

hw2

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Question 1:

```
abalone <- read.csv("abalone.csv")  
# adding age variable  
abalone['age'] <- c(abalone$rings+1.5)  
head(abalone)
```

```
##   type longest_shell diameter height whole_weight shucked_weight viscera_weight  
## 1    M         0.455    0.365  0.095      0.5140         0.2245         0.1010  
## 2    M         0.350    0.265  0.090      0.2255         0.0995         0.0485  
## 3    F         0.530    0.420  0.135      0.6770         0.2565         0.1415  
## 4    M         0.440    0.365  0.125      0.5160         0.2155         0.1140  
## 5    I         0.330    0.255  0.080      0.2050         0.0895         0.0395  
## 6    I         0.425    0.300  0.095      0.3515         0.1410         0.0775  
##   shell_weight rings  age  
## 1         0.150    15 16.5  
## 2         0.070     7  8.5  
## 3         0.210     9 10.5  
## 4         0.155    10 11.5  
## 5         0.055     7  8.5  
## 6         0.120     8  9.5
```

The distribution of age is the same as 'rings', shifted right by 1.5 units.

Question 2:

```
set.seed(1027)  
abalone_split <- initial_split(abalone, prop = 0.7, strata = age)  
abalone_train <- training(abalone_split)  
abalone_test  <- testing(abalone_split)
```

Question 3:

we should not use rings to predict age because age is a function of rings. rings would be a perfect predictor for age, creating bias towards the entire regression and the rest of the estimator coefficients.

```
abalone_train_mod <- abalone_train %>% select(-rings)  
abalone_recipe <-  
  recipe(age~., data = abalone_train_mod)
```

```

abalone_recipe %>%
  step_dummy(all_nominal_predictors()) %>%
  step_center(all_predictors()) %>%
  step_scale(all_predictors()) %>%
  step_interact(~ starts_with("type"):shucked_weight) %>%
  step_interact(~ longest_shell:diameter) %>%
  step_interact(~ shucked_weight:shell_weight)

```

```

## Recipe
##
## Inputs:
##
##      role #variables
## outcome      1
## predictor      8
##
## Operations:
##
## Dummy variables from all_nominal_predictors()
## Centering for all_predictors()
## Scaling for all_predictors()
## Interactions with starts_with("type"):shucked_weight
## Interactions with longest_shell:diameter
## Interactions with shucked_weight:shell_weight

```

Question 4:

```

lm_mod <-linear_reg() %>%
  set_engine("lm")

```

Question 5:

```

lm_wfl<-workflow() %>%
  add_model(lm_mod) %>%
  add_recipe(abalone_recipe)

```

Question 6:

```

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

```

```

## # A tibble: 10 x 5
##   term          estimate std.error statistic  p.value
##   <chr>          <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)    5.44      0.344     15.8 4.26e-54

```

```
## 2 typeI      -0.832    0.120    -6.91  5.91e-12
## 3 typeM       0.171    0.0985    1.73  8.33e- 2
## 4 longest_shell  0.649    2.10     0.309 7.58e- 1
## 5 diameter    10.2     2.60     3.94  8.41e- 5
## 6 height      8.79     1.65     5.32  1.14e- 7
## 7 whole_weight 10.2     0.939    10.8  8.69e-27
## 8 shucked_weight -21.4    1.06    -20.3  2.04e-85
## 9 viscera_weight -12.0    1.60    -7.46  1.13e-13
## 10 shell_weight  7.97     1.42     5.59  2.45e- 8
```

```
new <-data.frame(type="F", longest_shell = 0.50, diameter = 0.10,
                 height = 0.30, whole_weight = 4, shucked_weight = 1,
                 viscera_weight = 2, shell_weight = 1)
predict(lm_fit, new)
```

```
## # A tibble: 1 x 1
##   .pred
##   <dbl>
## 1  12.7
```

The predicted age for this abalone is 12.72 years old.

Question 7:

```
abalone_train_res<- predict(lm_fit, new_data = abalone_train %>%select(-age))
abalone_train_res<-bind_cols(abalone_train_res, abalone_train %>% select(
  age
))
abalone_metrics<-metric_set(rmse, rsq, mae)
abalone_metrics(abalone_train_res, truth = age,
                estimate=.pred)
```

```
## # A tibble: 3 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>      <dbl>
## 1 rmse    standard      2.17
## 2 rsq     standard      0.539
## 3 mae     standard      1.56
```

$R^2 = 0.5394$. This means that about 53.94% of the variation in age can be explained by the predictor variables in the regression created above.