

Homework 7

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This homework is due on April 12, 2021 at 11:00pm. Please submit as a pdf file on Canvas.

For all problems in this homework, we will work with the `penguins_clean` dataset, which is a cleaned-up version of the `penguins` dataset from the `palmerpenguins` package.

Note: This homework is about the contents of the plots. Don't worry about styling. It's OK to use the default theme and plot labeling.

```
library(palmerpenguins)

penguins_clean <- penguins %>%
  select(-year) %>% # remove the year column as it is distracting here
  na.omit()          # remove any rows with missing values

penguins_clean

## # A tibble: 333 x 7
##   species island bill_length_mm bill_depth_mm flipper_length_~ body_mass_g
##   <fct>   <fct>         <dbl>         <dbl>         <int>         <int>
## 1 Adelie  Torge~         39.1          18.7          181          3750
## 2 Adelie  Torge~         39.5          17.4          186          3800
## 3 Adelie  Torge~         40.3           18           195          3250
## 4 Adelie  Torge~         36.7          19.3          193          3450
## 5 Adelie  Torge~         39.3          20.6          190          3650
## 6 Adelie  Torge~         38.9          17.8          181          3625
## 7 Adelie  Torge~         39.2          19.6          195          4675
## 8 Adelie  Torge~         41.1          17.6          182          3200
## 9 Adelie  Torge~         38.6          21.2          191          3800
## 10 Adelie Torge~         34.6          21.1          198          4400
## # ... with 323 more rows, and 1 more variable: sex <fct>
```

Problem 1: (2 pts)

Perform a PCA of the `penguins_clean` dataset and make two plots: 1. A rotation plot of components 1 and 2; 2. A plot of the eigenvalues, showing the amount of variance explained by the various components.

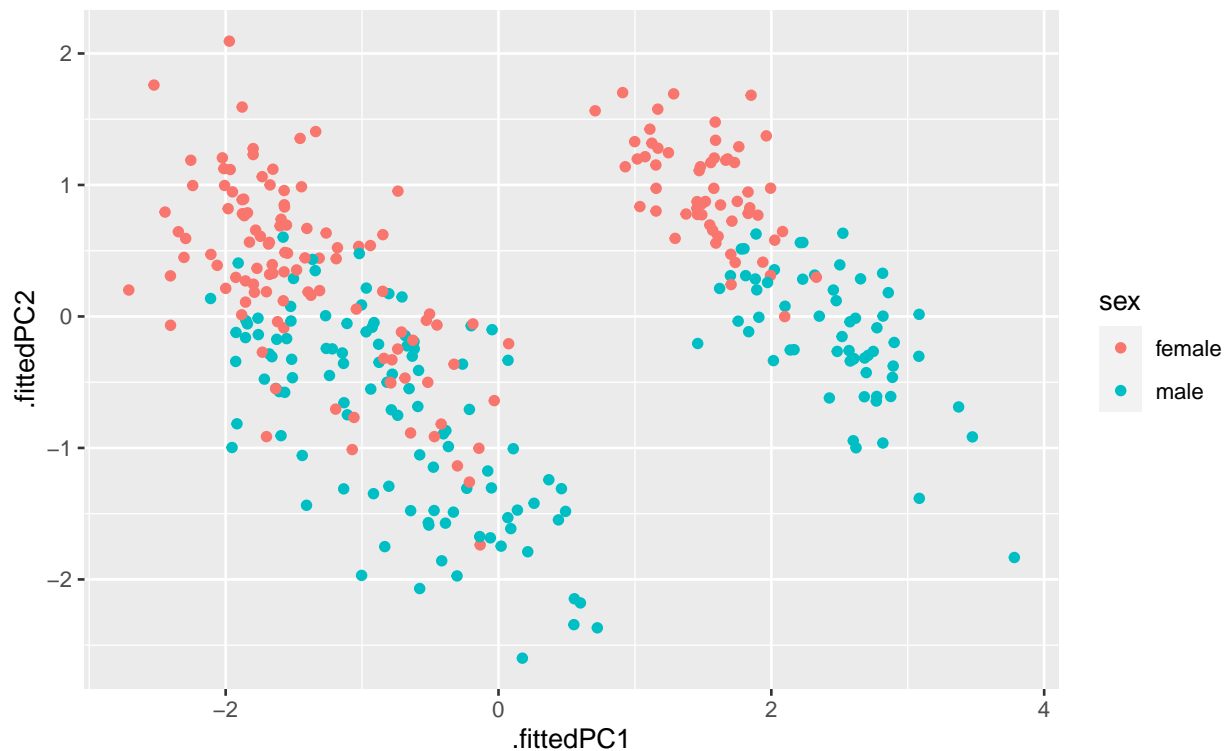
```
# your code goes here
#Plot1.
pca_fit <- penguins_clean %>%
  select(where(is.numeric)) %>%
  scale() %>%
  prcomp()

pca_fit

## Standard deviations (1, ..., p=4):
## [1] 1.6569115 0.8821095 0.6071594 0.3284579
##
```

```
## Rotation (n x k) = (4 x 4):
##           PC1      PC2      PC3      PC4
## bill_length_mm  0.4537532 -0.60019490 -0.6424951  0.1451695
## bill_depth_mm  -0.3990472 -0.79616951  0.4258004 -0.1599044
## flipper_length_mm 0.5768250 -0.00578817  0.2360952 -0.7819837
## body_mass_g      0.5496747 -0.07646366  0.5917374  0.5846861
```

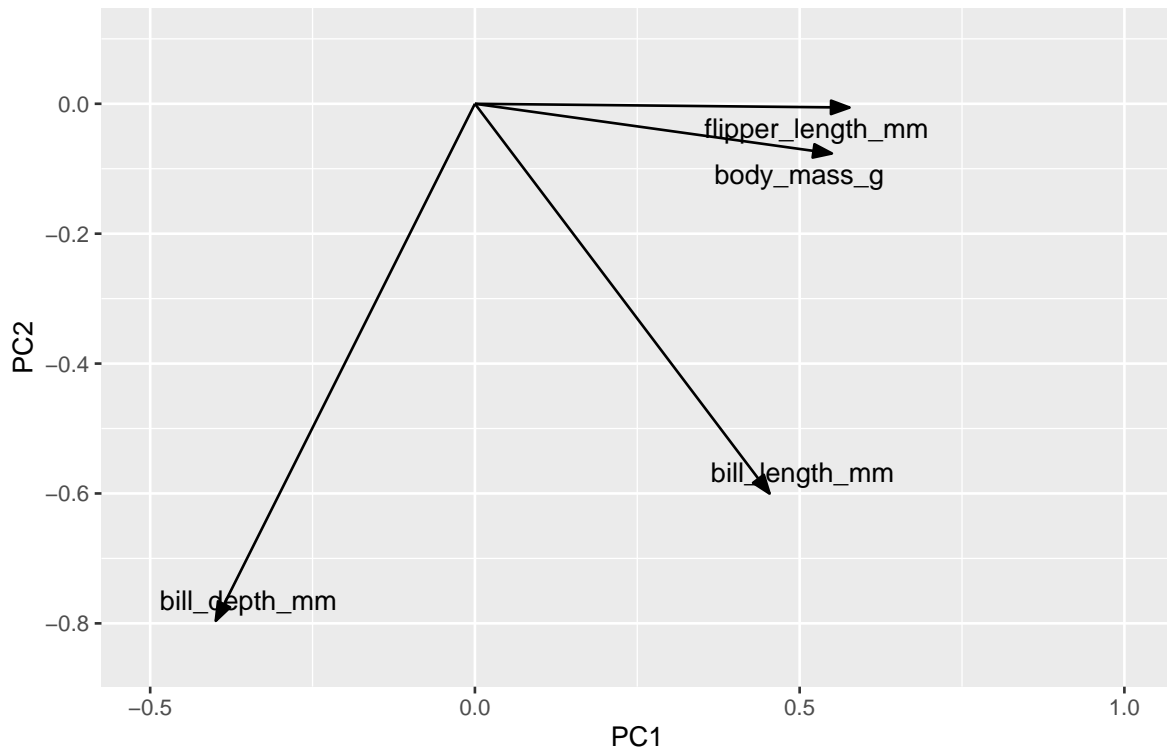
```
pca_fit %>%
  augment(penguins_clean) %>%
  ggplot(aes(.fittedPC1, .fittedPC2)) +
  geom_point(aes(color = sex))
```



```
arrow_style <- arrow(
  angle = 20, length = grid::unit(8, "pt"),
  ends = "first", type = "closed"
)

pca_fit %>%
  tidy(matrix = "rotation") %>% # extract rotation matrix
  pivot_wider(
    names_from = "PC", values_from = "value",
    names_prefix = "PC"
  ) %>%
  ggplot(aes(PC1, PC2)) +
  geom_segment(
    xend = 0, yend = 0,
    arrow = arrow_style
  ) +
  geom_text_repel(aes(label = column)) +
```

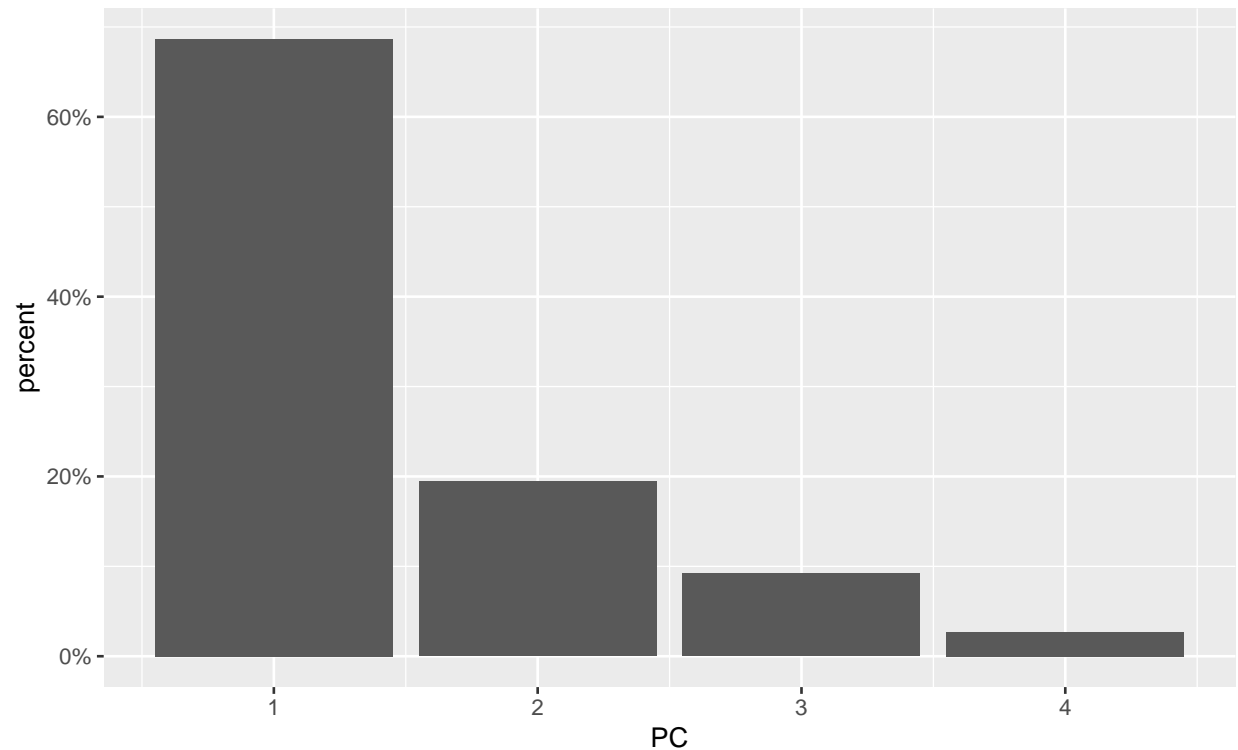
```
xlim(-0.50, 1) + ylim(-0.85, .1) +
coord_fixed()
```



```
#Plot2.
pca_fit %>%
  tidy(matrix = "eigenvalues")
```

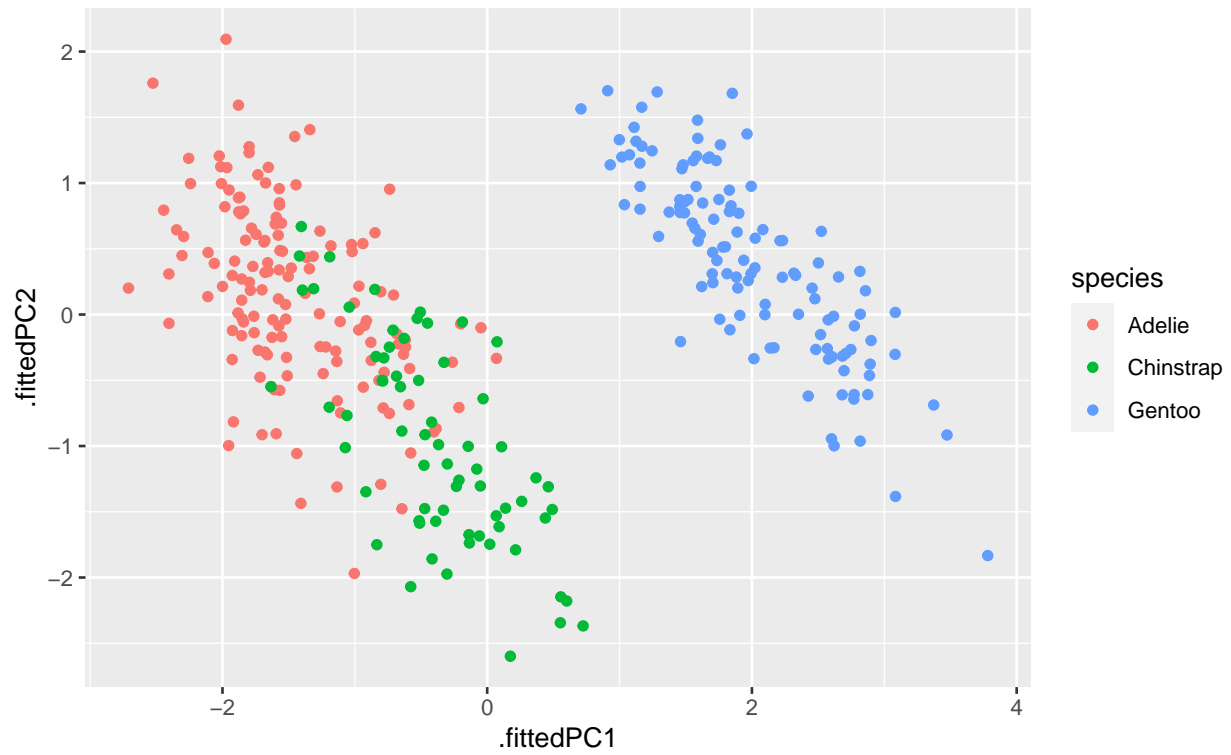
```
## # A tibble: 4 x 4
##   PC std.dev percent cumulative
##   <dbl> <dbl> <dbl>      <dbl>
## 1     1  1.66  0.686      0.686
## 2     2  0.882  0.195      0.881
## 3     3  0.607  0.0922     0.973
## 4     4  0.328  0.0270     1
```

```
pca_fit %>%
  tidy(matrix = "eigenvalues") %>%
  ggplot(aes(PC, percent)) +
  geom_col() +
  scale_x_continuous(breaks = 1:4) +
  scale_y_continuous(labels = scales::label_percent())
```



Problem 2: (4 pts) Make a scatter plot of PC 2 versus PC 1 and color by penguin species. Then use the rotation plot from Problem 1 to describe the physical characteristics by which the different penguin species differ. Finally, make one more scatter plot of the raw data that can support your interpretation of the PC analysis.

```
# your code goes here
pca_fit %>%
  augment(penguins_clean) %>%
  ggplot(aes(.fittedPC1, .fittedPC2)) +
  geom_point(aes(color = species))
```



```
#Plot2.
ggplot(penguins_clean, aes(body_mass_g, bill_length_mm, color = species)) +
  geom_point()
```

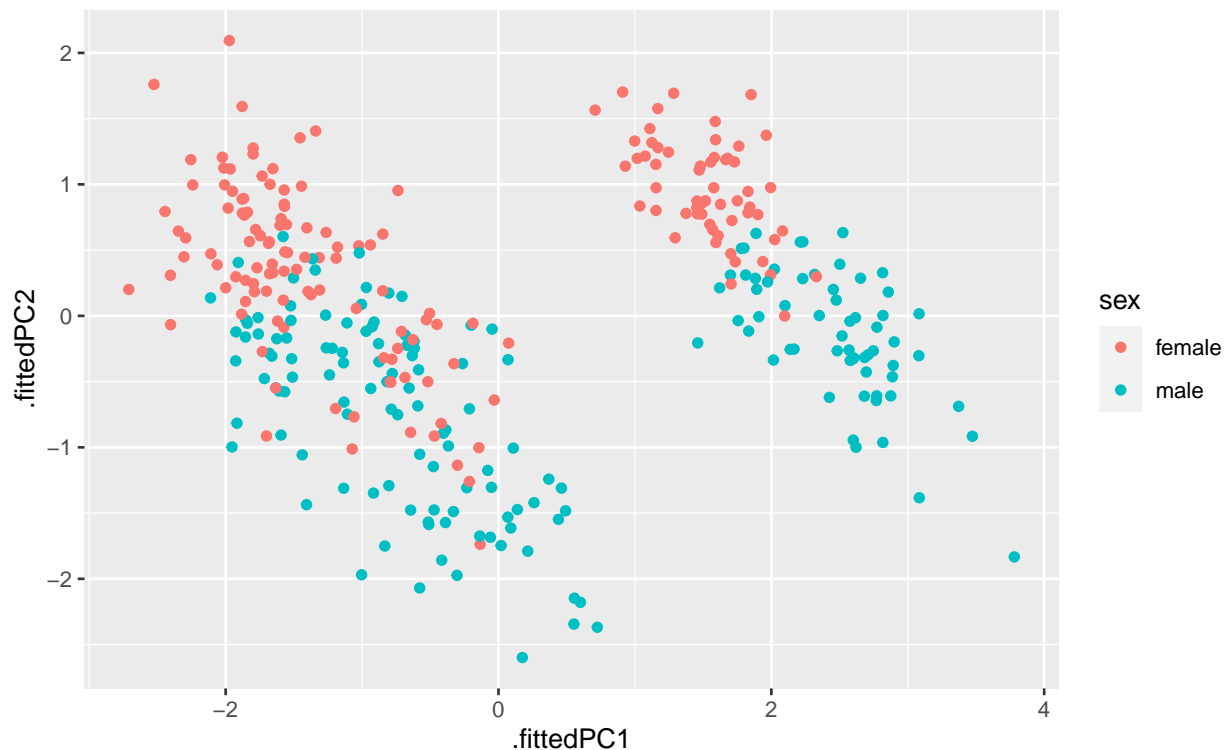


The rotation matrix shows that `flipper_length`, `body_mass`, and `bill_length` contribute positively to PC1 while `bill_depth` contributes negatively to PC1. Both `bill_depth` and `bill_length` contribute negatively to PC2. Considering this information, PC1 represents overall size of penguins while PC2 represents the size of beak. From the rotation matrix and the scatter plot for PC2 against PC1, this shows penguins with high body mass have relatively long bill lengths. Also, the scatter plot (PC2 against PC1) shows that Gentoo has relatively bigger body size compared to Chinstrap and Adelie. Chinstrap has bigger body size than Adelie. In terms of bill length, Adelie has shorter bill length than Chinstrap and Gentoo. This also found in the scatter plot using the raw data of bill length against body mass. In terms of body mass, this plot shows: Adelie < Chinstrap < Gentoo. Also, this plot shows that penguins with large body mass relatively have longer bill lengths.

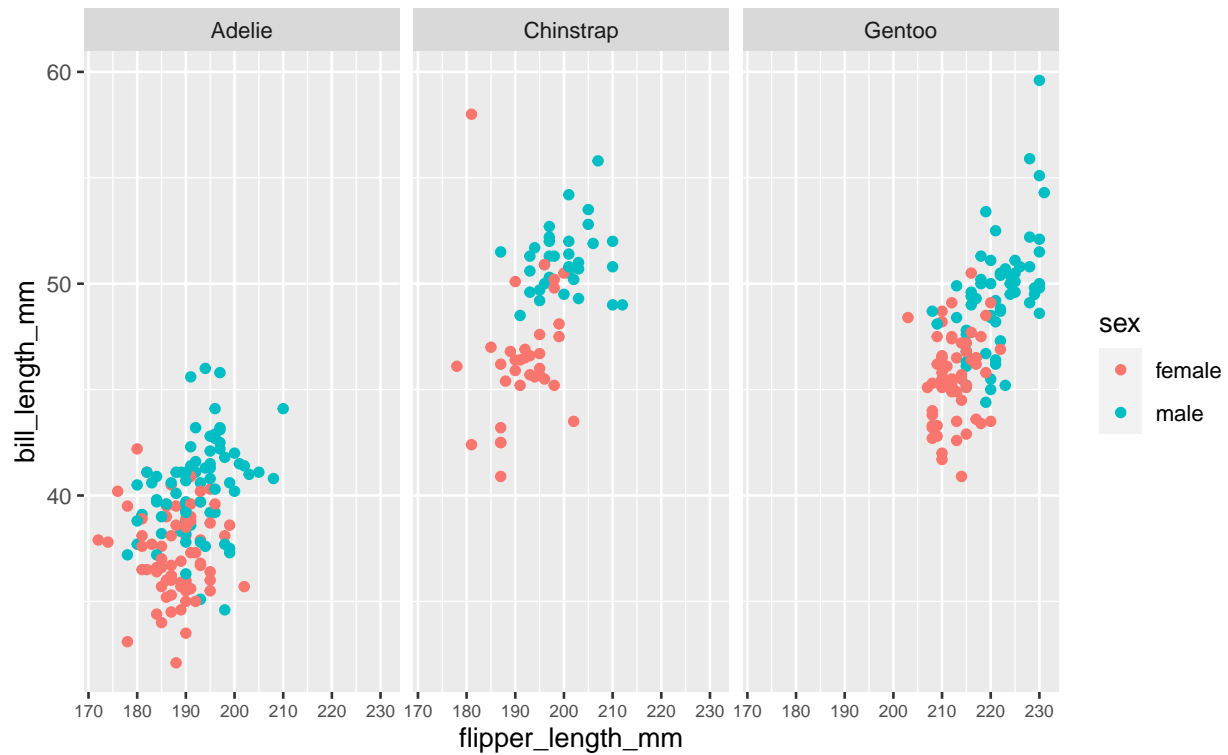
Problem 3: (4 pts) Again make a scatter plot of PC 2 versus PC 1, but now color by sex. Then use the rotation plot from Problem 1 to describe the physical characteristics by which the different penguin sexes differ. Finally, make one more scatter plot of the raw data that can support your interpretation of the PC analysis.

Hint: It helps to facet by penguin species.

```
# your code goes here
#Plot1.
pca_fit %>%
  augment(penguins_clean) %>%
  ggplot(aes(.fittedPC1, .fittedPC2)) +
  geom_point(aes(color = sex))
```



```
#Plot2.
ggplot(penguins_clean, aes(flipper_length_mm, bill_length_mm, color = sex)) +
  geom_point() +
  theme(axis.text.x = element_text(size = 7)) +
  facet_wrap(~species)
```



The rotation matrix shows that the *flipper_length* contributes on *PC1*; the *bill length* contributes on both *PC1* and *PC2*. The scatter plot of *PC 2* against *PC 1* by sex shows male penguins have bigger body size, flipper length, and bill length than female penguins when we look at the different scatter group. The scatter plot using the raw data for bill length against flipper length by sex shows male penguins are bigger than female penguins for flipper length and bill length. In terms of species, Adelie has the shortest lengths for both flipper and bill. Chinstrap and Gentoo have similar range for bill length but Gentoo has longer flipper than Chinstrap.