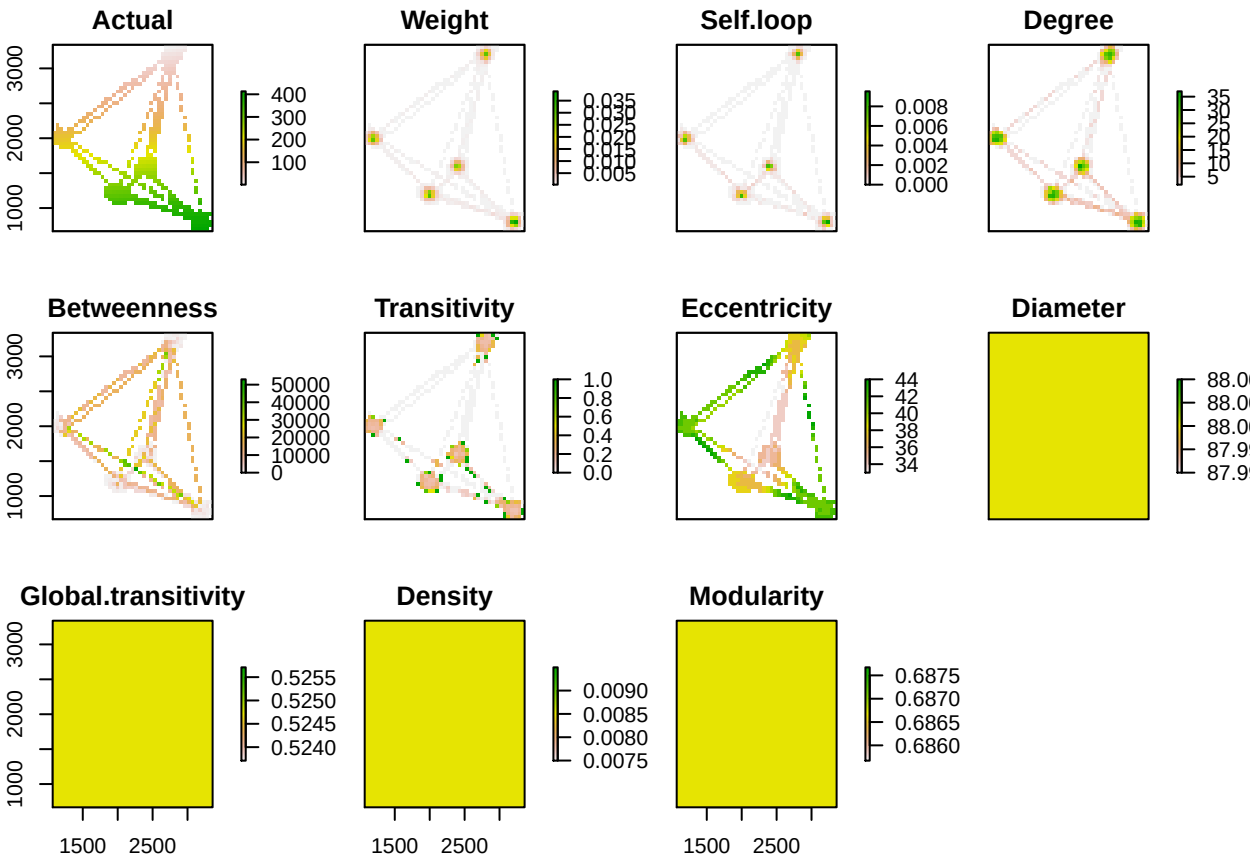
Calculation of network metrics - *adj2stack*

The function *adj2stack* takes the output of function *traj2adj* and calculates a series of node- and graph-level metrics. Each metric is stored as a individual raster and the output is a raster stack combining each metric. Graph-level metrics are also stored as a raster, each containing an unique value. The function *graphmet* extracts graph-level metrics. The function *val* extracts only the occupied cells (remove NA) in a raster and allows the calculation of statistics from node-level metrics.

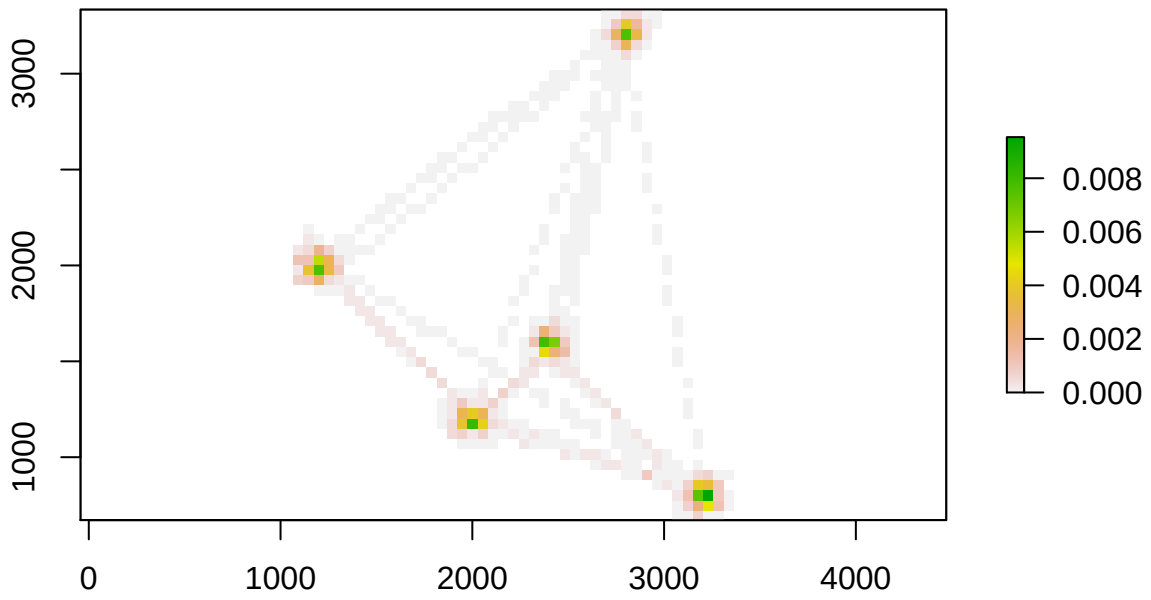
```
# Using multi-patches movement and median distance travelled
stck<-adj2stack(adj_patches,grph=T) #Plot the node-level metrics at the same time
```




```
plot(stck) #Plot also the graph-level metrics (not really useful)
```




```
plot(stck[[3]]) #Plot only one metric (degree)
```




```
graphmet(stck) # Extract graph-level metrics
```

```
##            Diameter Global.transitivity             Density
##        88.000000000         0.524710270         0.008498084
##          Modularity
##         0.686672390
```

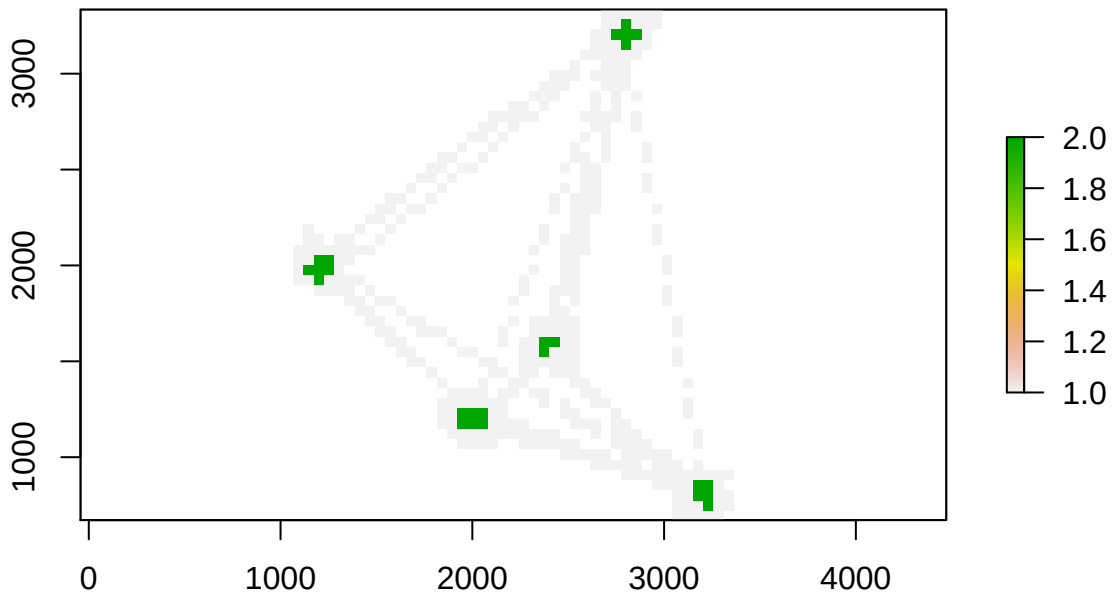
```
cv(val(stck, 4)) #Extract coefficient of variation of node-level betweenness.
```

```
## [1] 116.8318
```

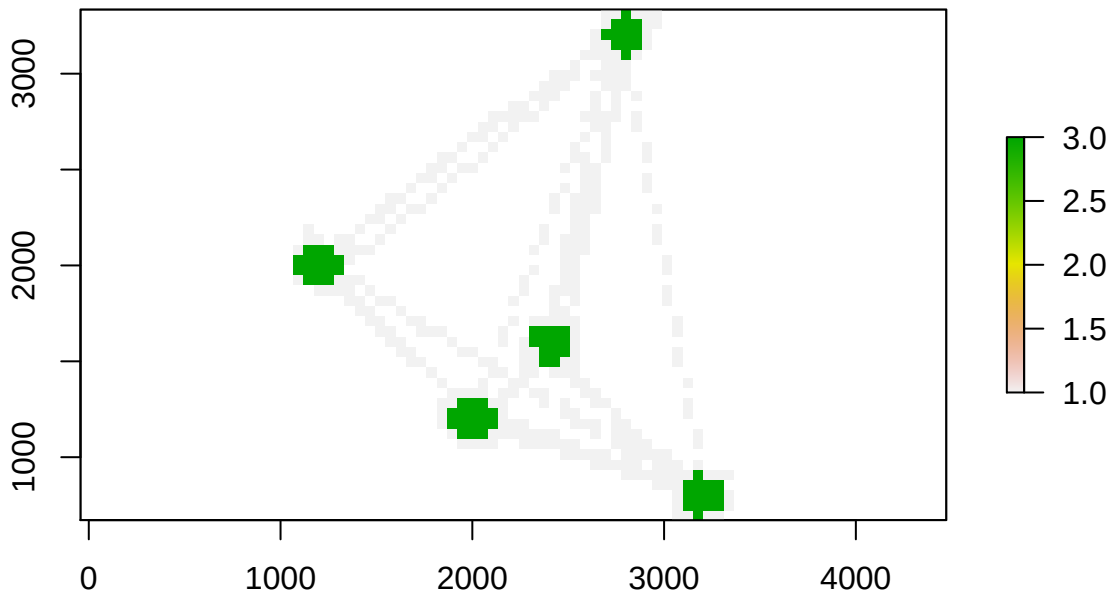
Clustering of node level metrics - *clustnet*

The function *clustnet* applies a normal mixture model to node-level metrics in order to cluster them into separate groups (default = 2). The function takes the output of function *adj2stack* with the user specifying the metric to cluster and the number of groups. Return a list containing output of function *Mclust* from package *mclust* and a raster displaying classification.

```
# Using multi-patches movement and median distance travelled
clust2<-clustnet(stck, id=3, nclust=2) # Clustering of degree in two groups
```



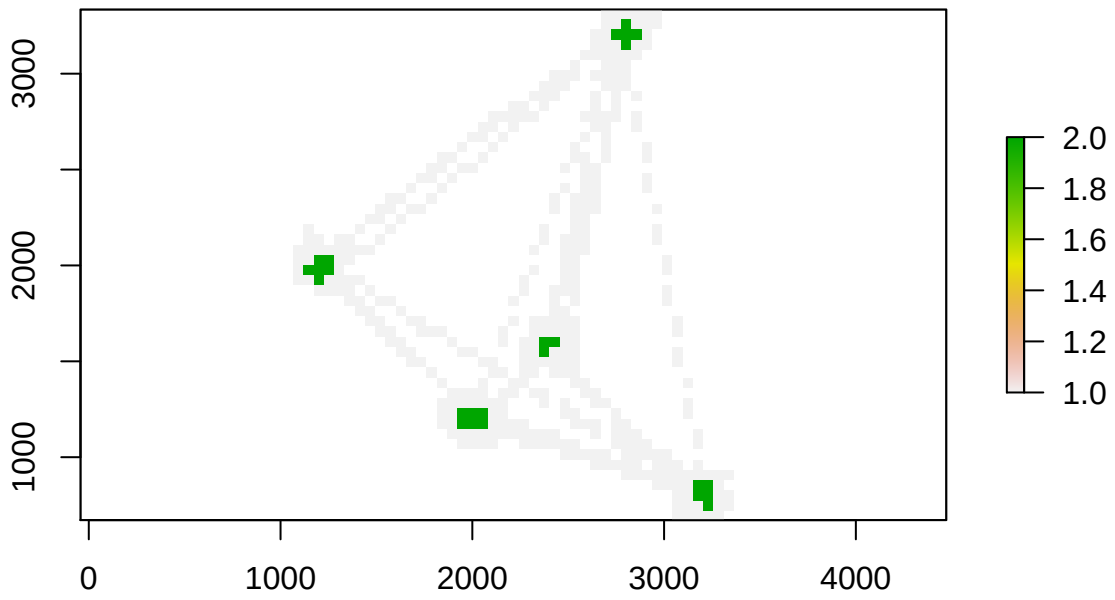

```
clust3<-clustnet(stck, id=4, nclust=3) #Clustering of betweenness in three groups
```




```
summary(clust2[[1]])
```

```
## ----------------------------------------------------
## Gaussian finite mixture model fitted by EM algorithm
## ----------------------------------------------------
##
## Mclust E (univariate, equal variance) model with 2 components:
##
##  log.likelihood   n df      BIC      ICL
##        2390.769 413  4 4757.445 4756.719
##
## Clustering table:
##   1   2
## 388  25
```

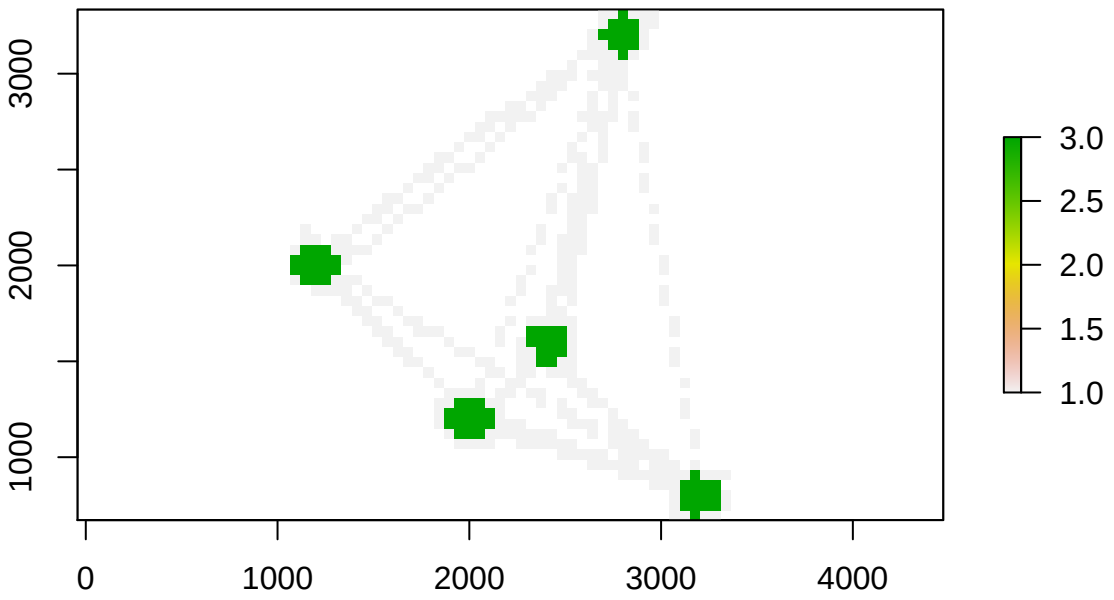
```
plot(clust2[[2]])
```




```
summary(clust3[[1]])
```

```
## ----------------------------------------------------
## Gaussian finite mixture model fitted by EM algorithm
## ----------------------------------------------------
##
## Mclust E (univariate, equal variance) model with 3 components:
##
##  log.likelihood   n df       BIC       ICL
##       -1269.247 413  6 -2574.635 -2895.684
##
## Clustering table:
##   1   2   3
## 342   0  71
```

```
plot(clust3[[2]])
```


3000
2000
1000
0
1000
2000
3000
4000
3.0
2.5
2.0
1.5
1.0