# PSTAT 131 Homework 4

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2022-04-26

### Question 1:

Splitting the data set

We split the data set and stratify on the variable survived.

```
set.seed(1004)
titanic_split <- initial_split(titanic, prop = 0.70, strata = survived)
titanic_train <- training(titanic_split)
titanic_test <- testing(titanic_split)</pre>
```

Verifying that the testing and training sets have the appropriate number of observations.

```
nrow(titanic_train)
## [1] 623
nrow(titanic_test)
## [1] 268
```

# Question 2:

Folding the training set

We fold the training set and use k-fold cross validation with k = 10.

```
titanic_folds <- vfold_cv(titanic_train, v = 10)
titanic_folds</pre>
```

```
## # 10-fold cross-validation
## # A tibble: 10 x 2
## splits id
## <list> <chr>
## 1 <split [560/63]> Fold01
## 2 <split [560/63]> Fold02
## 3 <split [560/63]> Fold03
## 4 <split [561/62]> Fold04
## 5 <split [561/62]> Fold05
```

```
## 6 <split [561/62]> Fold06
## 7 <split [561/62]> Fold07
## 8 <split [561/62]> Fold08
## 9 <split [561/62]> Fold09
## 10 <split [561/62]> Fold10
```

#### Question 3:

K-fold cross-validation splits the data into k number of folds to assess model performance. It is used to select the best model for our dataset. We use k-fold cross-validation to compare the best value to our model, rather than simply fitting and testing models. If we did use the entire training set, resampling method is called the validation set approach.

#### Question 4:

Setting up workflows for 3 models We will be setting up workflows for the following 3 models:

- A logistic regression with the glm engine;
- A linear discriminant analysis with the MASS engine;
- A quadratic discriminant analysis with the MASS engine.

```
# Logistic Regression
log_reg <- logistic_reg() %>%
  set_engine("glm") %>%
  set_mode("classification")

log_wkflow <- workflow() %>%
  add_model(log_reg) %>%
  add_recipe(titanic_recipe)
```

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

lda_wkflow <- workflow() %>%
  add_model(lda_mod) %>%
  add_recipe(titanic_recipe)
```

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

qda_wkflow <- workflow() %>%
  add_model(qda_mod) %>%
  add_recipe(titanic_recipe)
```

We will be fitting a total of 30 models across all folds since there are 3 models, Logistic Regression, Linear Discriminant Analysis, and Quadratic Discriminant Analysis and with 10 folds, there will be 30 models total.

#### Question 5:

Fitting the models into the folded data

```
log_fit <- log_wkflow %>%
  fit_resamples(titanic_folds)

lda_fit <- lda_wkflow %>%
  fit_resamples(titanic_folds)

qda_fit <- qda_wkflow %>%
  fit_resamples(titanic_folds)
```

# Question 6:

We use **collect\_metrics()** to print the mean and standard errors of the performance metric *accuracy* across all folds for each of the four models.

```
collect_metrics(log_fit)
## # A tibble: 2 x 6
##
    .metric .estimator mean
                                 n std_err .config
                        <dbl> <int>
    <chr>>
             <chr>
                                     <dbl> <chr>
## 1 accuracy binary
                        0.811 10 0.0153 Preprocessor1_Model1
## 2 roc_auc binary
                        0.845
                                 10 0.0156 Preprocessor1_Model1
collect_metrics(lda_fit)
## # A tibble: 2 x 6
    .metric .estimator mean
                                 n std_err .config
##
    <chr>>
             <chr>
                        <dbl> <int>
                                      <dbl> <chr>
## 1 accuracy binary
                        0.799
                                 10 0.0181 Preprocessor1_Model1
                                 10 0.0156 Preprocessor1_Model1
## 2 roc_auc binary
                        0.843
```

#### collect\_metrics(qda\_fit)

From above, we note that the best model is the logistic regression model. It has the lowest accuracy standard error at 0.01531636 and the highest mean accuracy at 0.8107015.

#### Question 7:

Now, we fit the Logistic Regression model to the entire training dataset.

0.819

```
log_fit <- fit(log_wkflow, titanic_train)</pre>
```

#### Question 8:

## 1 accuracy binary

Finally, we use predict(), bind\_cols(), and accuracy() to assess the model's performance on the testing data.

```
log_predict <- predict(log_fit, new_data = titanic_train, type = "prob")
log_predict <- bind_cols(log_predict, titanic_train %>% select(survived))
augment(log_fit, new_data = titanic_train) %>%
    accuracy(as.factor(survived), estimate = .pred_class)

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

From the above results, we have that our model's testing accuracy is estimated to be 0.8186196. This is a high number because the average accuracy across folds was 0.810701, which shows that our model's testing accuracy is higher than the data from the 10 folds.