

Homework 3

PSTAT 131/231

Contents

```
library(tidymodels)
library(tidyverse)
library(corrplot)
library(discrim)
library(poissonreg)
library(corr)
library(klaR)
titanic <- read.csv("titanic.csv")
titanic$survived <- factor(titanic$survived, levels = c("Yes", "No"))
titanic$pclass <- factor(titanic$pclass)
head(titanic)

##   passenger_id survived pclass
## 1             1       No     3
## 2             2      Yes     1
## 3             3      Yes     3
## 4             4      Yes     1
## 5             5       No     3
## 6             6       No     3
##                                     name   sex age sib_sp parch
## 1           Braund, Mr. Owen Harris male  22     1     0
## 2 Cumings, Mrs. John Bradley (Florence Briggs Thayer) female 38     1     0
## 3          Heikkinen, Miss. Laina female 26     0     0
## 4    Futrelle, Mrs. Jacques Heath (Lily May Peel) female 35     1     0
## 5          Allen, Mr. William Henry male  35     0     0
## 6          Moran, Mr. James male   NA     0     0
##   ticket   fare cabin embarked
## 1 A/5 21171 7.2500   <NA>      S
## 2   PC 17599 71.2833    C85      C
## 3 STON/O2. 3101282 7.9250   <NA>      S
## 4        113803 53.1000    C123      S
## 5        373450  8.0500   <NA>      S
## 6        330877  8.4583   <NA>      Q
```

Question1 Split the data, stratifying on the outcome variable, `survived`. You should choose the proportions to split the data into. Verify that the training and testing data sets have the appropriate number of observations. Take a look at the training data and note any potential issues, such as missing data.

Why is it a good idea to use stratified sampling for this data?

```

set.seed(1979)
titanic_split <- initial_split(titanic, prop = 0.8,
                                strata = survived)
titanic_train <- training(titanic_split)
titanic_test <- testing(titanic_split)
c(nrow(titanic_train), nrow(titanic_test))

```

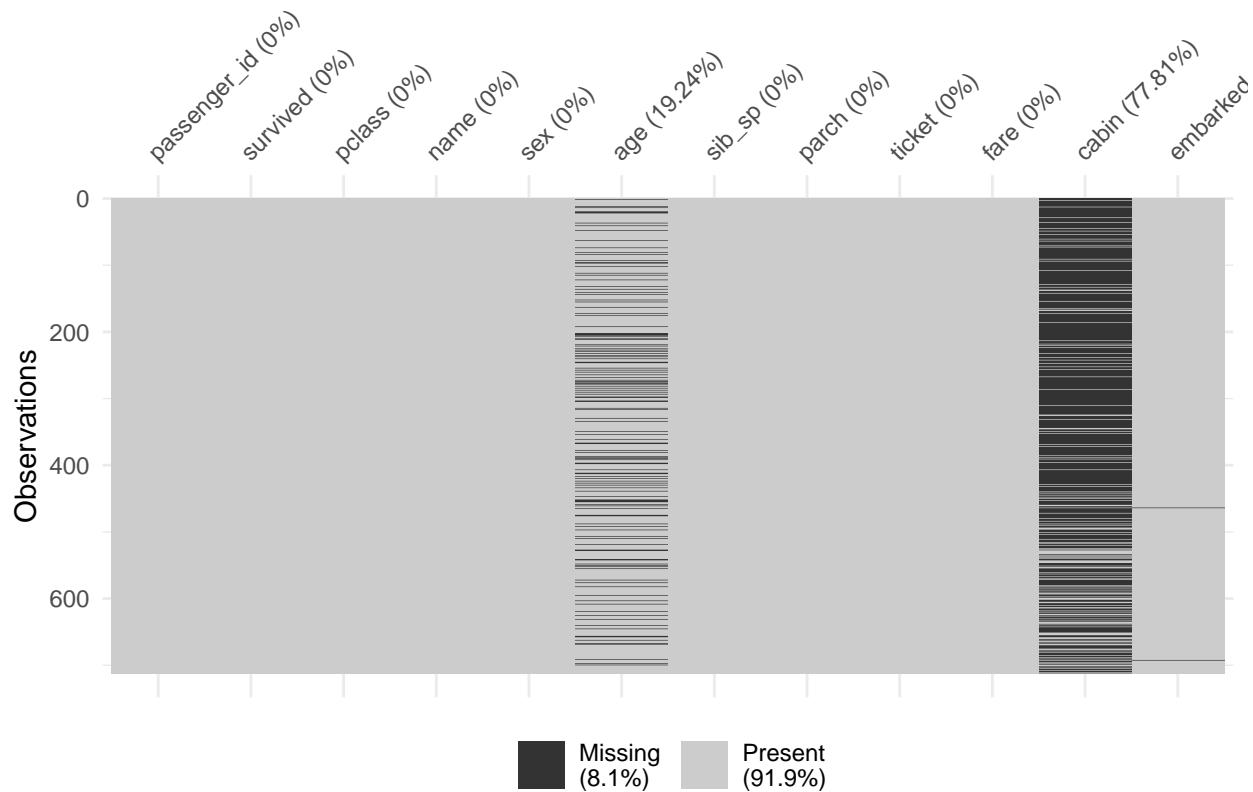
```
## [1] 712 179
```

The training data sets have 712 observation and testing data sets have 179 observation. They both have the appropriate number of observation

```

library(naniar)
vis_miss(titanic_train)

```

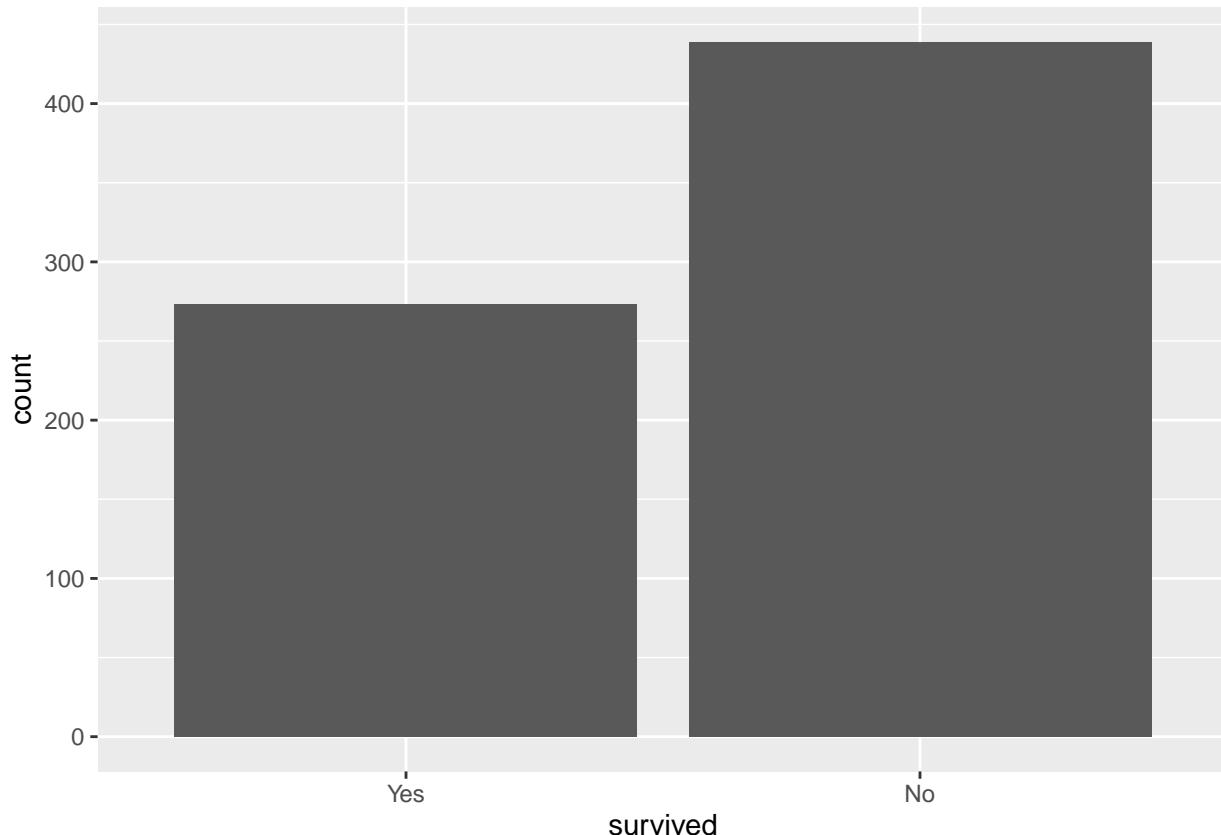


We have 77.81% missing value in cabin and 19.24% in age.

Why is it a good idea to use stratified sampling for this data? + Stratified sampling can make a representative amount of each level (No/Yes) of variable (survived) is included in the training and testing data set.

Question2 Using the **training** data set, explore/describe the distribution of the outcome variable **survived**.

```
titanic_train %>%
  ggplot(aes(x = survived)) +
  geom_bar()
```

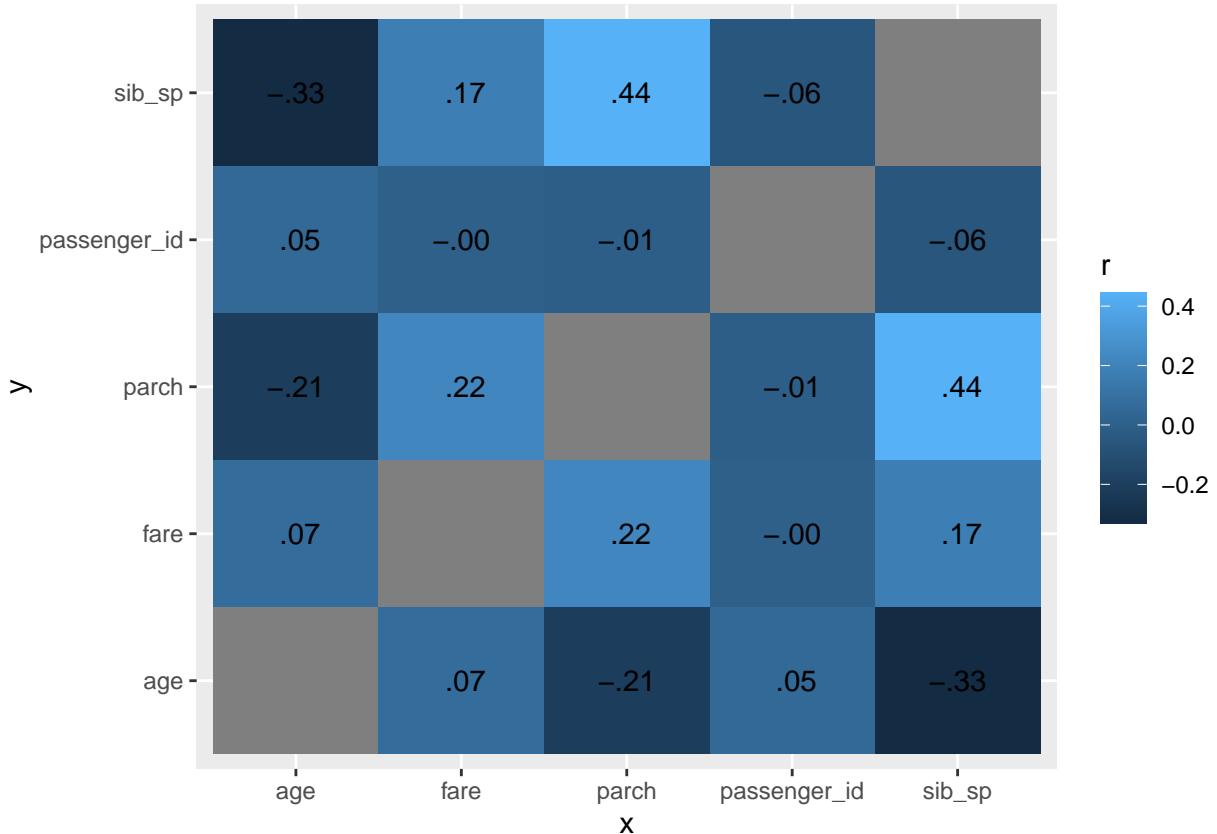


There are more died passenger than survived, with 40% dead and 60% survived.

Question3 Using the **training** data set, create a correlation matrix of all continuous variables. Create a visualization of the matrix, and describe any patterns you see. Are any predictors correlated with each other? Which ones, and in which direction?

```
library(corr)
cor_titanic <- titanic_train %>%
  select_if(is.numeric) %>%
  correlate()

cor_titanic %>%
  stretch() %>%
  ggplot(aes(x, y, fill = r)) +
  geom_tile() +
  geom_text(aes(label = as.character(fashion(r))))
```



We can also use ggplot and the geom_tile() function to create a heatmap-style correlation plot. Parch is positive correlated with sib_sp with correlation 0.44. Parch is positive correlated with fare with correlation 0.22. Sib_sp is negative correlated with age with correlation -0.33. Sib_sp is positive correlated with fare with correlation 0.17. Most of variables are not correlated with each other with correlation close to 0.

Question4 Using the **training** data, create a recipe predicting the outcome variable **survived**. Include the following predictors: ticket class, sex, age, number of siblings or spouses aboard, number of parents or children aboard, and passenger fare.

```
simple_titanic_recipe <-  
  recipe(survived ~ pclass+sex+age+sib_sp+parch+fare, data = titanic_train)%>%  
  step_impute_linear(age) %>%  
  step_dummy(all_nominal_predictors()) %>%  
  step_interact(~ fare:starts_with("sex") + age:fare)
```

Question5 Specify a **logistic regression** model for classification using the "glm" engine. Then create a workflow. Add your model and the appropriate recipe. Finally, use **fit()** to apply your workflow to the **training** data.

```
log_model <- logistic_reg() %>%  
  set_engine("glm")  
  
log_wflow <- workflow() %>%  
  add_model(log_model) %>%  
  add_recipe(simple_titanic_recipe)
```

```
log_fit <- fit(log_wf, titanic_train)
```

Question6 Repeat **Question 5**, but this time specify a linear discriminant analysis model for classification using the "MASS" engine.

```
lda_mod <- discrim_linear() %>%
  set_mode("classification") %>%
  set_engine("MASS")

lda_wf <- workflow() %>%
  add_model(lda_mod) %>%
  add_recipe(simple_titanic_recipe)

lda_fit <- fit(lda_wf, titanic_train)
```

Question7 Repeat **Question 5**, but this time specify a quadratic discriminant analysis model for classification using the "MASS" engine.

```
qda_mod <- discrim_quad() %>%
  set_mode("classification") %>%
  set_engine("MASS")

qda_wf <- workflow() %>%
  add_model(qda_mod) %>%
  add_recipe(simple_titanic_recipe)

qda_fit <- fit(qda_wf, titanic_train)
```

Question8 Repeat **Question 5**, but this time specify a naive Bayes model for classification using the "klaR" engine. Set the `usekernel` argument to FALSE.

```
nb_mod <- naive_Bayes() %>%
  set_mode("classification") %>%
  set_engine("klaR") %>%
  set_args(usekernel = FALSE)

nb_wf <- workflow() %>%
  add_model(nb_mod) %>%
  add_recipe(simple_titanic_recipe)

nb_fit <- fit(nb_wf, titanic_train)
```

Question9 Now you've fit four different models to your training data.

Use `predict()` and `bind_cols()` to generate predictions using each of these 4 models and your **training** data. Then use the *accuracy* metric to assess the performance of each of the four models.

Which model achieved the highest accuracy on the training data?

```

predict(log_fit, new_data = titanic_train, type = "prob") %>%
  bind_cols(titanic_train)

## # A tibble: 712 x 14
##   .pred_Yes .pred_No passenger_id survived pclass name      sex    age sib_sp
##   <dbl>    <dbl>        <int> <fct>    <fct>  <chr>     <chr> <dbl> <int>
## 1 0.112    0.888       1 No      3 Braund, M~ male    22     1
## 2 0.0756   0.924       5 No      3 Allen, Mr~ male   35     0
## 3 0.111    0.889       6 No      3 Moran, Mr~ male  NA     0
## 4 0.324    0.676       7 No      1 McCarthy, ~ male  54     0
## 5 0.134    0.866       8 No      3 Palsson, ~ male  2      3
## 6 0.182    0.818      13 No     3 Saunderc~ male  20     0
## 7 0.0495   0.950      14 No     3 Andersson~ male 39     1
## 8 0.794    0.206      15 No     3 Vestrom, ~ fema~ 14     0
## 9 0.0774   0.923      17 No     3 Rice, Mas~ male  2      4
## 10 0.463   0.537      19 No     3 Vander Pl~ fema~ 31     1
## # ... with 702 more rows, and 5 more variables: parch <int>, ticket <chr>,
## #   fare <dbl>, cabin <chr>, embarked <chr>

predict(qda_fit, new_data = titanic_train, type = "prob") %>%
  bind_cols(titanic_train)

## # A tibble: 712 x 14
##   .pred_Yes .pred_No passenger_id survived pclass name      sex    age sib_sp
##   <dbl>    <dbl>        <int> <fct>    <fct>  <chr>     <chr> <dbl> <int>
## 1 0.00549  0.995       1 No      3 Braund, ~ male  22     1
## 2 0.00373  0.996       5 No      3 Allen, ~ male 35     0
## 3 0.00525  0.995       6 No      3 Moran, ~ male NA     0
## 4 0.135    0.865       7 No      1 McCarth~ male 54     0
## 5 0.000124 1.00        8 No      3 Palsson~ male  2      3
## 6 0.00875  0.991      13 No     3 Saunderc~ male 20     0
## 7 0.550    0.450      14 No     3 Andersss~ male 39     1
## 8 0.503    0.497      15 No     3 Vestrom~ fema~ 14     0
## 9 0.000000685 1.00      17 No     3 Rice, M~ male  2      4
## 10 0.232   0.768      19 No     3 Vander ~ fema~ 31     1
## # ... with 702 more rows, and 5 more variables: parch <int>, ticket <chr>,
## #   fare <dbl>, cabin <chr>, embarked <chr>

predict(lda_fit, new_data = titanic_train, type = "prob") %>%
  bind_cols(titanic_train)

## # A tibble: 712 x 14
##   .pred_Yes .pred_No passenger_id survived pclass name      sex    age sib_sp
##   <dbl>    <dbl>        <int> <fct>    <fct>  <chr>     <chr> <dbl> <int>
## 1 0.0694   0.931       1 No      3 Braund, M~ male  22     1
## 2 0.0481   0.952       5 No      3 Allen, Mr~ male 35     0
## 3 0.0688   0.931       6 No      3 Moran, Mr~ male NA     0
## 4 0.260    0.740       7 No      1 McCarthy, ~ male 54     0
## 5 0.0878   0.912       8 No      3 Palsson, ~ male 2      3
## 6 0.110    0.890      13 No     3 Saunderc~ male 20     0
## 7 0.0322   0.968      14 No     3 Andersson~ male 39     1
## 8 0.841    0.159      15 No     3 Vestrom, ~ fema~ 14     0

```

```

##   9    0.0551    0.945          17 No      3      Rice, Mas~ male    2    4
## 10    0.567     0.433          19 No      3      Vander Pl~ fema~  31    1
## # ... with 702 more rows, and 5 more variables: parch <int>, ticket <chr>,
## #   fare <dbl>, cabin <chr>, embarked <chr>

# naive Bayes model
predict(nb_fit, new_data = titanic_train, type = "prob") %>%
  bind_cols(titanic_train)

## # A tibble: 712 x 14
##   .pred_Yes .pred_No passenger_id survived pcclass name       sex     age sib_sp
##   <dbl>     <dbl>        <int> <fct>   <fct>   <chr>     <chr> <dbl> <int>
## 1 0.0155     0.984         1 No     3 Braund, ~ male    22    1
## 2 0.0152     0.985         5 No     3 Allen, M~ male   35    0
## 3 0.0158     0.984         6 No     3 Moran, M~ male  NA    0
## 4 0.573      0.427         7 No     1 McCarthy~ male  54    0
## 5 0.000278    1.00        8 No     3 Palsson,~ male  2     3
## 6 0.0176     0.982        13 No    3 Saunderc~ male  20    0
## 7 0.0960     0.904        14 No    3 Andersso~ male  39    1
## 8 0.475      0.525        15 No    3 Vestrom,~ fema~ 14    0
## 9 0.00000328  1.00        17 No    3 Rice, Ma~ male  2     4
## 10 0.383     0.617        19 No    3 Vander P~ fema~ 31    1
## # ... with 702 more rows, and 5 more variables: parch <int>, ticket <chr>,
## #   fare <dbl>, cabin <chr>, embarked <chr>

log_reg_acc <- augment(log_fit, new_data = titanic_train) %>%
  accuracy(truth = survived, estimate = .pred_class)

lda_acc <- augment(lda_fit, new_data = titanic_train) %>%
  accuracy(truth = survived, estimate = .pred_class)

qda_acc <- augment(qda_fit, new_data = titanic_train) %>%
  accuracy(truth = survived, estimate = .pred_class)
qda_acc

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>        <dbl>
## 1 accuracy binary     0.760

nb_acc <- augment(nb_fit, new_data = titanic_train) %>%
  accuracy(truth = survived, estimate = .pred_class)
nb_acc

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>        <dbl>
## 1 accuracy binary     0.760

```

```

accuracies <- c(log_reg_acc$.estimate, lda_acc$.estimate,
                 nb_acc$.estimate, qda_acc$.estimate)
models <- c("Logistic Regression", "LDA", "Naive Bayes", "QDA")
results <- tibble(accuracies = accuracies, models = models)
results %>%
  arrange(-accuracies)

## # A tibble: 4 x 2
##   accuracies models
##       <dbl> <chr>
## 1     0.808 Logistic Regression
## 2     0.799 LDA
## 3     0.760 Naive Bayes
## 4     0.760 QDA

```

Logistic Regression achieved highest accuracy on the training data

Question 10 Fit the model with the highest training accuracy to the **testing** data. Report the accuracy of the model on the **testing** data.

Again using the **testing** data, create a confusion matrix and visualize it. Plot an ROC curve and calculate the area under it (AUC).

How did the model perform? Compare its training and testing accuracies. If the values differ, why do you think this is so?

```

predict(log_fit, new_data = titanic_test, type = "prob")

## # A tibble: 179 x 2
##   .pred_Yes .pred_No
##       <dbl>    <dbl>
## 1     0.819    0.181
## 2     0.219    0.781
## 3     0.415    0.585
## 4     0.0322   0.968
## 5     0.324    0.676
## 6     0.111    0.889
## 7     0.757    0.243
## 8     0.292    0.708
## 9     0.382    0.618
## 10    0.814    0.186
## # ... with 169 more rows

multi_metric <- metric_set(accuracy, sensitivity, specificity)

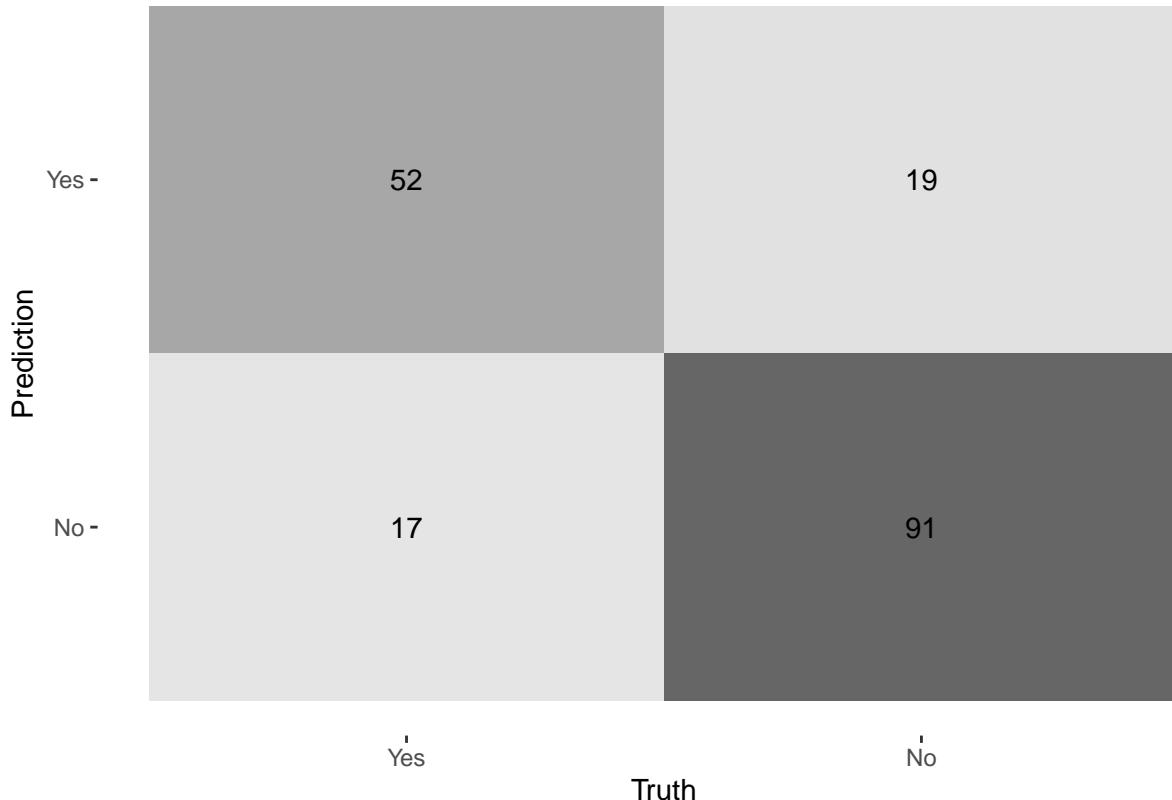
augment(log_fit, new_data = titanic_test) %>%
  multi_metric(truth = survived, estimate = .pred_class)

## # A tibble: 3 x 3
##   .metric      .estimator .estimate
##   <chr>        <chr>        <dbl>
## 1 accuracy    binary      0.799
## 2 sensitivity binary      0.754
## 3 specificity binary      0.827

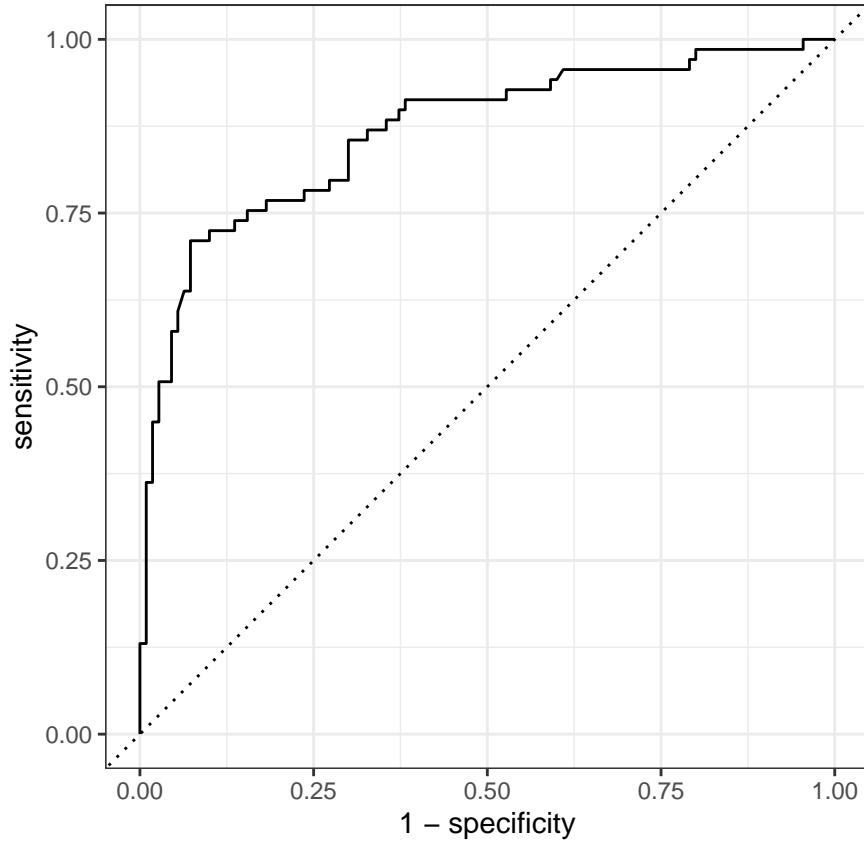
```

The accuracy of the model on the testing data is 0.7988827

```
augment(log_fit, new_data = titanic_test) %>%
  conf_mat(truth = survived, estimate = .pred_class) %>%
  autoplot(type = "heatmap")
```



```
augment(log_fit, new_data = titanic_test) %>%
  roc_curve(survived, .pred_Yes) %>%
  autoplot()
```



```
augment(log_fit, new_data = titanic_test) %>%
  roc_auc(survived, .pred_Yes)
```

```
## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>       <dbl>
## 1 roc_auc binary     0.867
```

The model performs well, and the area under ROC-curve is 0.86. The accuracy between training and testing is similar and both close to 80.