Stat 123 Summer 2023 Midterm 1b

2023-05-31

All answers are to be done with R commands.

Question 1

Download the data file artists.csv from Brightspace and save it in a directory of your choice.

Read the data file as a data frame and name it artist.

```
#a. Find the sum of the artists' height and display the result below.
artists <- read.csv('artists.csv')
head(artists)</pre>
```

```
##
        Name Height Weight Eye_Colour Martial_Arts
## 1
         Tom
              1.83
                     75.3
                                  1
## 2
              1.72 70.8
                                  3
                                              1
        Josh
## 3 Michelle
              1.55 62.5
                                  2
                                              1
                                              0
## 4
              1.76 71.2
       Peter
                    60.8
                                              1
## 5
       Emily
              1.50
```

```
#b. Find the average of the artists' weight and display the result below.
artists_weight <- artists[, "Weight"]
avg_artsits_weigth <- mean(artists_weight)
avg_artsits_weigth</pre>
```

```
## [1] 68.12
```

Question 2

You will use the built-in data chickwts and the dplyr package for this question.

```
#a. Load the dplyr package using a R command.

library(dplyr)
```

```
##
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
##
## filter, lag
```

```
## The following objects are masked from 'package:base':
##
## intersect, setdiff, setequal, union
```

```
#b. Display the first 8 observations (rows) in the dataset.
chick <- chickwts
chickwts[1:8, ]</pre>
```

```
weight
                 feed
## 2
        160 horsebean
## 3
        136 horsebean
## 4
        227 horsebean
## 5
        217 horsebean
## 6
        168 horsebean
## 7
        108 horsebean
## 8
        124 horsebean
```

```
#. For the following questions, you MUST use dplyr piping commands to complete the tasks.

#c. Find out how many chickens receive the meatmeal feed.
meal <- chickwts['feed'] %>%
  filter(feed == 'meatmeal')

length(meal[, 'feed'])
```

```
## [1] 11
```

#d. Create a new data frame from chickwts named chickwt2 that contains chickens with
weight more than 150. Do not show the results.
chickwt2 <- chickwts['weight'] %>% filter(weight > 150)
length(chickwt2[,'weight'])

[1] 64

#e. Display the chickwt2 data frame in ascending order by weight.
chickwt2_sorted <- chickwt2 %>%
 arrange(weight)

chickwt2_sorted

```
##
      weight
## 1
         153
## 2
         158
## 3
         160
## 4
         168
## 5
         169
## 6
         171
## 7
         179
## 8
         181
## 9
         193
## 10
         199
## 11
         203
## 12
         206
## 13
         213
## 14
         216
## 15
         217
## 16
         222
## 17
         226
## 18
         227
## 19
         229
## 20
         230
## 21
         242
## 22
         243
## 23
         244
## 24
         248
## 25
         248
## 26
         250
## 27
         257
## 28
         257
## 29
         258
## 30
         260
## 31
         260
## 32
         263
## 33
         267
## 34
         271
## 35
         271
## 36
         283
## 37
         295
## 38
         297
## 39
         303
## 40
         309
## 41
         315
## 42
         316
## 43
         318
## 44
         318
## 45
         320
## 46
         322
## 47
         325
## 48
         327
## 49
         329
## 50
         332
## 51
         334
## 52
         339
## 53
         340
## 54
         341
## 55
         344
## 56
         352
## 57
         359
## 58
         368
## 59
         379
## 60
         380
## 61
         390
## 62
         392
## 63
         4\,0\,4
## 64
         423
#f. Display the mean weight of chickens grouped by their feed type.
```

```
#f. Display the mean weight of chickens grouped by their feed type.
group <- chickwts %>%
  group_by(feed) %>%
  summarize(mean_weight = mean(weight))
group
```

```
## # A tibble: 6 × 2
            mean_weight
    feed
##
    <fct>
                    <dbl>
## 1 casein
                     324.
## 2 horsebean
                     160.
## 3 linseed
                     219.
## 4 meatmeal
                     277.
## 5 soybean
                     246.
## 6 sunflower
                     329.
```

Once you are satisfied with your results, knit this file as a html and submit it to Brightspace. If you are running out of time and the file does not knit, add the character # at the beginning of the commands that did not run. Then knit it as a html file and submit.