Assignment-2

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Loading required packages

Used supressMessages with library as without it, I was getting this error while knitting: "! Package inputenc Error: Unicode char √ (U+221A) (inputenc) not set up for use with LaTeX."

```
suppressMessages(library(tidyverse))
suppressMessages(library(ggplot2))
surveys <- read.csv("surveys.csv", header = T, sep = ",")
View(surveys)
suppressMessages(library(gapminder))</pre>
```

1. Write R code to extract the survey observations for the first three months of 1990 using the filter() function. (5 points)

```
winter surveys 90 <- surveys %>% filter((year == "1990"), (month %in%
c("1","2","3")))
View(winter surveys 90)
head(winter surveys 90)
## record id month day year plot id species id sex hindfoot length weight
##1
      16879 1 6 1990
                         1
                               DM F
                                           37
                                                35
                               OL M
##2
      16880 1 6 1990
                         1
                                           21
                                                28
##3
     16881 1 6 1990
                         6
                               PF M
                                           16
                                                7
     16882 1 6 1990
## 4
                         23
                                RM F
                                            17
                                                 9
##5
                                            17
     16883 1 6 1990
                         12
                                RM M
                                                 10
## 6 16884 1 6 1990
                         24
                                RM M
                                            17
```

2. Sort the 1990 winter surveys data by descending order of record ID, then by ascending order of weight. (10 points)

Taking the winter surveys data as the data of the first three months

```
arrange_record <- arrange(winter_surveys_90, desc(record_id), weight)
View(arrange_record)
head(arrange_record)
```

```
## record id month day year plot id species id sex hindfoot length weight
      17369
                                DM F
## 1
              3 30 1990
                          8
                                            36
                                                 39
                                DM F
##2
      17368
              3 30 1990
                          11
                                             35
                                                 41
      17367 3 30 1990
##3
                          4
                                DM M
                                             37
                                                 44
##4
                          11
                                             37
      17366 3 30 1990
                                DM M
                                                46
##5
      17365
              3 30 1990
                          4
                                DM F
                                            38
                                                 48
                                                 51
## 6 17364 3 30 1990
                          8
                                DM M
                                            36
```

Note that the echo = FALSE parameter was added to the code chunk to prevent printing of the R code that generated the plot.

3. Write code that returns the record_id, sex and weight of all surveyed individuals of Reithrodontomys montanus (RO), (10 points)

```
RO data <- surveys %>% filter(species id == "RO") %>% select(species id,
record id, sex, weight)
head(RO_data)
## species id record id sex weight
## 1
         RO
              18871 F
                         11
##2
         RO
              33397 M
                          8
                         9
##3
         RO
              33556 M
## 4
              33565 F
                          8
         RO
##5
         RO
                         11
              34517 M
##6
         RO
              35402 F
                         12
```

4. Write code that returns the average weight and hindfoot length of Dipodomys merriami (DM) individuals observed in each month (irrespective of the year). Make sure to exclude NA values. (10 points)

```
DM <- filter(surveys, species id =="DM")
View(DM)
months DM <- group by(DM, month)
View(months DM)
month sum <- summarise(months DM, weight mean = mean(weight, na.rm
= T), hindfoot length mean = mean(hindfoot length, na.rm = T))
month sum
## # A tibble: 12 x 3
##
     month weight mean hindfoot length mean
                             <dbl>
## <int>
              <dbl>
## 1 1
             42.9
                           36.1
       2
## 2
             44.0
                           36.2
## 3
       3
             45.2
                           36.1
```

```
## 4 4
            44.8
                         36.2
       5
## 5
            43.2
                         35.8
## 6
       6
            41.5
                         36.0
## 7
     7
            41.9
                         35.7
## 8 8
            41.8
                         35.8
## 9
       9
            43.3
                         35.8
## 10 10
             42.5
                          36.0
## 11
       11
             42.4
                          35.9
## 12 12
             43.0
                          36.0
```

5. Write code that determines the number of individuals by species observed in the winter of 1990. (15 points)

Taking the winter surveys data as the data of the first three months

```
Groups <- summarize(group by(winter surveys 90, species id),
num individuals =n()
head(Groups)
## # A tibble: 6 x 2
## species id num individuals
## <fctr>
                   <int>
##1""
                    1
## 2 AB
                    25
##3 AH
                     4
                     3
## 4 BA
## 5 DM
                    132
## 6 DO
                    65
```

6. Create a dataframe named gapminder_df and mutate it to contain a column that contains the gross domestic product for each row in the data frame. (5 points)

```
gapminder df <- as.data.frame(gapminder)</pre>
View(gapminder df)
gapminder df <- mutate(gapminder df, GDP = pop*gdpPercap)</pre>
head(gapminder df)
##
      country continent year lifeExp
                                     pop gdpPercap
                                                       GDP
## 1 Afghanistan
                   Asia 1952 28.801 8425333 779.4453 6567086330
                   Asia 1957 30.332 9240934 820.8530 7585448670
## 2 Afghanistan
## 3 Afghanistan
                   Asia 1962 31.997 10267083 853.1007 8758855797
## 4 Afghanistan
                   Asia 1967 34.020 11537966 836.1971 9648014150
                   Asia 1972 36.088 13079460 739.9811 9678553274
## 5 Afghanistan
## 6 Afghanistan Asia 1977 38.438 14880372 786.1134 11697659231
```

7. Calculate the Mean GDP for Cambodia for the years within the dataset. (15 points)

```
summarize(gapminder_df, country = "Cambodia", mean_GDP = mean(GDP))
## country mean_GDP
## 1 Cambodia 186809560507
```

8. Find the year with the maximum life expectancy for countries in Asia and arrange them in descending order by year, The result should contain the country's name, the year and the life expectancy. (15 points)

```
gapminder df %>% filter(continent == "Asia") %>% group by(country) %>%
filter(lifeExp == max(lifeExp)) %>% select(country,year,lifeExp) %>%
arrange(desc(year))
## # A tibble: 33 x 3
## # Groups: country [33]
## country year lifeExp
## <fctr>
             <int> <dbl>
## 1 Afghanistan 2007 43.8
## 2 Bahrain
                 2007 75.6
## 3 Bangladesh 2007 64.1
## 4 Cambodia 2007 59.7
                 2007 73.0
## 5 China
## 6 Hong Kong, China 2007 82.2
## 7 India
                2007 64.7
## 8 Indonesia
                  2007 70.6
## 9 Iran
                2007 71.0
                2007 80.7
## 10 Israel
## # ... with 23 more rows
```

The year with the maximum life expectancy for countries in Asia is 2007 #9. Count the number of observations per continent. (5 points)

10. Compute the average and median life expectancy and GDP per capita by continent for the years 1952 and 2007. Should we be optimistic given the results? (10 points)

```
gapminder df %>% filter(year %in% c(1952,2007)) %>%
group by(continent, year) %>% summarize(mean lifeExp = mean(lifeExp),
med lifeExp = median(lifeExp), avg GDP = mean(gdpPercap), med GDP =
median(gdpPercap))
## # A tibble: 10 x 6
## # Groups: continent [?]
     continent year mean lifeExp med lifeExp avg GDP med GDP
##
     <fctr> <int>
                      <dbl>
                               <dbl> <dbl> <dbl>
## 1 Africa
             1952
                      39.1
                              38.8
                                    1253
                                           987
## 2 Africa
             2007
                      54.8
                              52.9
                                    3089
                                          1452
## 3 Americas 1952
                        53.3
                                54.7 4079
                                            3048
## 4 Americas 2007
                        73.6
                                72.9 11003 8948
## 5 Asia
                              44.9 5195
            1952
                      46.3
                                         1207
## 6 Asia
            2007
                              72.4 12473 4471
                      70.7
## 7 Europe
              1952
                       64.4
                               65.9 5661
                                            5142
                               78.6 25054 28054
## 8 Europe
              2007
                       77.6
## 9 Oceania 1952
                       69.3
                                69.3 10298 10298
## 10 Oceania 2007
                                80.7 29810 29810
                        80.7
```

We should be optimistic as the average and median life expectancy and GDP per capita by continent are increasing.