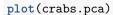
Selvstudie 1 - PCA/C4.5

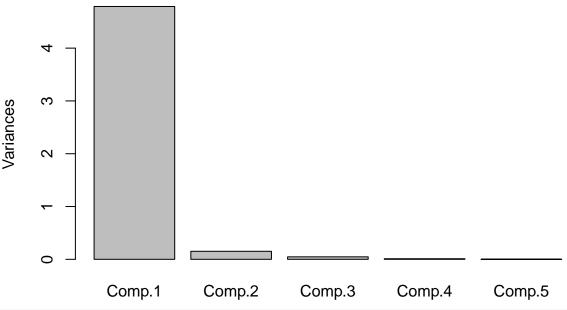
Nicholas Fitzhugh
9/2/2017

Opgave 1 (PCA)

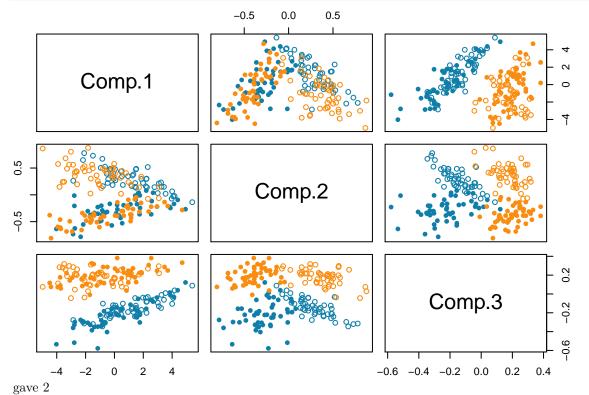
```
library(MASS)
library(tidyverse)
## Warning: package 'tidyverse' was built under R version 3.3.2
## Loading tidyverse: ggplot2
## Loading tidyverse: tibble
## Loading tidyverse: tidyr
## Loading tidyverse: readr
## Loading tidyverse: purrr
## Loading tidyverse: dplyr
## Warning: package 'ggplot2' was built under R version 3.3.2
## Warning: package 'tidyr' was built under R version 3.3.2
## Conflicts with tidy packages ------
## filter(): dplyr, stats
## lag():
            dplyr, stats
## select(): dplyr, MASS
data(crabs); head(crabs)
    sp sex index FL RW CL
## 1 B M 1 8.1 6.7 16.1 19.0 7.0
## 2 B M
              2 8.8 7.7 18.1 20.8 7.4
## 3 B M
              3 9.2 7.8 19.0 22.4 7.7
## 4 B M
             4 9.6 7.9 20.1 23.1 8.2
## 5 B M
             5 9.8 8.0 20.3 23.0 8.2
              6 10.8 9.0 23.0 26.5 9.8
## 6 B M
crabs.pca <- princomp(crabs[,4:8], cor = TRUE)</pre>
loadings(crabs.pca)
##
## Loadings:
     Comp.1 Comp.2 Comp.3 Comp.4 Comp.5
## FL -0.452 -0.138 0.531 0.697
## RW -0.428 0.898
## CL -0.453 -0.268 -0.310
                                -0.792
## CW -0.451 -0.181 -0.653
                                 0.575
## BD -0.451 -0.264 0.443 -0.707 0.176
##
                 Comp.1 Comp.2 Comp.3 Comp.4 Comp.5
## SS loadings
                   1.0
                          1.0
                                1.0
                                       1.0
                                              1.0
                   0.2
                                 0.2
                                       0.2
                                              0.2
## Proportion Var
                          0.2
## Cumulative Var
                   0.2
                          0.4
                                 0.6
                                       0.8
                                              1.0
```



crabs.pca

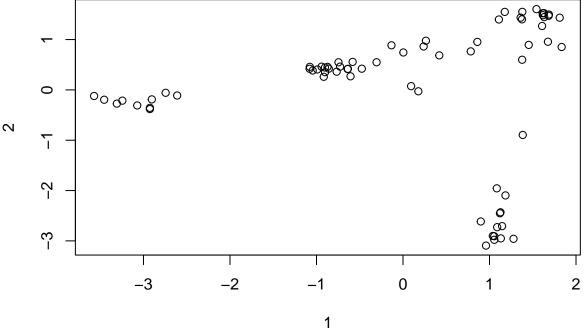


```
num_cols <- c("FL","RW","CL","CW","BD")
ccol <- function(sp) ifelse(sp=="B","#0f7fa9","#fa8d0f")
cpch <- function(sx) 1+ 15*(crabs$sex=="M")
pairs(crabs.pca$scores[,1:3], col=ccol(crabs$sp), pch=cpch(crabs$sex))</pre>
```

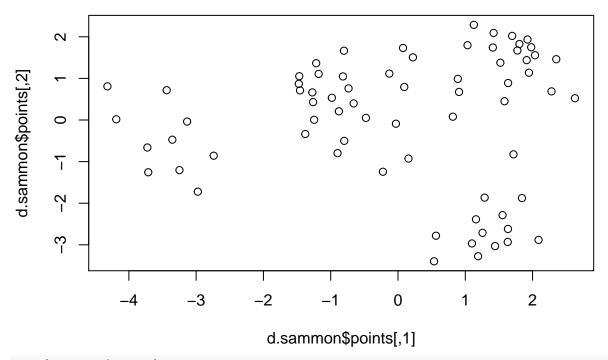


##Op-

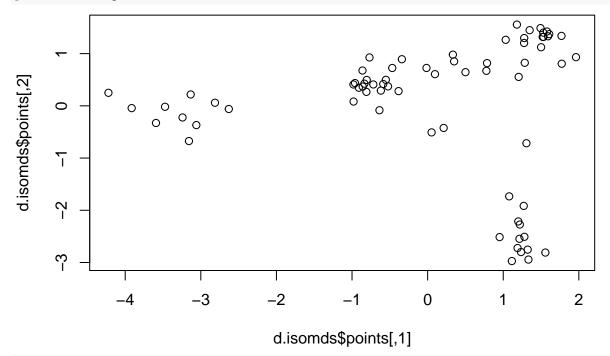
```
aims_freq <- read.csv("aims_freq.csv", header = TRUE)</pre>
d <- dist(aims_freq[,4:100])</pre>
d.mds <- cmdscale(d)</pre>
d.sammon \leftarrow sammon(d, y = d.mds)
## Initial stress
                          : 0.07568
## stress after 10 iters: 0.04823, magic = 0.092
## stress after 20 iters: 0.03480, magic = 0.213
## stress after 30 iters: 0.02931, magic = 0.150
## stress after 40 iters: 0.02914, magic = 0.500
## stress after 50 iters: 0.02909, magic = 0.500
d.isomds <- isoMDS(d)</pre>
## initial value 9.861148
## iter 5 value 7.054416
## final value 6.902732
## converged
plot(d.mds, xlab = "1", ylab = "2")
```



plot(d.sammon\$points)

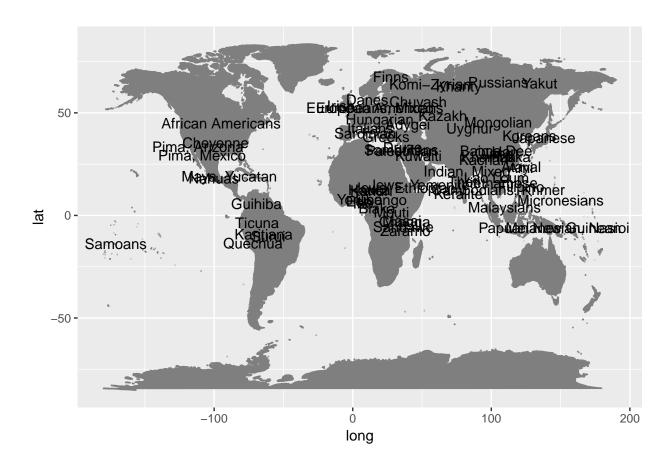


plot(d.isomds\$points)



```
ggplot() + borders("world", colour="gray50", fill="gray50") +
geom_text(data = aims_freq, aes(x = long, y = lat, label = pop))
##
```

```
## Attaching package: 'maps'
## The following object is masked from 'package:purrr':
##
## map
```

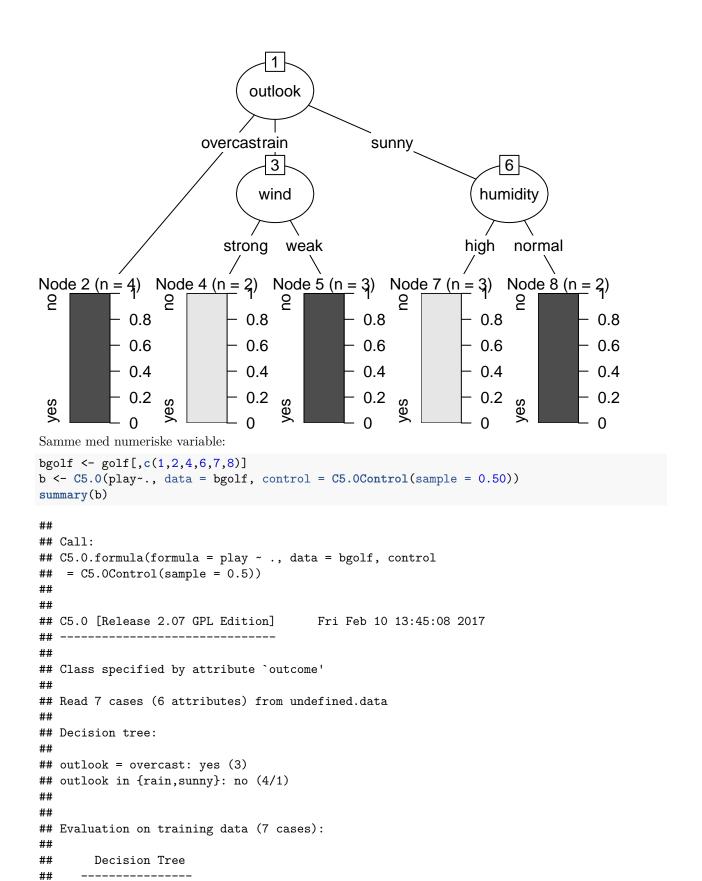


C4.5

Opgave 3

```
golf <- read.csv("golf.csv", header = TRUE)

a)
library(C50)
cgolf <- golf[,c("outlook","temperature","humidity","wind","play")]
a <- C5.0(play~.,data=cgolf)
plot(a)</pre>
```



##

##

Size

Errors

```
1(14.3%) <<
##
##
##
##
       (a)
            (b)
                    <-classified as
##
##
         3
                    (a): class no
##
               3
                    (b): class yes
##
##
##
    Attribute usage:
##
##
   100.00% outlook
##
##
## Evaluation on test data (7 cases):
##
##
        Decision Tree
##
##
      Size
              Errors
##
         2
##
             4(57.1%)
                         <<
##
##
##
       (a)
            (b)
                    <-classified as
##
##
         2
                    (a): class no
             1
##
         4
                    (b): class yes
##
##
## Time: 0.0 secs
```

Opgave 3.B

Se nederst 3.A

Opgave 4

```
sp_sex <- c()
for(i in 1:200){
    if(crabs[i,2]=="M"){
        if(crabs[i,1]=="B"){
            sp_sex <- c(sp_sex, "MB")
        }
        else{
            sp_sex <- c(sp_sex, "MO")
        }
    }
    if(crabs[i,2]=="F"){
        if(crabs[i,1]=="B"){
            sp_sex <- c(sp_sex, "FB")
        }
        else{</pre>
```

```
sp_sex <- c(sp_sex, "FO")
}
}
crabs2 <- data.frame(crabs, sp_sex)</pre>
```

De numeriske variable er:

```
crabs3 <- crabs2[4:9]
e <- C5.0(as.factor(sp_sex)~., data = crabs3)
plot(e)</pre>
```

