PHW251 Problem Set 4

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For this problem set you will tidy up a dataset of 500 individuals. We also want to calculate each individual's BMI and appropriately categorize them.

Load your data (500_Person_Gender_Height_Weight.csv):

Question 1

Clean the column headers to be all lower case, have no spaces, and rename "Location information" to location.

```
data <- data %>%
  rename(
    location = `Location information`,
    gender = GENDER
)

colnames(data)
```

[1] "location" "gender" "height" "weight"

Question 2

Create a new variable that calculates BMI for each individual.

You will need to navigate the different system of measurements (metric vs imperial). Only the United States is using imperial.

• BMI calculation and conversions:

```
- metric: BMI = weight(kg)/[height(m)]^2

- imperial: BMI = 703 * weight(lbs)/[height(in)]^2

- 1 foot = 12 inches

- 1 cm = 0.01 meter
```

Although there's many ways you can accomplish this task, we want you to use an if_else() to calculate BMI with the appropriate formula based on each person's location.

```
# head(data)

data <- data %>%
  mutate(
    bmi = if_else(
        location %in% c("United Kingdom","Taiwan"),
        weight/((height*0.01)^2),
        703 * weight/((height*12)^2)
    )
    )
head(data)
```

```
## # A tibble: 6 x 5
##
                     gender height weight
     location
                                             bmi
##
     <chr>>
                     <chr>>
                             <dbl>
                                     <dbl> <dbl>
## 1 New York
                     Male
                              5.71
                                      212.
                                            31.7
## 2 United Kingdom Male
                            189
                                       87
                                            24.4
## 3 New York
                     Female
                              6.07
                                      243.
                                            32.1
## 4 Taiwan
                     Female 195
                                      104
                                            27.4
## 5 Taiwan
                                            27.5
                     Male
                            149
                                       61
## 6 Taiwan
                     Male
                            189
                                      104
                                            29.1
```

Question 3

Create a new variable that categorizes BMI with case_when():

• Underweight: BMI below 18.5

Normal: 18.5-24.9Overweight: 25.0-29.9Obese: 30.0 and Above

```
data <- data %>% mutate(
    bmi_cat =
        case_when(
        bmi < 18.5 ~ "Underweight",
        bmi <= 24.9 ~ "Normal",
        bmi <= 29.9 ~ "Overweight",
        TRUE ~ "Obese"
    )
)
head(data)</pre>
```

```
## # A tibble: 6 x 6
##
     location
                    gender height weight
                                            bmi bmi_cat
     <chr>
                            <dbl> <dbl> <dbl> <chr>
##
                    <chr>>
## 1 New York
                    Male
                             5.71
                                     212. 31.7 Obese
## 2 United Kingdom Male
                           189
                                      87
                                           24.4 Normal
## 3 New York
                             6.07
                                     243.
                                           32.1 Obese
                    Female
## 4 Taiwan
                    Female 195
                                     104
                                           27.4 Overweight
## 5 Taiwan
                    Male
                           149
                                           27.5 Overweight
                                      61
## 6 Taiwan
                    Male
                           189
                                     104
                                           29.1 Overweight
```

Could we have used if_else()?

Yes, but we would have the specify the lower bound as well for each line in contrast with case_when(), where the lines are run sequentially and the first match in the function would be the output.

Question 4

Arrange your data first by location and then by descending order of BMI.

```
data <- data %>%
 arrange(location, desc(bmi))
head(data)
## # A tibble: 6 x 6
    location gender height weight
                                    bmi bmi_cat
##
    <chr>>
             <chr>
                     <dbl> <dbl> <dbl> <chr>
                      4.66
## 1 Colorado Female
                             351. 78.8 Obese
## 2 Colorado Female
                                   74.6 Obese
                      4.59
                             322.
## 3 Colorado Male
                      4.72
                             320.
                                   70.1 Obese
## 4 Colorado Female
                     4.95
                                   69.4 Obese
                             348.
## 5 Colorado Female 4.66
                             302.
                                   67.9 Obese
## 6 Colorado Male
                      4.95
                             340.
                                   67.7 Obese
```

Question 5

Use a dplyr method to remove the height, weight, and BMI columns from your data.

Optional Challenge

Perform all the actions in this problem set with one dpylr call.

```
data <- read_csv("~/phw251_fall24_cv/week 5/500_Person_Gender_Height_Weight.csv") %>%
  rename(
    location = `Location information`,
    gender = GENDER
  ) %>%
  mutate(
    bmi = if_else(
     location %in% c("United Kingdom", "Taiwan"),
      weight/((height*0.01)^2),
     703 * weight/((height*12)^2)
    )
  ) %>%
  mutate(
  bmi_cat =
    case_when(
     bmi < 18.5 ~ "Underweight",
     bmi <= 24.9 ~ "Normal",</pre>
     bmi <= 29.9 ~ "Overweight",
      TRUE ~ "Obese"
    )
) %>%
  arrange(location, desc(bmi)) %>%
  select(location, gender, bmi_cat)
## Rows: 500 Columns: 4
## -- Column specification -----
## Delimiter: ","
## chr (2): Location information, GENDER
## dbl (2): height, weight
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
head(data)
## # A tibble: 6 x 3
##
    location gender bmi_cat
             <chr> <chr>
     <chr>
## 1 Colorado Female Obese
## 2 Colorado Female Obese
## 3 Colorado Male Obese
## 4 Colorado Female Obese
## 5 Colorado Female Obese
## 6 Colorado Male Obese
```