

4.7 Characteristic Hull Polygons (CHP)

Manual of Applied Spatial Ecology

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Now we are going to get into another class of home range estimators that use polygons created by Delaunay triangulation of a set of relocations and then removing a subset of the resulting triangles. These polygons can have concave edges, be composed of disjoint regions, and contain empty portions of unused space within hull interiors. This estimator has been described in the `adehabitatHR` package and evaluated on black-footed albatross (*Phoebastria nigripes*; Downs and Horner 2009). Polygon-based estimators may be a useful method for a variety of species but research has been limited.

1. Exercise 4.7 - Download and extract zip folder into your preferred location
2. Set working directory to the extracted folder in R under Session - Set Working Directory...
3. Now open the script “CHPscript.Rmd” and run code directly from the script
4. First we need to load the packages needed for the exercise

```
library(adehabitatHR)
library(maptools)
```

5. Now let's have a separate section of code to include projection information we will use throughout the exercise. In previous versions, these lines of code were within each block of code

```
utm17.crs <- CRS("+proj=utm +zone=17N +ellps=WGS84")
```

```
## Warning in showSRID(uprojargs, format = "PROJ", multiline = "NO", prefer_proj
## = prefer_proj): Discarded datum Unknown based on WGS84 ellipsoid in Proj4
## definition
```

6. Load panther dataset

```
#Creates a Spatial Points Data Frame for 2 animals by ID
twocats <- read.csv("pantherjitter.csv", header=T)
data.xy = twocats[c("X", "Y")]
```

```
#Creates class Spatial Points for all locations
xysp <- SpatialPoints(data.xy)
proj4string(xysp) <- CRS("+proj=utm +zone=17N +ellps=WGS84")
```

```
#Creates a Spatial Data Frame from
sppt<-data.frame(xysp)
```

```
#Creates a spatial data frame of ID
idsp<-data.frame(twocats[1])
```

```
#Merges ID data frame with GPS locations data frame
#Data frame is called "idsp" comparable to the "relocs" from puechabon dataset
coordinates(idsp)<-sppt
```

```
head(as.data.frame(idsp))
```

```
##   CatID      X      Y
## 1   121 494155.6 2904240
## 2   121 498182.3 2905598
## 3   121 498476.2 2905114
## 4   121 499210.5 2905661
## 5   121 499467.3 2905533
## 6   121 502960.9 2904391
```

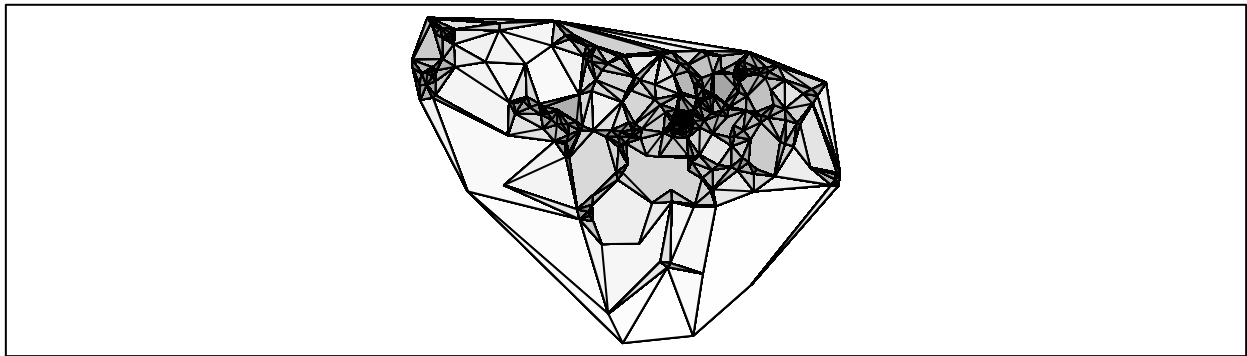
```
#Results for above code
```

```
#Home Range estimation
res <- CharHull(idsp[,1])
class("res")
```

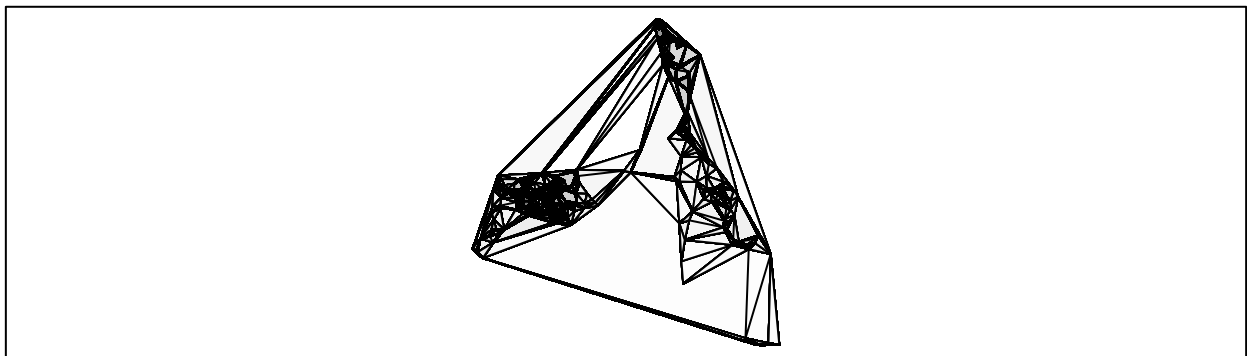
```
## [1] "character"
```

```
#Displays the home range
plot(res)
```

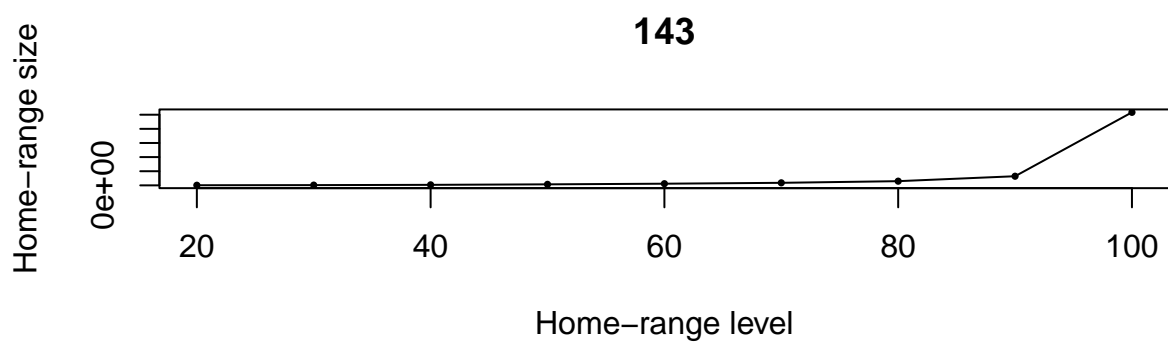
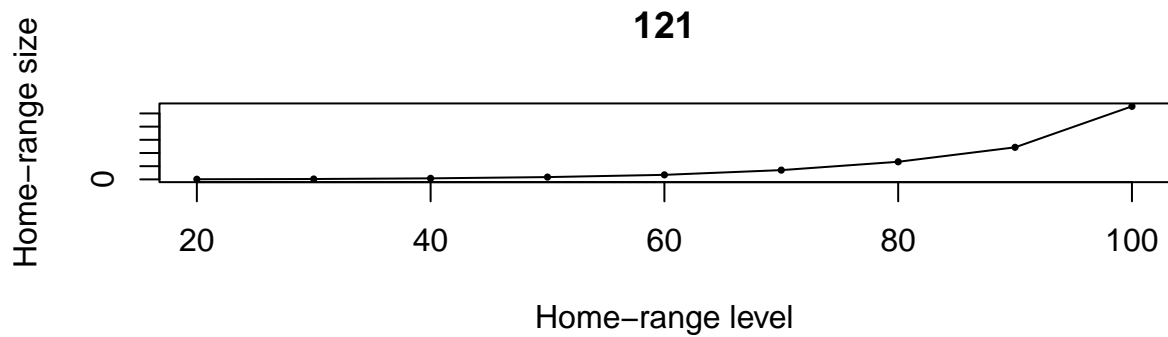
121



143



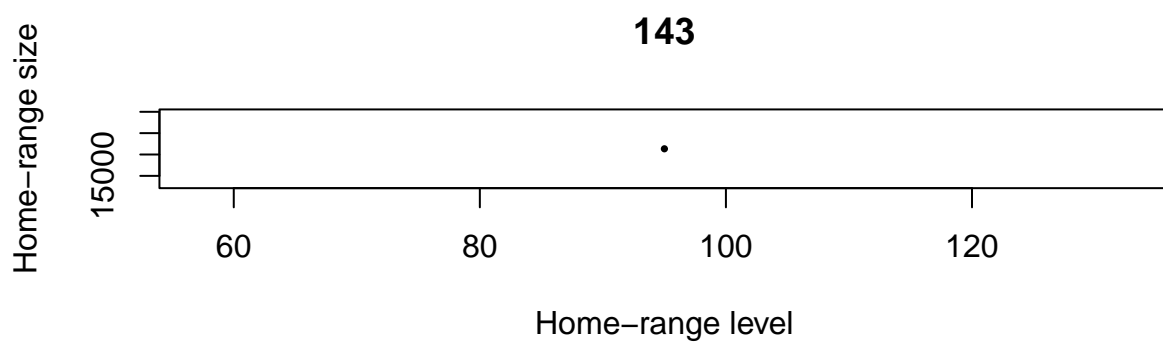
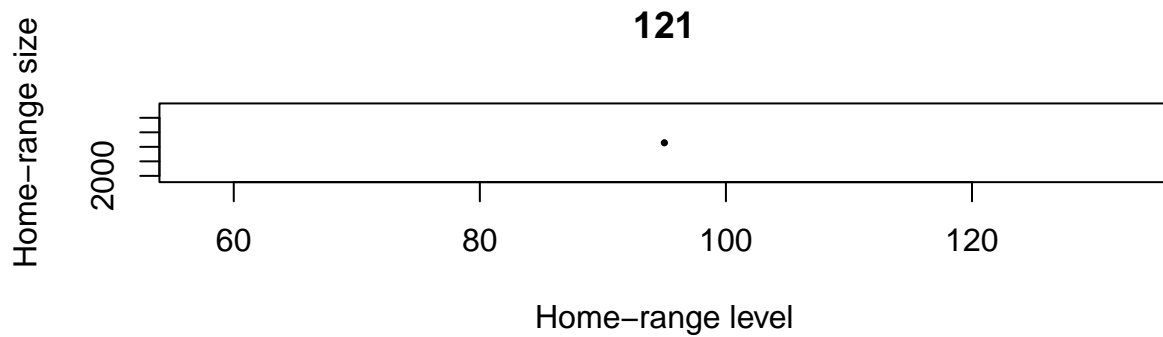
```
#Computes the home range size for 20-100 percent
MCHu2hrsize(res)
```



```
##          121          143
## 20      10.70262    150.2998
## 30      28.73158    342.9798
## 40      79.15079    688.7553
## 50     172.95317   1418.7588
## 60     343.12024   2319.1578
## 70     700.94072   3574.5525
## 80    1333.94110   5910.8221
## 90    2431.27046  12912.0320
## 100  5538.22402 103361.1336
```

#OR

#Computes the home range size for 95 percent
MCHu2hrsize(res, percent=95)



```
##          121      143
## 95 3139.295 21335.9
```

```
#OR use
```

```
ver <- getverticeshr(res, percent=90)
ver
```

```
## Object of class "SpatialPolygonsDataFrame" (package sp):
```

```
##
```

```
## Number of SpatialPolygons: 2
```

```
##
```

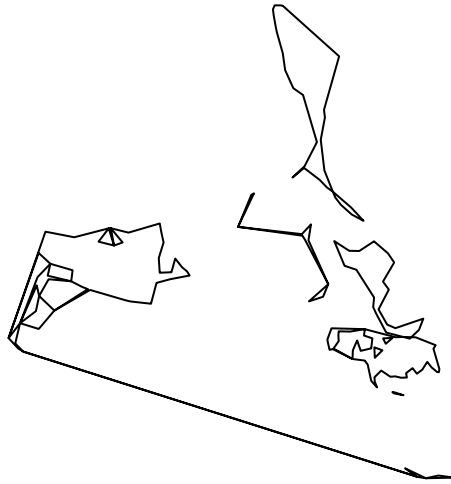
```
## Variables measured:
```

```
##   id    area
```

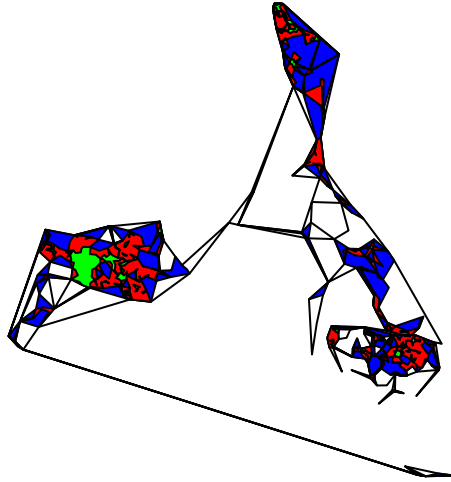
```
## 1 121 2413.518
```

```
## 2 143 12845.298
```

```
plot(ver)
```



```
ver50 <- getverticeshr(res, percent=50)
ver80 <- getverticeshr(res, percent=80)
ver90 <- getverticeshr(res, percent=90)
ver95 <- getverticeshr(res, percent=95)
plot(ver95)
plot(ver90, add=TRUE, col="blue")
plot(ver80, add=TRUE, col="red")
plot(ver50, add=T, col="green")
```



*#The object uu below is the single-linkage cluster analysis that estimates
#home range returned as a list of SpatialPolygonDataFrame objects (one per animal)*

```
uu <- clusthr(idsp)
class(uu)
```

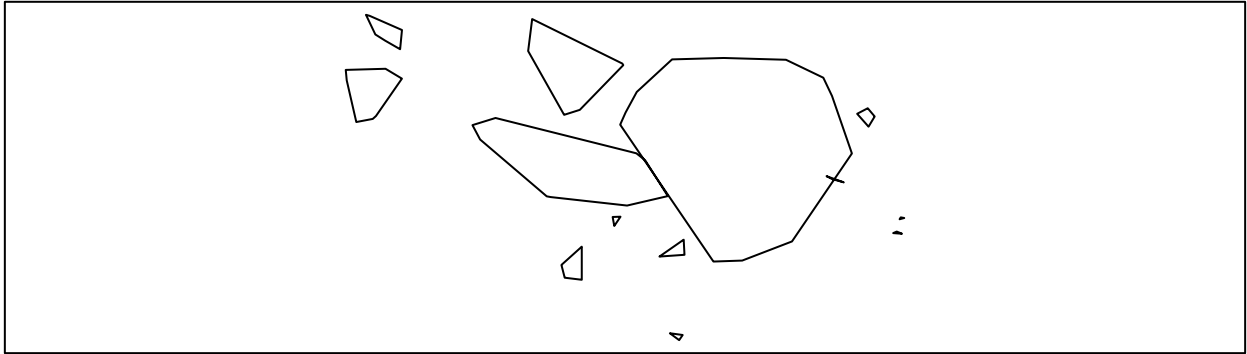
```
## [1] "MCHu"
```

```
#[1] "MCHu"
```

```
plot(uu, percent=95)
```

```
#Returns home range of Cat143 and 95% HR  
plot(uu[[1]][250,], add=T, col="green")
```

121



143

