Multivariate Statistical Analysis

Homework 3

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Problem 1

```
swissbanknotes <- read.csv("Data_csv/SwissBankNotes.csv")</pre>
swissbanknotes$class <- factor(swissbanknotes$class)</pre>
banknote_type <- swissbanknotes$class</pre>
(a)
swissbanknotes2 <- scale(swissbanknotes[, -7])</pre>
summary(swissbanknotes2)
##
        length
                        height_left
                                                             inner_frame_lower
                                           height_right
##
  Min.
           :-2.91060
                       Min.
                              :-3.1064
                                          Min.
                                                 :-2.3672
                                                             Min.
                                                                    :-1.5350
   1st Qu.:-0.78608
                       1st Qu.:-0.6135
                                          1st Qu.:-0.6348
                                                             1st Qu.:-0.8428
## Median : 0.01062
                       Median : 0.2174
                                          Median : 0.1077
                                                             Median :-0.2198
                               : 0.0000
                                                 : 0.0000
## Mean
          : 0.00000
                       Mean
                                          Mean
                                                             Mean
                                                                   : 0.0000
##
  3rd Qu.: 0.54175
                       3rd Qu.: 0.7714
                                          3rd Qu.: 0.6645
                                                             3rd Qu.: 0.8186
## Max.
           : 3.72855
                               : 2.4333
                                          Max.
                                                 : 2.8299
                                                             Max.
                                                                  : 2.2723
##
  inner_frame_upper
                           diagonal
## Min.
           :-3.67459
                               :-2.32889
##
  1st Qu.:-0.68560
                       1st Qu.:-0.85354
  Median :-0.06289
                       Median :-0.02907
##
  Mean
          : 0.00000
                       Mean
                              : 0.00000
    3rd Qu.: 0.68435
                       3rd Qu.: 0.88218
  Max.
           : 2.05431
                               : 1.66324
                       Max.
pca_banknotes <- princomp(swissbanknotes2)</pre>
summary(pca_banknotes)
## Importance of components:
##
                              Comp.1
                                        Comp.2
                                                  Comp.3
                                                              Comp.4
                                                                         Comp.5
## Standard deviation
                           1.7119668 1.1276938 0.9298857 0.66896923 0.51704305
## Proportion of Variance 0.4909264 0.2130140 0.1448388 0.07496145 0.04477948
## Cumulative Proportion 0.4909264 0.7039403 0.8487791 0.92374054 0.96852002
##
                               Comp.6
## Standard deviation
                           0.43351526
## Proportion of Variance 0.03147998
## Cumulative Proportion 1.00000000
```

The cumulative proportion of the first two components is 0.704.

(b)

pca_banknotes\$loadings[, 1]

```
## length height_left height_right inner_frame_lower
## 0.006987029 -0.467758161 -0.486678705 -0.406758327
## inner_frame_upper diagonal
## -0.367891118 0.493458317
```

Component 1

The largest elements (in absolute value) correspond to height_left, height_right, diagonal length. So it's an index of height / diagonal length, where diagonal length has a negative sign. Observations with large negative component scores ksi_{i1} are shorter banknotes with longer diagonal whereas observations with large positive component scores ksi_{i1} are taller banknotes with a shorter diagonal.

pca_banknotes\$loadings[, 2]

```
## length height_left height_right inner_frame_lower
## 0.81549497 0.34196711 0.25245860 -0.26622878
## inner_frame_upper diagonal
## -0.09148667 0.27394074
```

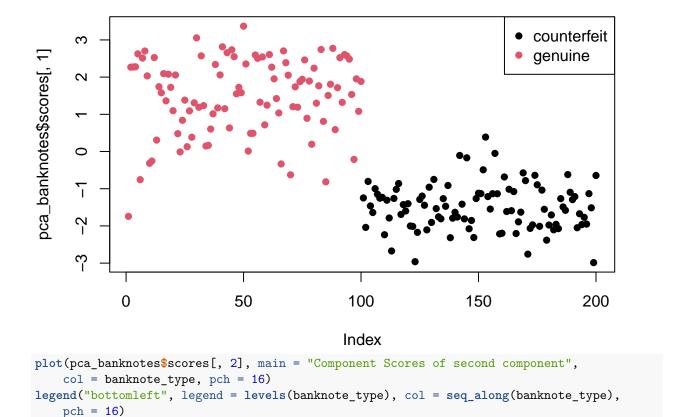
Component 2

The largest element (in absolute value) corresponds to length. So it's an index of banknote length. Observations with large negative component scores ksi_{i2} are shorter banknotes whereas Observations with large positive component scores ksi_{i2} are longer banknotes.

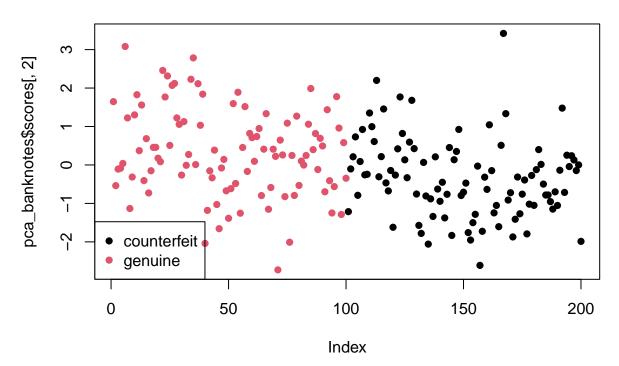
(c)

```
plot(pca_banknotes$scores[, 1], main = "Component Scores of first component",
        col = banknote_type, pch = 16)
legend("topright", legend = levels(banknote_type), col = seq_along(banknote_type),
        pch = 16)
```

Component Scores of first component



Component Scores of second component



The two groups for counterfeit banknotes and genuine banknotes are pretty well separated for both components.

Problem 2

```
turtles <- read.csv("Data_csv/turtles.csv")
turtle_sex <- factor(turtles$sex)
turtles <- turtles[-1]</pre>
```

(a)

Before applying log we check if there are any zero values.

The cumulative proportion of the first two components is 0.992.

(b)

```
pca_turtles$loadings[, 1]
## length width height
```

Component 1

0.5785513 0.5784099 0.5750829

The largest elements (in absolute value) correspond to length, width and height So it's a positive index of length, width and height. Observations with large negative component scores ksi_{i1} are turtles with smaller length, width and height whereas Observations with large positive component scores ksi_{i1} are turtles with larger length, width and height.

```
pca_turtles$loadings[, 2]

## length width height

## 0.3936563 0.4195001 -0.8179575
```

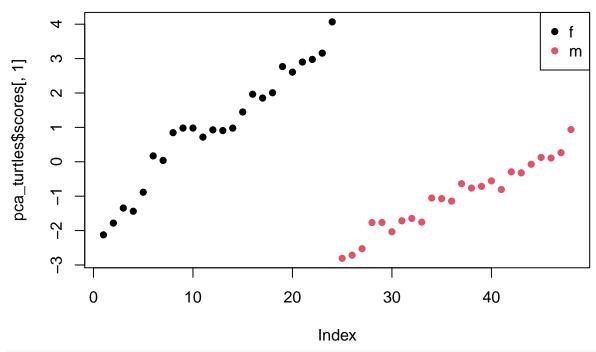
Component 2

The largest element (in absolute value) corresponds to height. So it's a negative index of turtle height. Observations with large negative component scores ksi_{i2} are taller turtles whereas observations with large positive component scores ksi_{i2} are shorter turtles.

(c)

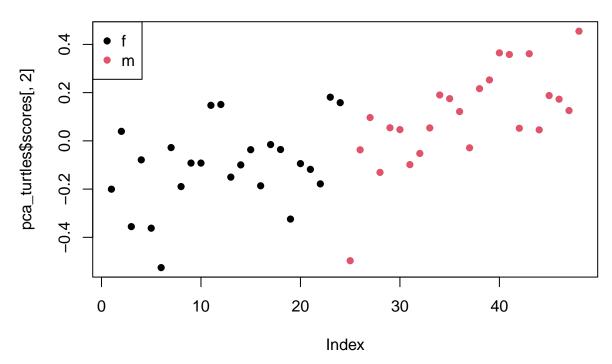
```
legend("topright", legend = levels(turtle_sex), col = seq_along(turtle_sex),
    pch = 16)
```

Component Scores of first component



```
plot(pca_turtles$scores[, 2], main = "Component Scores of second component",
        col = turtle_sex, pch = 16)
legend("topleft", legend = levels(turtle_sex), col = seq_along(turtle_sex),
        pch = 16)
```

Component Scores of second component



The two groups for female turtles and male turtles are pretty well separated for both components.