# DS5100\_HW11

Eashan Kaw 4/6/2023

# Metadata

Course: DS 5100

Module: 11 R Programming 2 Topic: HW on Tidyverse

Author: R.C. Alvarado (adapted)
Date: 07 October 2022 (revised)

## Student Info

Name: Eashan Kaw Net ID: elk7ed

File GitHub URL: https://github.com/elkaw/DS5100-2023-01-0/blob/main/lessons/M10 RBasics/M11-HW.ipynb

# Instructions

In your private course repo use this notebook to write code that performs the tasks below.

Save your notebook in the M11 directory.

Remember to add and commit these files to your repo.

Then push your commits to your repo on GitHib.

Be sure to fill out the Student Info block above.

To submit your homework, save your results as a PDF and upload it to GradeScope.

**TOTAL POINTS: 7** 

# Overview

In this homework, you will work with the Abalone dataset (https://archive.ics.uci.edu/ml/datasets/Abalone) from the UCI Machine Learning Repository.

To get started, download and import the abalone.data dataset from this URL:

https://archive.ics.uci.edu/ml/machine-learning-databases/abalone/abalone.data (https://archive.ics.uci.edu/ml/machine-learning-databases/abalone/abalone.data)

You can pass the URL directly to read.csv() and that there is no header row.

Note: The instruction to print in the questions below can be accomplished either through the print() function or by displaying a value directly.

**TOTAL POINTS: 7** 

# **Tasks**

#### Task 0

(0 points)

Get the dataset.

```
# CODE HERE
#install.packages("tinytex")
library(tinytex)
library(dplyr)
```

```
## Registered S3 methods overwritten by 'tibble':
## method from
## format.tbl pillar
## print.tbl pillar
```

```
##
## 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
```

```
abalone_df <- read.csv('/Users/eashankaw/Documents/Continuing Education/DS5100_programming_FOR_DATA_SCIENCE/abalo
ne.data', header = F)
head(abalone_df)</pre>
```

```
## V1 V2 V3 V4 V5 V6 V7 V8 V9
## 1 M 0.455 0.365 0.095 0.5140 0.2245 0.1010 0.150 15
## 2 M 0.350 0.265 0.090 0.2255 0.0995 0.0485 0.070 7
## 3 F 0.530 0.420 0.135 0.6770 0.2565 0.1415 0.210 9
## 4 M 0.440 0.365 0.125 0.5160 0.2155 0.1140 0.155 10
## 5 I 0.330 0.255 0.080 0.2050 0.0895 0.0395 0.055 7
## 6 I 0.425 0.300 0.095 0.3515 0.1410 0.0775 0.120 8
```

#### Task 1

(1 point)

Print the number of rows in the dataset.

```
# CODE HERE
print(nrow(abalone_df))
```

```
## [1] 4177
```

#### Task 2

(1 point)

The rightmost column is the number of rings. Print the maximum number of rings

```
# CODE HERE
print(abalone_df %>% select(V9) %>% max)
```

```
## [1] 29
```

## Task 3

(1 point)

The leftmost column is the gender with these values: M: male, F: female, I: infant.

Apply the filter() function from tidyverse to select only rows where gender is infant, and print the number of records.

```
# CODE HERE
#print(abalone_df$M %>% unique)
print(nrow(abalone_df %>% filter(abalone_df$V1 == "I")))
```

```
## [1] 1342
```

# Task 4

(1 point)

Apply the filter() function from tidyverse to select only rows where gender is infant or male, and print the number of records.

```
# CODE HERE
print(nrow(abalone_df %>% filter(abalone_df$V1 == "I" | abalone_df$V1 == "M")))
```

```
## [1] 2870
```

# Task 5

(1 point)

Call the table() function on the abalone genders to find out how many of each gender are present.

Print the result.

```
# CODE HERE
print(abalone_df$V1 %>% table)
```

```
## .
## F I M
## 1307 1342 1528
```

## Task 6

(1 point)

Compute the mean value of column 2 (V2) grouped by gender.

V2 is the longest shell measurement.

Requirements: use the %>% operator to chain commands, and the group\_by() and summarize() functions.

```
# CODE HERE
abalone_df %>% group_by(abalone_df$V1) %>% summarize(mean(abalone_df$V2))
```

```
## Warning in pillar::colonnade(df, has_row_id = if (star) "*" else TRUE,
## needs_dots = needs_dots): `...` must be empty.
```

# Task 7

(1 point)

Compute the MEDIAN value of longest shell measurement for only the males.

Requirements: use the %>% operator to chain commands.

```
# CODE HERE
print(abalone_df %>% filter(abalone_df$V1 == 'M') %>% select(V2) %>% summarize(median(abalone_df$V2)))
```

```
## median(abalone_df$V2)
## 1 0.545
```