# PSTAT 131 Homework 2

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## Linear Regression

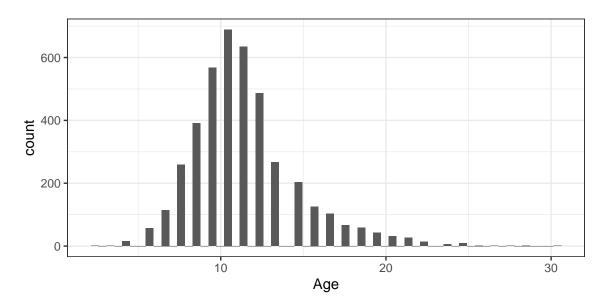
## Question 1:

Predicting abalone age by computing number of rings + 1.5 and adding the age variable into the data set. I renamed the variable Class Number of Rings to Rings for simplicity.

```
abalone <- rename(abalone, Rings = Class_number_of_rings)
abalone <- mutate(abalone, Age = Rings + 1.5)
```

Here, we will assess and describe the distribution of Age.

```
abalone %>%
  ggplot(aes(x = Age)) +
  geom_histogram(bins = 60) +
  theme_bw()
```



From the histogram above, we see that is a spike at Age of 10 and it is skewed right.

#### Question 2:

Splitting the abalone data into a training set and a testing set using stratified sampling with a proportion of 80%.

#### Question 3:

Using the **training** data, I create a recipe predicting the outcome variable, age, with all other predictor variables. \*We do not include rings to predict age because the variable age stemmed from rings, and is only rings + 1.5.

Steps for the recipe:

- 1. dummy code any categorical predictors
- 2. create interactions between
  - type and shucked\_weight
  - longest shell and diameter
  - · shucked weight and shell weight
- 3. center all predictors,
- 4. scale all predictors.

### Question 4:

Creating and storing a linear regression object using the 1m engine.

```
lm_model <- linear_reg() %>%
set_engine("lm")
```

#### Question 5:

Now, we create a workflow by 1. setting up an empty workflow,

- 2. adding the model created in Question 4, and
- 3. adding the recipe created in Question 3

```
lm_wflow <- workflow() %>%
  add_model(lm_model) %>%
  add_recipe(abalone_recipe)
```

#### Question 6:

Using fit() object to predict the age of a hypothetical female abalone with length = 0.50, diameter = 0.10, height = 0.30,  $whole_weight = 4$ ,  $shucked_weight = 1$ ,  $viscera_weight = 2$ ,  $shell_weight = 1$ .

```
# fitting the linear model
lm_fit <- fit(lm_wflow, abalone_train)

lm_fit %>%
    extract_fit_parsnip() %>%
    tidy()
```

```
## # A tibble: 14 x 5
##
      term
                                    estimate std.error statistic p.value
##
      <chr>
                                       <dbl>
                                                 <dbl>
                                                           <dbl>
                                                                     <dbl>
##
   1 (Intercept)
                                      11.5
                                                0.0376
                                                           304.
                                                                 0
## 2 Length
                                       0.499
                                                0.288
                                                            1.73 8.36e- 2
## 3 Diameter
                                       2.08
                                                0.313
                                                            6.63 3.90e-11
                                       0.245
                                                0.0695
                                                            3.53 4.27e- 4
## 4 Height
## 5 Whole_weight
                                       5.20
                                                0.399
                                                           13.1 5.34e-38
## 6 Shucked_weight
                                      -4.38
                                                0.258
                                                          -17.0 5.80e-62
## 7 Viscera_weight
                                      -0.995
                                                0.160
                                                           -6.23 5.10e-10
                                                            7.86 5.03e-15
## 8 Shell_weight
                                       1.70
                                                0.217
## 9 Sex I
                                                0.115
                                                           -8.08 8.79e-16
                                      -0.925
                                                           -3.15 1.67e- 3
## 10 Sex_M
                                      -0.329
                                                0.104
## 11 Sex_I_x_Shucked_weight
                                                0.0868
                                                            5.83 6.05e- 9
                                       0.506
## 12 Sex_M_x_Shucked_weight
                                                            3.59 3.35e- 4
                                       0.399
                                                0.111
## 13 Length_x_Diameter
                                      -2.85
                                                0.407
                                                           -6.99 3.23e-12
## 14 Shucked_weight_x_Shell_weight
                                                0.210
                                                           -1.26 2.08e- 1
                                      -0.264
```

```
## # A tibble: 1 x 1
## .pred
## <dbl>
## 1 23.3
```

The predicted age of the specifications above is  $\approx 23.32552$ .

#### Question 7:

Now I assess the model's performance using the yardstick package:

- 1. Create a metric set that includes  $R^2$ , RMSE (root mean squared error), and MAE (mean absolute error).
- 2. Use predict() and bind\_cols() to create a tibble of the model's predicted values from the training data along with the actual observed ages (these are needed to assess your model's performance).
- 3. Finally, apply the metric set to the tibble, report the results, and interpret the  $R^2$  value.

```
# creating a metric set
abalone metrics <- metric set(rmse, rsq, mae)
# creating a tibble of predicted values
abalone_train_res <- predict(lm_fit, new_data = abalone_train %>% select(-Age, -Rings))
abalone_train_res <- bind_cols(abalone_train_res, abalone_train %>% select(Age))
# applying the metric set to the tibble
abalone_metrics(abalone_train_res, truth = Age, estimate = .pred)
## # A tibble: 3 x 3
##
     .metric .estimator .estimate
             <chr>>
                            <dbl>
     <chr>>
                            2.17
## 1 rmse
             standard
                            0.553
## 2 rsq
             standard
## 3 mae
             standard
                            1.56
```

From above, we see that the RMSE  $\approx 2.1706344$ ,  $R^2 \approx 0.5534585$ , MAE  $\approx 1.5611926$ . The estimate of  $R^2$  implies that 53.50% of the variability in the response is explained by the predictors.