Multivariate Statistical Analysis

Homework 3

Lucas Fellmeth, Helen Kafka, Sven Bergmann

02/28/24

Problem 1

```
swissbanknotes <- read.csv("../Data_csv/SwissBankNotes.csv")
swissbanknotes$class <- factor(swissbanknotes$class)
banknote_type <- swissbanknotes$class</pre>
```

(a)

```
swissbanknotes2 <- scale(swissbanknotes[, -7])
summary(swissbanknotes2)</pre>
```

```
##
        length
                        height_left
                                          height_right
                                                            inner_frame_lower
           :-2.91060
                              :-3.1064
                                                 :-2.3672
                                                                   :-1.5350
##
   Min.
                       Min.
                                         Min.
   1st Qu.:-0.78608
                       1st Qu.:-0.6135
                                         1st Qu.:-0.6348
                                                            1st Qu.:-0.8428
  Median : 0.01062
                       Median : 0.2174
                                                            Median :-0.2198
                                         Median : 0.1077
   Mean
          : 0.00000
                       Mean
                              : 0.0000
                                         Mean
                                                 : 0.0000
                                                            Mean
                                                                   : 0.0000
   3rd Qu.: 0.54175
                       3rd Qu.: 0.7714
                                                            3rd Qu.: 0.8186
##
                                          3rd Qu.: 0.6645
## Max.
           : 3.72855
                       Max.
                              : 2.4333
                                                 : 2.8299
                                                            Max.
                                                                   : 2.2723
                                         Max.
##
   inner_frame_upper
                          diagonal
## Min.
          :-3.67459
                              :-2.32889
                       Min.
  1st Qu.:-0.68560
                       1st Qu.:-0.85354
## Median :-0.06289
                       Median :-0.02907
          : 0.00000
                              : 0.00000
   3rd Qu.: 0.68435
##
                       3rd Qu.: 0.88218
   Max.
           : 2.05431
                       Max.
                              : 1.66324
```

```
pca_banknotes <- princomp(swissbanknotes2)
summary(pca_banknotes)</pre>
```

```
## 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
```

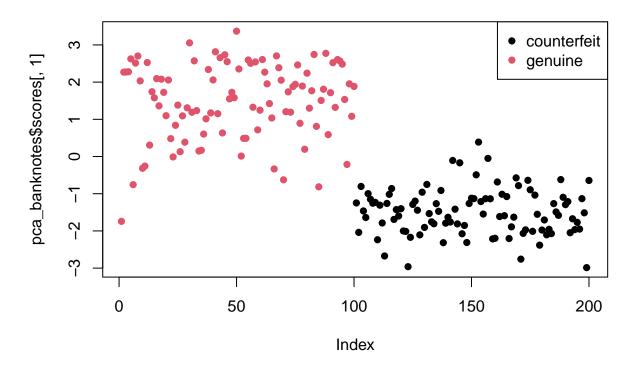
Component 2

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)

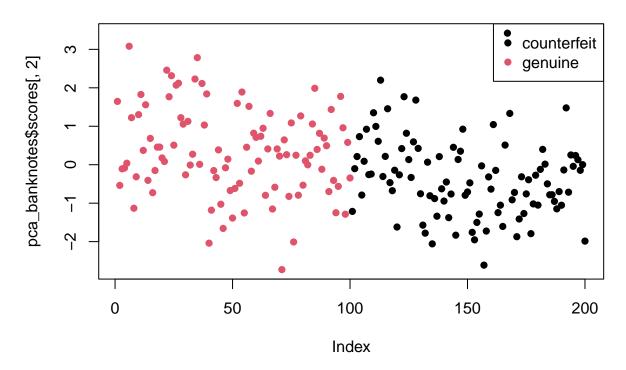
```
plot(pca_banknotes$scores[, 1], main = "Component Scores of first component",
      col = banknote_type, pch = 16)
legend("topright", legend = levels(banknote_type), col = seq_along(banknote_type),
      pch = 16)
```

Component Scores of first component



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

Component Scores of second component



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

Problem 2

summary(pca_turtles)

```
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.

```
which(turtles == 0)

## integer(0)

turtles2 <- log(turtles)
turtles2 <- scale(turtles2)
pca_turtles <- princomp(turtles2)</pre>
```

```
## Importance of components:

## Comp.1 Comp.2 Comp.3

## Standard deviation 1.6942473 0.21117868 0.149765480

## Proportion of Variance 0.9771826 0.01518177 0.007635642

## Cumulative Proportion 0.9771826 0.99236436 1.000000000
```

The cumulative proportion of the first two components is 0.992.

(b)

```
pca_turtles$loadings[, 1]

## length width height
## 0.5785513 0.5784099 0.5750829
```

Component 1

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
```

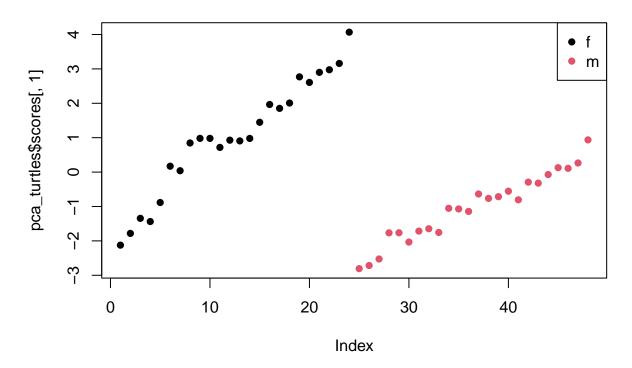
Component 2

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)

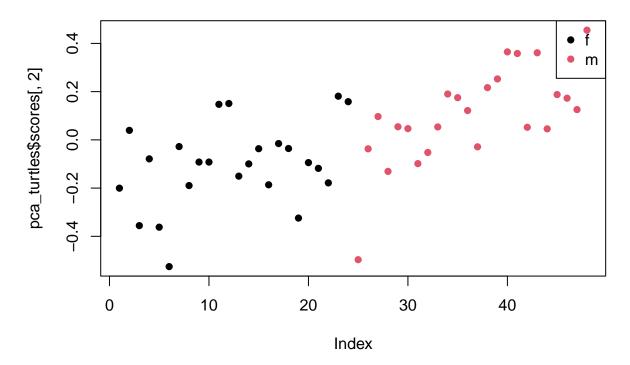
```
plot(pca_turtles$scores[, 1], main = "Component Scores of first component",
        col = turtle_sex, pch = 16)
legend("topright", legend = levels(turtle_sex), col = seq_along(turtle_sex),
        pch = 16)
```

Component Scores of first component



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