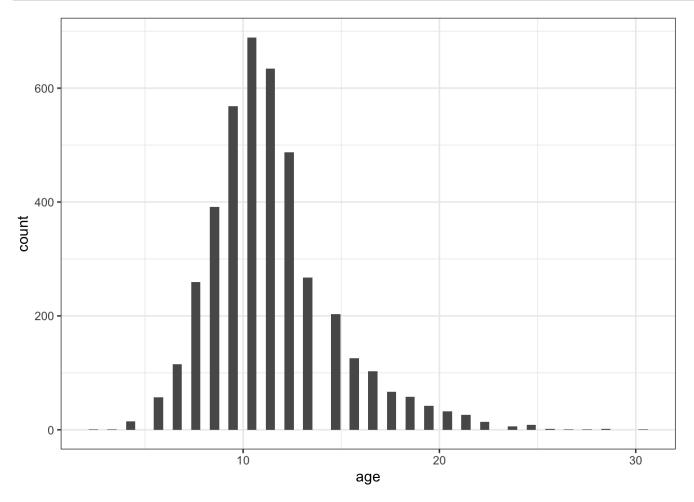
Homework2

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Question 1

```
setwd("/Users/safiyaalavi/Desktop/PSTAT 131/Homework 2/homework-2/data")
abalone_df <- read_csv("abalone.csv")
age <- abalone_df[,9]+1.5
abalone_df <- cbind(abalone_df, age)
names(abalone_df)[10] <- "age"
abalone_df %>%
    ggplot(aes(x = age)) +
    geom_histogram(bins = 60) +
    theme_bw()
```



The distribution of age predictor in the abalone data is clearly normally distributed. The distribution has positive skewness, or in other words the data is skewed to the right. The mean looks to be about 11.

Question 2

```
set.seed(3435)
abalone_split <- initial_split(abalone_df, prop = 3/4, strata = age)
abalone_train <- training(abalone_split)
abalone_test <- testing(abalone_split)</pre>
```

Question 3

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

```
## Recipe
##
## Inputs:
##
        role #variables
##
##
    outcome
## predictor
##
## Operations:
##
## Dummy variables from all_nominal_predictors()
## Interactions with starts_with("type"):shucked_weight + longest_shell...
## Centering for all_predictors()
## Scaling for all predictors()
```

We do not use rings to predict age because age is directly based on rings. Age is equal to 1.5 plus the value of rings, hence we cannot use rings to predict age because the information is the same.

Question 4

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

Question 5

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

Question 6

```
lm_fit <- fit(lm_wflow, abalone_train)
lm_fit %>% extract_fit_parsnip() %>% tidy()
```

```
## # A tibble: 14 × 5
##
    term
                                  estimate std.error statistic p.value
##
     <chr>
                                     <dbl>
                                              <dbl>
                                                        <dbl>
                                                                 <dbl>
## 1 (Intercept)
                                    11.4
                                              0.0382
                                                        299. 0
## 2 longest shell
                                     0.485
                                              0.288
                                                         1.68 9.22e- 2
## 3 diameter
                                     1.96
                                              0.313
                                                          6.24 4.91e-10
## 4 height
                                                          3.12 1.80e- 3
                                     0.218
                                              0.0698
## 5 whole_weight
                                     4.22
                                              0.406
                                                        10.4 6.66e-25
## 6 shucked weight
                                              0.264
                                                        -14.6 1.06e-46
                                    -3.85
                                                        -5.11 3.49e- 7
## 7 viscera weight
                                    -0.817
                                              0.160
## 8 shell weight
                                     1.95
                                              0.224
                                                        8.69 5.57e-18
                                    -0.934
## 9 type I
                                              0.118
                                                        -7.90 3.94e-15
## 10 type M
                                    -0.243
                                              0.106
                                                        -2.29 2.21e- 2
## 11 type_I_x_shucked_weight
                                              0.0890
                                                         5.46 5.23e- 8
                                    0.486
## 12 type_M_x_shucked_weight
                                     0.317
                                              0.113
                                                          2.81 4.99e- 3
## 13 longest shell x diameter
                                    -2.62
                                              0.404
                                                        -6.50 9.40e-11
## 14 shucked_weight_x_shell_weight
                                   -0.272
                                              0.204
                                                        -1.34 1.82e- 1
```

```
new_abalone <- tibble(type = "F", longest_shell = 0.50, diameter = 0.1, height = 0.3, wh
ole_weight = 4, shucked_weight = 1, viscera_weight = 2, shell_weight = 1, rings = 0)
predict(lm_fit, new_data = new_abalone)</pre>
```

```
## # A tibble: 1 × 1
## .pred
## <dbl>
## 1 22.6
```

Question 7