P8139 HW3

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3/28/2022

```
Load packages
```

```
library(tidyverse)
library(knitr)
```

Import data

```
data = read.table(file="PCA_data.csv", header=TRUE, row.names=1, sep=",")
```

PCA fitting and exploration

```
data_pca = prcomp(data, retx=TRUE, center=TRUE, scale=TRUE)
scores = data_pca$x
```

a) Calculate score using only loadings and data, and compare results with scores

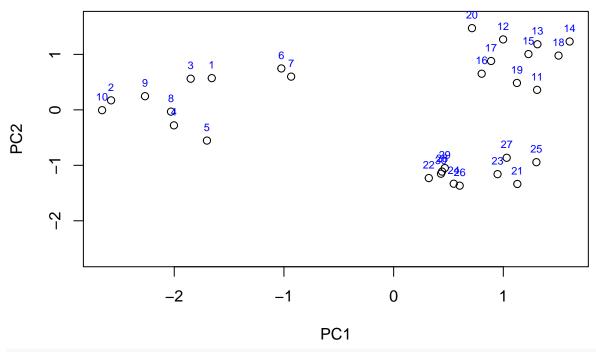
```
# extract loadings from the data_pca object
loadings = data_pca$rotation
# compute mean math, biology and chemistry grades
means =
  data %>%
  summarize(math = mean(math),
            bio = mean(bio),
            chem = mean(chem))
# compute math, biology and chemistry grade variances
vars =
  data %>%
  summarize(math = var(math),
            bio = var(bio),
            chem = var(chem))
# compute the standardized math, biology and chemistry grades
# by subtracting them with their means and dividing by
# their variances
std_data =
 data %>%
 mutate(
   math = (math - means$math)/sqrt(vars$math),
   bio = (bio - means$bio)/sqrt(vars$bio),
   chem = (chem - means$chem)/sqrt(vars$chem)
  ) %>%
  as.matrix()
```

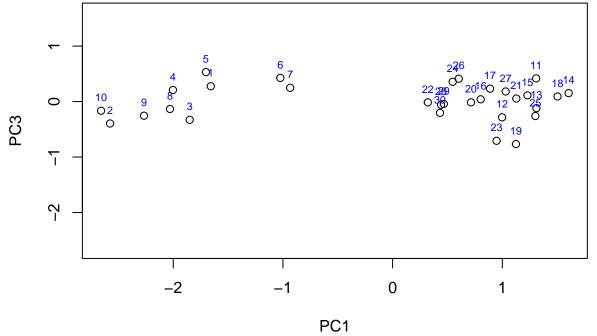
```
# multiply the matrices to obtain the PCs values for each observation
calc_scores = std_data %*% loadings
```

b) Percent Variance Explained by each PC

PC1	PC2	PC3
0.6692802	0.2971136	0.0336061

c) PCA Plots





```
# iii) PC2 vs PC3
plot(scores[,2:3],
     xlim=c(min(scores[,2:3]),
```

```
max(scores[,2:3])),
ylim=c(min(scores[,2:3]),
max(scores[,2:3])))
text(scores[,2], scores[,3], rownames(scores), col="blue", cex=0.7, pos=3)
```

