Exercise 6 - Self Organising Map (in R)

November 15, 2017

First model map output:

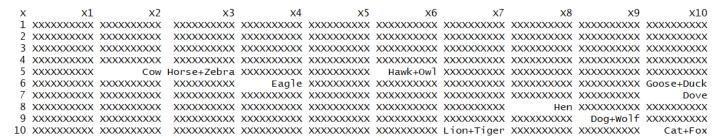


Figure 1 - Output map from SOM algorithm

First model human-clustered map:

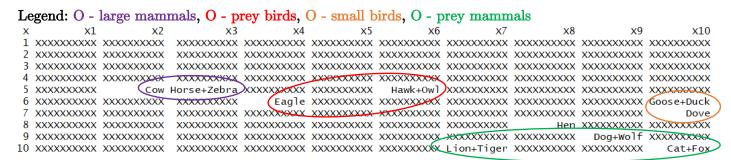


Figure 2 - Manually annotated map from SOM algorithm

Second model map output:

X	x1	x2	x 3	x4	x5	x6	x 7	x8	x9	x10
1	XXXXXXXXX	XXXXXXXXX	XXXXXXXXX	Tiger	XXXXXXXXX	XXXXXXXXX	XXXXXXXXX	XXXXXXXXX	XXXXXXXXX	Cat
2	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	xxxxxxxxxx	XXXXXXXXXX	xxxxxxxxxx	XXXXXXXXX
3	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXX
4	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	Fox	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXX
5	Eagle	XXXXXXXXXX	XXXXXXXXX							
6	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	Cow
7	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	Owl+Hawk	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	Lion	XXXXXXXXXX	XXXXXXXXX
8	XXXXXXXXX	Dove	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXXX	XXXXXXXXX	Wolf	XXXXXXXXX	XXXXXXXXX
9	XXXXXXXXX	XXXXXXXXXX	XXXXXXXXX							
10	Hen	XXXXXXXXXX	XXXXXXXXXX	Duck+Goose	XXXXXXXXXX	XXXXXXXXXX	Dog	XXXXXXXXX	XXXXXXXXXX	Horse+Zebra

Figure 3 - Output map from refined SOM algorithm

Final model map output:

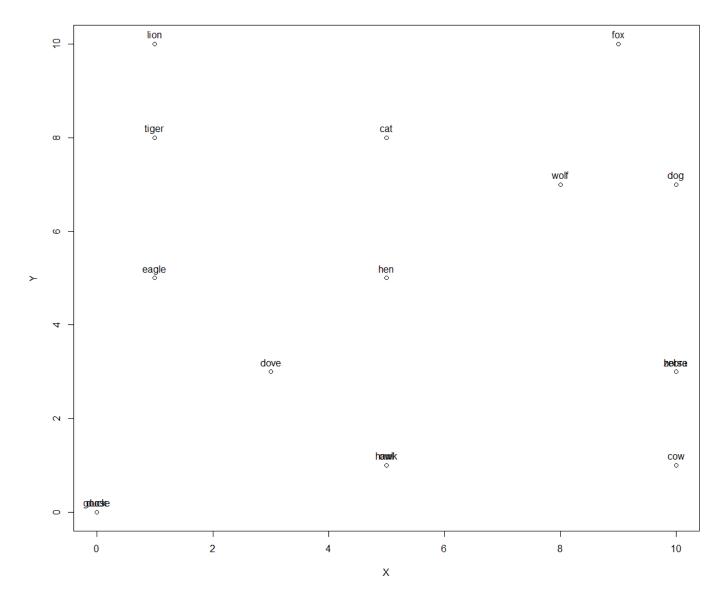


Figure 4 - Output map from final SOM algorithm

```
Script:
#Author: Thomas Hollis
#Subject: Bachelor Thesis
#1. Data Import
data <- read.csv("R/AnimalsSOMnolab.csv")</pre>
#2. Global variables
t <- 1
d <- matrix(0, 10, 10)</pre>
pattern_size <- 13</pre>
map <- matrix(list(), 10, 10)</pre>
positionx <- integer(16)</pre>
positiony <- integer(16)</pre>
SOM <- matrix(0,2,16)
colnames(SOM) <- c("dove", "hen", "duck", "goose", "owl", "hawk", "eagle", "fox",</pre>
"dog", "wolf", "cat", "tiger", "lion", "horse", "zebra", "cow")
rownames(SOM) <- c("X", "Y")</pre>
#3. Set random map weights
for (i in 1:10)
  for (j in 1:10)
    map[[i,j]] <- runif(13, 0, 1)</pre>
}
#4. Pick animal at random
for (t in 1:10000)
  r <- round(runif(1, 1, 16))
  x <- data[r,]
  X \leftarrow t(x)
    for (i in 1:10)
    for (j in 1:10)
      w <- map[[i,j]]</pre>
      W \leftarrow t(w)
      d[i,j] <- sqrt((as.numeric(x)-as.numeric(W))%*%(as.numeric(X)-as.numeric(w)));</pre>
    }
  }
  distance \leftarrow d[1,1]
```

for (i in 1:10)

for (j in 1:10)

u <- i v <- j

} }

if (d[i,j] < distance)</pre>

distance <- d[i,j]</pre>

{

```
}
  #setting learning parameters
  theta <- 1;
  alpha <- 100/(200+t);
  for (i in 1:10)
    for (j in 1:10)
    {
      eta \leftarrow \exp(-(((i-u)^2)+(j-v)^2)/2*theta^2)
      map[[i,j]] <- map[[i,j]] + alpha*eta*(X-map[[i,j]])</pre>
    }
  }
}
#TESTING PHASE
for (test in 1:16)
  y <- data[test,]</pre>
  Y \leftarrow t(y)
  for (i in 1:10)
    for (j in 1:10)
      w <- map[[i,j]]</pre>
      W \leftarrow t(w)
      d[i,j] <- sqrt((as.numeric(y)-as.numeric(W))%*%(as.numeric(Y)-as.numeric(w)))</pre>
  }
  distance \leftarrow d[1,1]
  for (i in 1:10)
    for (j in 1:10)
      if (d[i,j] < distance)</pre>
         distance <- d[i,j]</pre>
         positionx[test] <- i</pre>
         positiony[test] <- j</pre>
      }
    }
  }
}
#Mapping each animal
for(i in 1:16)
  SOM[1,i] <- positionx[i]</pre>
  SOM[2,i] <- positiony[i]</pre>
}
plot(t(SOM))
text(t(SOM), labels = colnames(SOM), pos = 3)
```