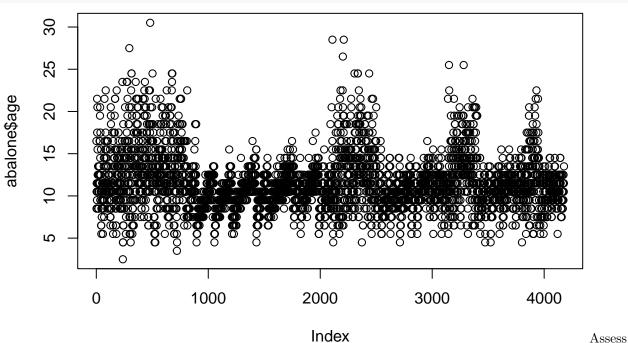
HW2

2022-10-04

1. Predict abalone age

```
abalone$age <- abalone$rings + 1.5
plot(abalone$age)
```



and describe the distribution of age. Majority of their ages seem to be between 8 to 13.

2.

```
set.seed(1738)

abalone_split <- initial_split(abalone, prop = 0.7, strata = age)
abalone_train <- training(abalone_split)
abalone_test <- testing(abalone_split)</pre>
```

3. Create a recipe predicting the outcome variable, age, with all other predictor variables. Explain why you shouldn't use rings to predict age.

```
abalone_recipe <- recipe(age ~ type + longest_shell + diameter + height + whole_weight + shucked_weight step_dummy(all_nominal_predictors()) %>% step_interact(~starts_with("type"):shucked_weight) %>% step_interact(~longest_shell:diameter) %>% step_interact(~shucked_weight:shell_weight) %>% step_interact(~shucked_weight:shell_weight) %>% step_center(longest_shell, diameter, height, whole_weight, shucked_weight, viscera_weight, shell_weight step_scale(longest_shell, diameter, height, whole_weight, shucked_weight, viscera_weight, shell_weight abalone_recipe
```

```
## Recipe
##
## Inputs:
##
##
         role #variables
##
      outcome
##
   predictor
##
## Operations:
##
## Dummy variables from all_nominal_predictors()
## Interactions with starts_with("type"):shucked_weight
## Interactions with longest_shell:diameter
## Interactions with shucked_weight:shell_weight
## Centering for longest_shell, diameter, height, whole_weight, ...
## Scaling for longest_shell, diameter, height, whole_weight, ...
Rings are directly correlated with age, so we can't use rings to predict age.
  4. Create and store a linear regression object using the "lm" engine.
lm_model <- linear_reg() %>%
  set_engine("lm")
  5. Workflow set up
lm_wkflow <- workflow() %>%
  add_model(lm_model) %>%
  add_recipe(abalone_recipe)
lm_wkflow
## == Workflow =====
## Preprocessor: Recipe
## Model: linear_reg()
##
## 6 Recipe Steps
##
## * step_dummy()
## * step_interact()
## * step_interact()
## * step_interact()
## * step_center()
## * step_scale()
##
## Linear Regression Model Specification (regression)
## Computational engine: lm
  6. Use your fit() object to predict the age of a hypothetical female abalone with longest_shell = 0.50, diam-
    eter = 0.10, height = 0.30, whole_weight = 4, shucked_weight = 1, viscera_weight = 2, shell_weight
lm_fit <- fit(lm_wkflow, abalone_train)</pre>
lm_fit %>%
  extract_fit_parsnip() %>%
```

```
tidy()
## # A tibble: 14 x 5
                                     estimate std.error statistic p.value
##
      term
##
      <chr>
                                        <dbl>
                                                   <dbl>
                                                             <dbl>
                                                                      <dbl>
## 1 (Intercept)
                                      18.3
                                                   1.02
                                                            17.8
                                                                   1.06e-67
## 2 longest_shell
                                       0.0463
                                                  0.318
                                                             0.146 8.84e- 1
## 3 diameter
                                                             6.67 3.00e-11
                                       2.35
                                                  0.353
## 4 height
                                       0.578
                                                  0.103
                                                             5.59 2.42e-8
## 5 whole_weight
                                       5.93
                                                  0.450
                                                            13.2
                                                                   1.20e-38
## 6 shucked_weight
                                      -4.60
                                                  0.281
                                                           -16.4
                                                                   8.09e-58
## 7 viscera_weight
                                      -1.18
                                                  0.173
                                                            -6.86 8.63e-12
## 8 shell_weight
                                                  0.235
                                                            4.27 1.99e- 5
                                       1.01
## 9 type_I
                                      -1.66
                                                  0.267
                                                            -6.23 5.41e-10
## 10 type_M
                                      -0.247
                                                            -1.07 2.86e- 1
                                                  0.231
## 11 type_I_x_shucked_weight
                                       3.94
                                                  0.818
                                                             4.82 1.53e- 6
## 12 type_M_x_shucked_weight
                                       0.856
                                                  0.472
                                                             1.81 6.99e- 2
## 13 longest_shell_x_diameter
                                                            -6.05 1.63e- 9
                                     -28.6
                                                  4.73
                                                            -0.538 5.90e- 1
## 14 shucked_weight_x_shell_weight -1.03
                                                  1.91
new_obs <- tibble(</pre>
  longest\_shell = c(0.5), diameter = c(0.10), height = c(0.30), whole_weight = c(4), shucked_weight = c(4)
new_pred <- new_obs %>%
  bind_cols(lm_fit %>%
              predict(new_data = new_obs))
new_pred
## # A tibble: 1 x 9
     longest_shell diameter height whole_weight shuck~1 visce~2 shell~3 type .pred
##
                       <dbl> <dbl>
                                           <dbl>
                                                   <dbl>
                                                            <dbl>
                                                                    <dbl> <chr> <dbl>
             <dbl>
               0.5
                         0.1
                                0.3
                                                        1
                                                                2
                                                                        1 F
                                                                                  21.9
## # ... with abbreviated variable names 1: shucked_weight, 2: viscera_weight,
       3: shell_weight
Predicted Age: 21.94
  7.
library(yardstick)
new_metric <- metric_set(rsq, rmse, mae)</pre>
preds <- abalone_train %>%
  select(age) %>%
  bind_cols(lm_fit %>%
              predict(abalone_train))
preds %>%
  new_metric(age, .pred)
## # A tibble: 3 x 3
##
     .metric .estimator .estimate
##
     <chr>
             <chr>>
                             <dbl>
## 1 rsq
             standard
                             0.554
## 2 rmse
             standard
                             2.16
             standard
                             1.55
```

R^2: 0.55 Root Mean Square Error: 2.16 Mean Absolute Error: 1.55

$ m R^2$ depicts how well the predicted values fit the actual 55% of the observed variabilty is explained by our model.	