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# Final Project

Code ▼

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## Introduction

My project will outline predictor variables of interest that could potentially predict the severity of a car accident.

## What's in this dataset? What is it?

To begin, the dataset is a countrywide record of car accidents from 49 different states. The data set covers a wide time frame where it tracks monthly and daily data spanning from 2016 to 2021. It's important to note that the data was collected from various sources—such as the US and state departments of transportation, law enforcement agencies, traffic cameras, and traffic sensors, etc. There's a wide variety of data, with over 1.5 accidents, or observations, that have been recorded with the highest frequencies of recorded accidents being from the state of California. This model contain various variables, but most notably, one variable contained is the severity attribute which spans from 1-4, where 1 is the lowest severity and 4 is the highest. This severity is determined by it's impact on traffic, or the shortest delay on oncoming traffic as a result of the accident.

## Why even create the model?

I'll list further details on the variables below, but essentially, the original data set contained nearly 47 different variables. All these variables levels in importance based on what one might be interested in predicting, but as I noted previously, I want to see what factors may be especially important in predicting severity of a certain accident. Many times we are forced to drive in various different weather conditions and usually we bat a blind eye towards the dangers that may result from these conditions. Weather conditions

such as wind speed, weather conditions (rain, snow, etc.), night/day, temperature, and maybe even the presence of certain road factors (roundabouts, cross ways, etc.). It may be important to know the relationship of these accidents and how they might vary in severity to potentially raise awareness to future drivers.

## Loading Data and Packages

The project was found through Kaggle, and as I noted before, there were nearly 1.5 million observations in the data. Therefore, I wasn't able to import the raw data as a whole into R, therefore I must note that I did some light preliminary cleaning of the data set by filtering by only what I will be using as the subject of my analysis, 2021 California data. Therefore, after filtering out blank inputs, and grouping by 2021 and California exclusive cases, we now have a new data set that I can do further cleaning on. Let's clean the data a bit before displaying the full table and splitting the data for further analysis.

Hide

```
library(ggplot2)
library(readr)
library(tidymodels)
library(tidyverse)
library(ISLR)
library(ISLR2)
library(discrim)
library(janitor)
library(poissonreg)
library(pROC)
library(corr)
library(corrplot)
library(purrr)
library(caret)
library(dplyr)
library(binom)
library(themis)
library(randomForest)
library(parsnip)
library(rpart)
library(rpart.plot)
library(ranger)
library(vip)
library(xgboost)
library(kknn)
library(rpart)
```

Hide

```
accidents_data <- read_csv("D:/UCSB/PSTAT131/Final Project/data/accidents1.csv")
```

## Data Cleaning

Below, we will be removing some unneeded variables, renaming some columns, and other techniques for needed clarity in reading the data. Note, there is no need to remove missing/blank observations since I have already made sure they were not included through excel and before importing the csv.

- Check for any remaining na's

Hide

```
accidents <- na.omit(accidents_data)
sum(is.na(accidents))
```

```
## [1] 0
```

- Remove the start\_time and state column since I had already filtered that completely in excel. variables

Hide

```
accidents <- accidents %>%
  select(-Start_Time,-State,-Wind_Direction)
```

- Cleaning the variable names for clarity

Hide

```
accidents <- janitor::clean_names(accidents)
```

- Now I must factor encode all the variables that are boolean values.

Hide

```
accidents$severity <- factor(accidents$severity)
accidents$side <- factor(accidents$side)
accidents$weather_condition <- factor(accidents$weather_condition)
accidents$bump <- factor(accidents$bump)
accidents$crossing <- factor(accidents$crossing)
accidents$railway <- factor(accidents$railway)
accidents$stop <- factor(accidents$stop)
accidents$traffic_signal <- factor(accidents$traffic_signal)
accidents$sunrise_sunset <- factor(accidents$sunrise_sunset)
```

Now the data is cleaned, and we can proceed with the analysis

## Variables

Below are noteworthy variables that will be used in our analysis (note that there are 17 variables, but I chose only select few important ones to display here. A full, detailed explanation of each variable can be found in the codebook in the zip file.)

<b>Table 1: Variable Details</b>
----------------------------------

Variable Name	Description	Type
severity	Shows the severity of the accident, a number between 1 and 4, where 1 indicates the least impact on traffic (i.e., short delay as a result of the accident) and 4 indicates a significant impact on traffic (i.e., long delay).	Factor
Distance(mi)	The length of the road extent affected by the accident	Numeric
visibility_mi	Shows visibility (in miles).	Numeric
wind_speed_mph	Shows wind speed (in miles per hour).	Numeric
precipitation_in	Shows precipitation amount in inches, if there is any.	Numeric
weather_condition	Shows the weather condition (rain, snow, thunderstorm, fog, etc.)	Factor
crossing	Presence of crossing in a nearby location.	Factor
stop	Presence of stop in a nearby location.	Factor
traffic_signal	Presence of traffic signal in a nearby location.	Factor
sunrise_sunset	Shows the period of day (i.e. day or night) based on sunrise/sunset.	Factor

## Data Split

I split my data using a proportion of 80/20, where the training data is 80% of my total data, and the testing data covers the other 20%. Because I had a large imbalance in the response variable, severity, then I must use stratified sampling (which you can see in the EDA portion).

Hide

```
accidents_split <- initial_split(accidents, prop = 0.70, strata = severity)
accidents_train <- training(accidents_split)
accidents_test <- testing(accidents_split)

# save(accidents, accidents_recipe, accidents_train, file = "D:/UCSB/PSTAT131/Final Project/R
_Scripts/processed\\accidents.rda")
# save(accidents_train, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/processed\\acciden
ts_train.rda")
# save(accidents_test, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/processed\\accident
s_test.rda")
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/processed\\accidents.rda")
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/processed\\accidents_train.rda")
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/processed\\accidents_test.rda")
```

Now I run a sanity check on the dimensions of the testing and training set to see if they match. And as we can see below, they do match in dimensions so there aren't any issues here. Now let's proceed to the next section, exploratory data analysis.

Hide

```
dim(accidents)
```

```
## [1] 30605    17
```

Hide

```
dim(accidents_train) + dim(accidents_test)
```

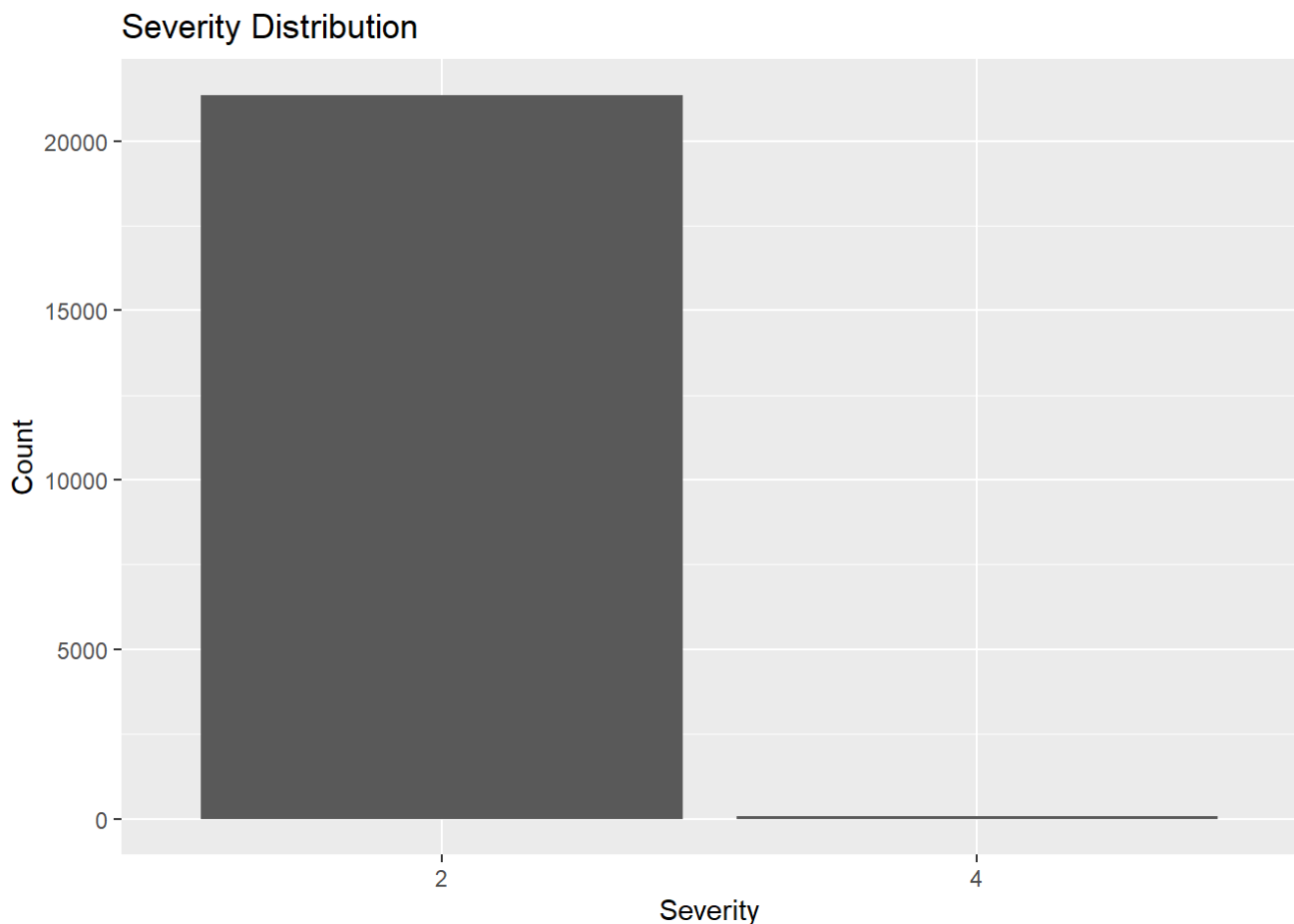
```
## [1] 30605    34
```

## Exploratory Data Analysis

The proceeding section will be exploratory data analysis on the training set to find any potential patterns and expectations we might have before we proceed with fitting the models.

### Severity Distribution

Code



Here is the proportion of the response variable with a severity of 4.

Hide

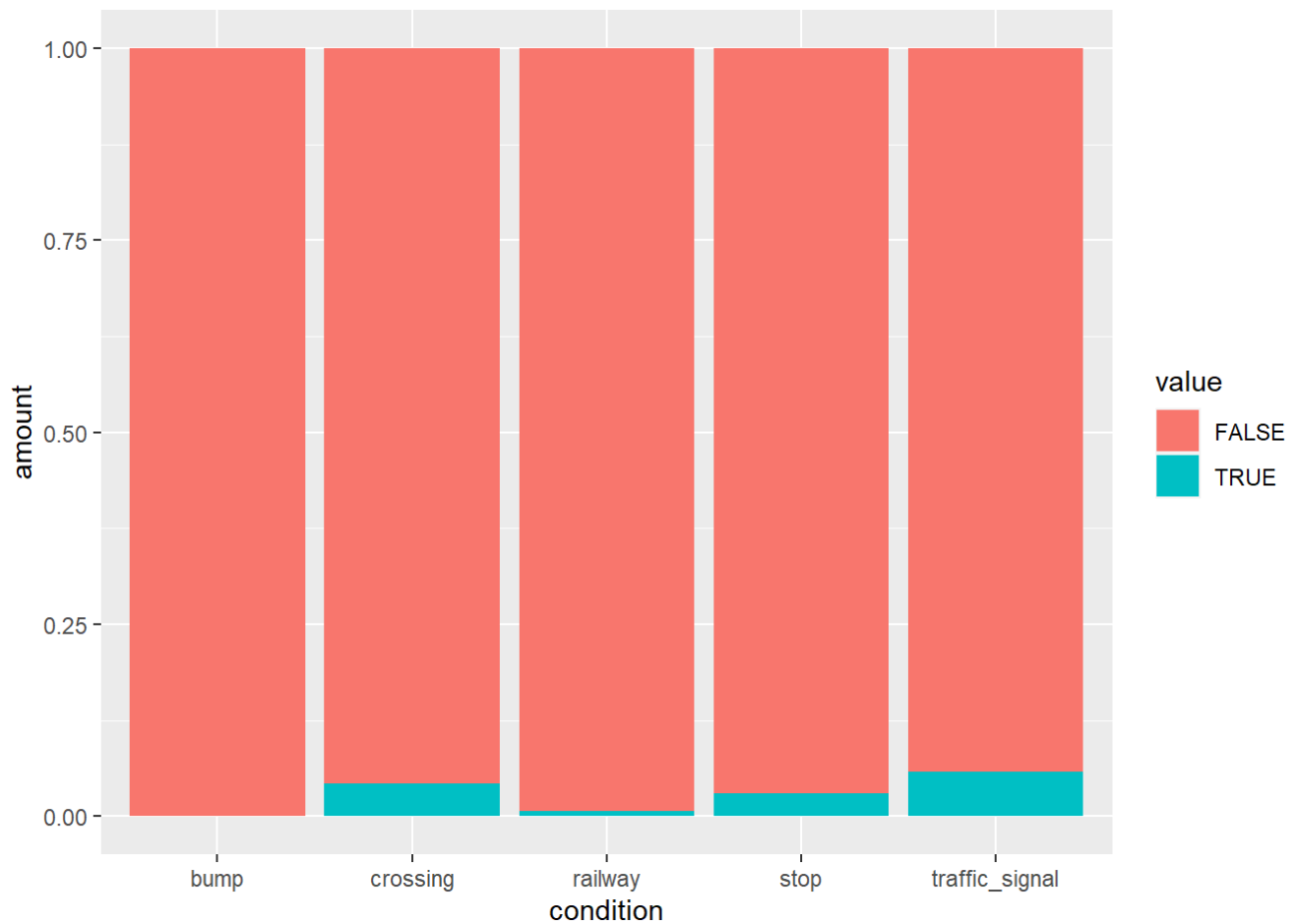
```
sum(accidents_train$severity=="4")/nrow(accidents_train)
```

```
## [1] 0.002894086
```

Notice that we find ourselves an extremely prevalent class unbalance in our response observations, therefore, something such as down sampling will be extremely valuable later on when creating the recipe. Also, notice how the response variables only contain 2 and 4 despite the range being from 1-4, therefore it appears all the accidents were either mild or extreme in terms of severity, so therefore our recipe will be a binary classification models.

## Exploring the Location landmarks Factor Variables

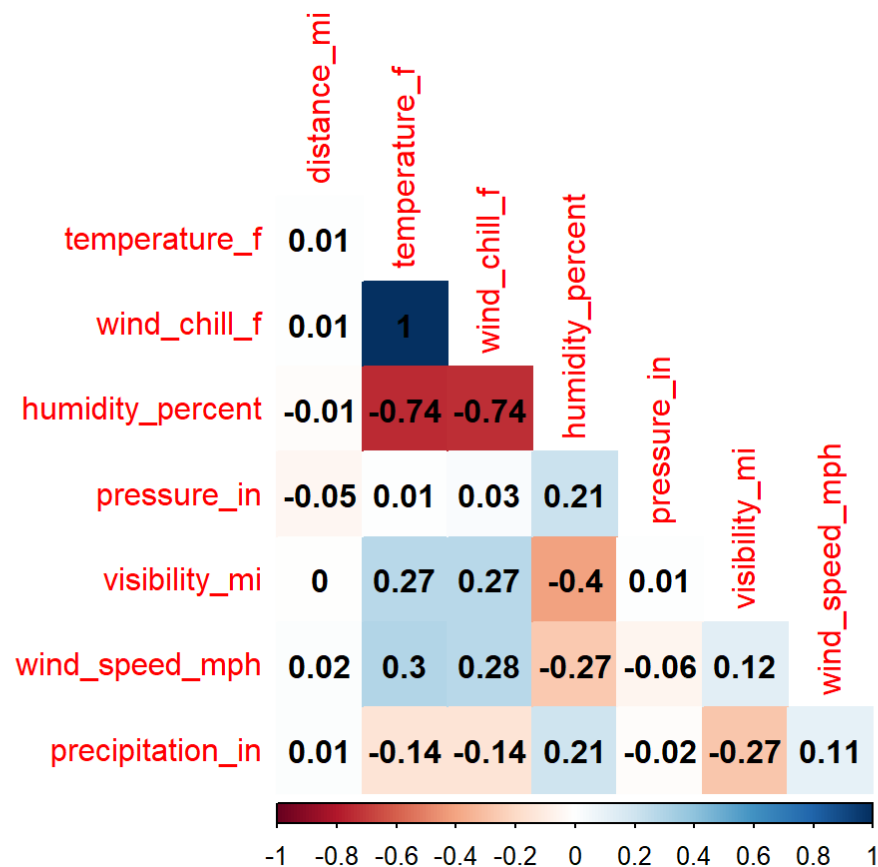
Code



The first thing we can notice already is the that crossing, traffic signals, and stop related traffic incidents are much more frequent than bumps and railway accidents. Therefore, we might be able to say that since there is a lower amount of railway and bump accidents in both categories, it might be two variables that won't probably be too important in determining the accidents. Additionally, just using logic, it would make sense that these two factors are something that doesn't seem exclusive to severe accidents as well. Being a daily driver myself, most accidents just generally tend to stray away from bumps and railways as they aren't populated parts of the road.

## Potential Predictor Correlation

[Code](#)

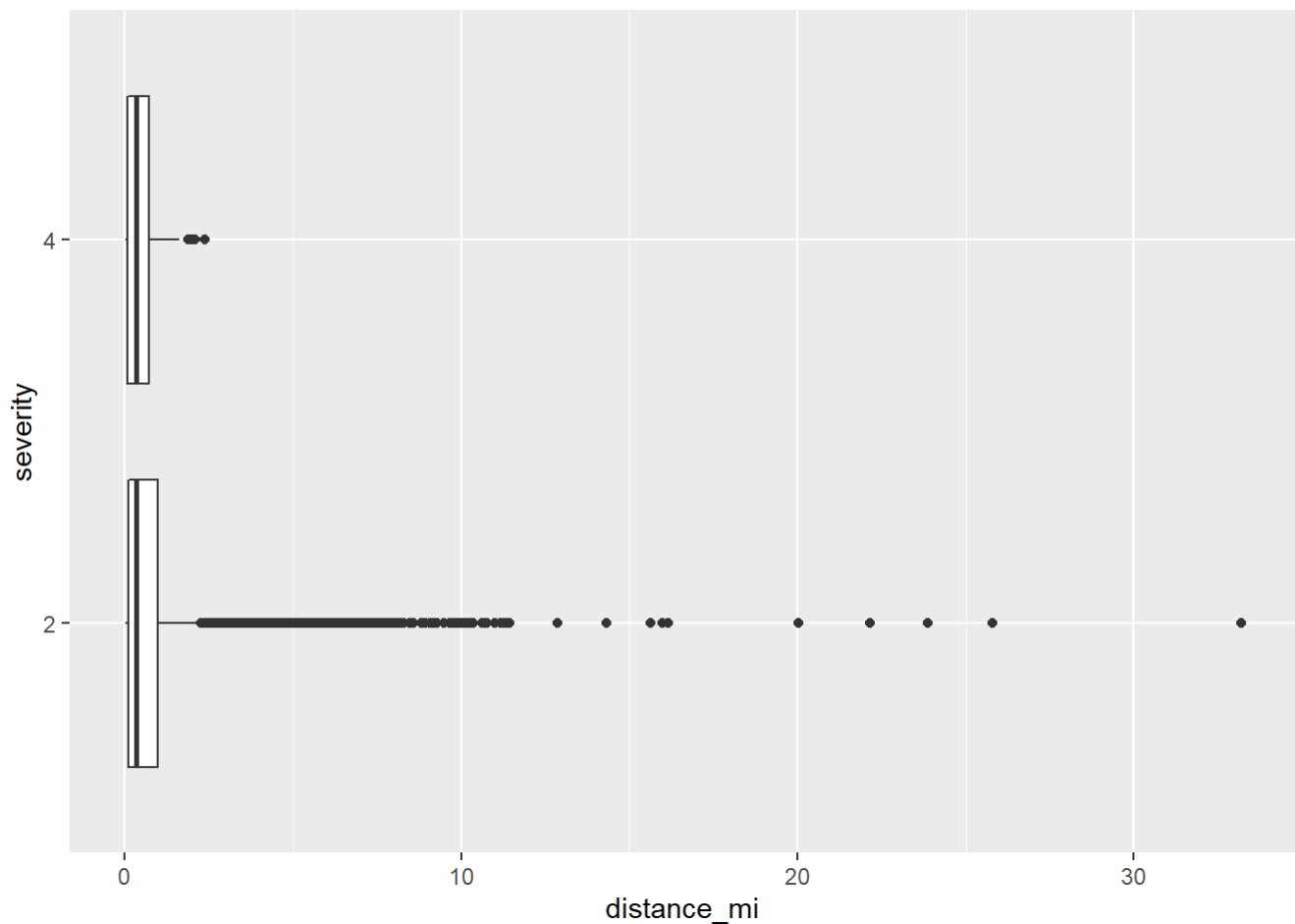


I wanted to create a correlation matrix to find any potential correlations between the numeric variables before proceeding with the analysis. I believe this is important so we can take our model with a grain of salt and maybe interpret the data differently depending on what variables are “significant.” Already, we see that temp/wind\_chill are 100% correlation which is pretty self explanatory, as wind temperature will increase at overall temperature increases. Similar with the highly negative correlated wind\_chill/humidity as we’d expect that humidity increase at the temperature increases, so the wind would be less chilly. Though what I find interesting is that humidity/temperature are negatively correlated, I’d think that humidity increases at temperature increase. Otherwise, as you can see, distance isn’t quite correlated with anything, which is a bit odd since that’s essentially what we are checking for. The severity of the accident, in terms of traffic block up (distance of traffic stopped being one of them potentially). So it seems that the numeric weather factors don’t seem to be all too correlated so far, which might be something to keep in mind when actually running our logistic model to find any significant variables. As for the other variables, nothing seems to be too exceptional in terms of correlation that is specifically noteworthy.

## Road Sidedness and distance\_mi/visibility\_mi

Code

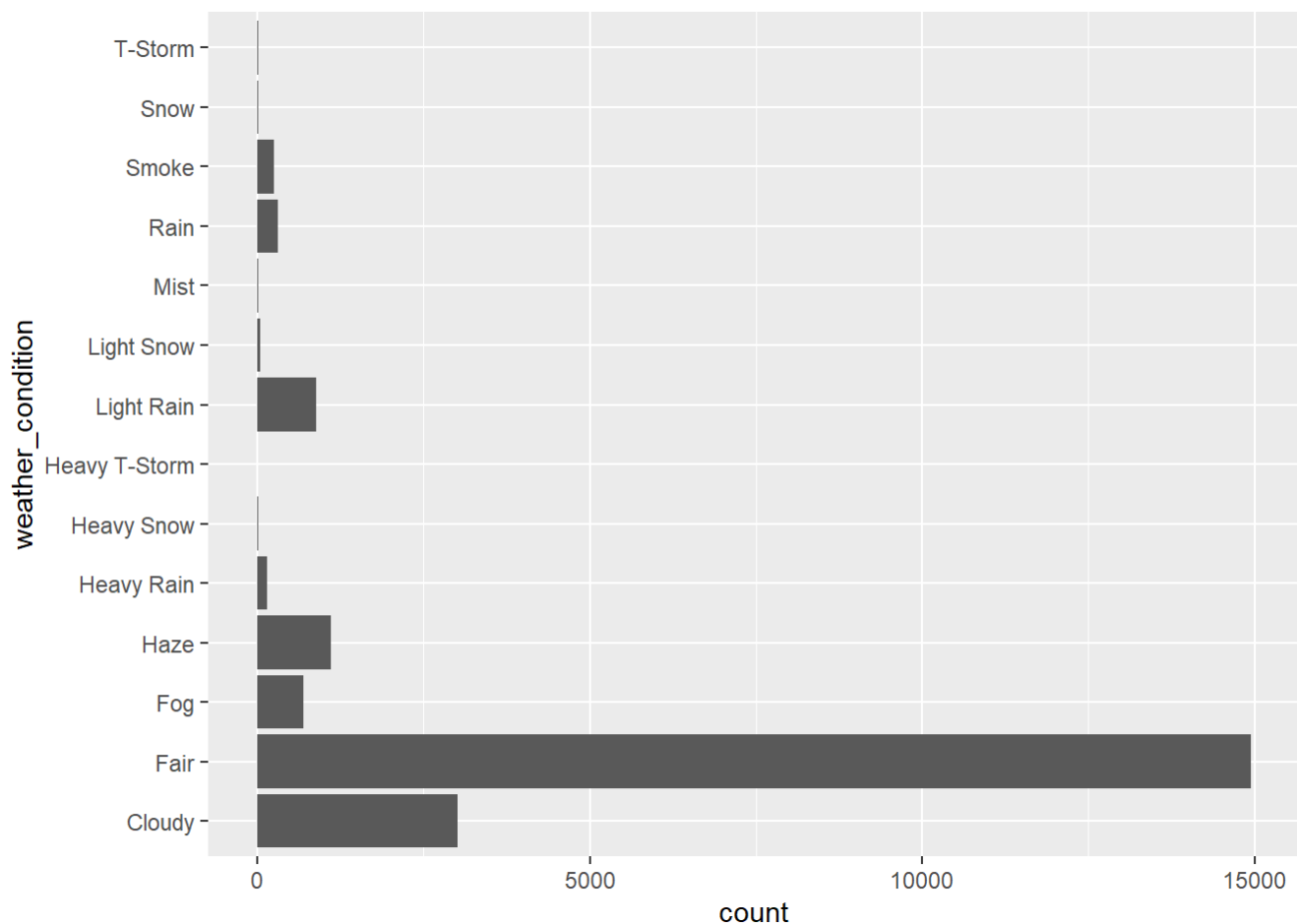




As I noted previously, the severity of an accident is based largely on how much of traffic the accident ends up holding up. Therefore, we'd have to notions that we'd think that a longer distance of traffic that's held up by the accident would mean a higher severity and they would be correlated. We actually see that there appears to be no correlation and there are pretty extreme outliers in the severity 2 category, which might affect how if our models consider distance as significant or valuable, but remember, this is only a preliminary check and distance can certainly still be a significant variable.

# Weather conditions

Code

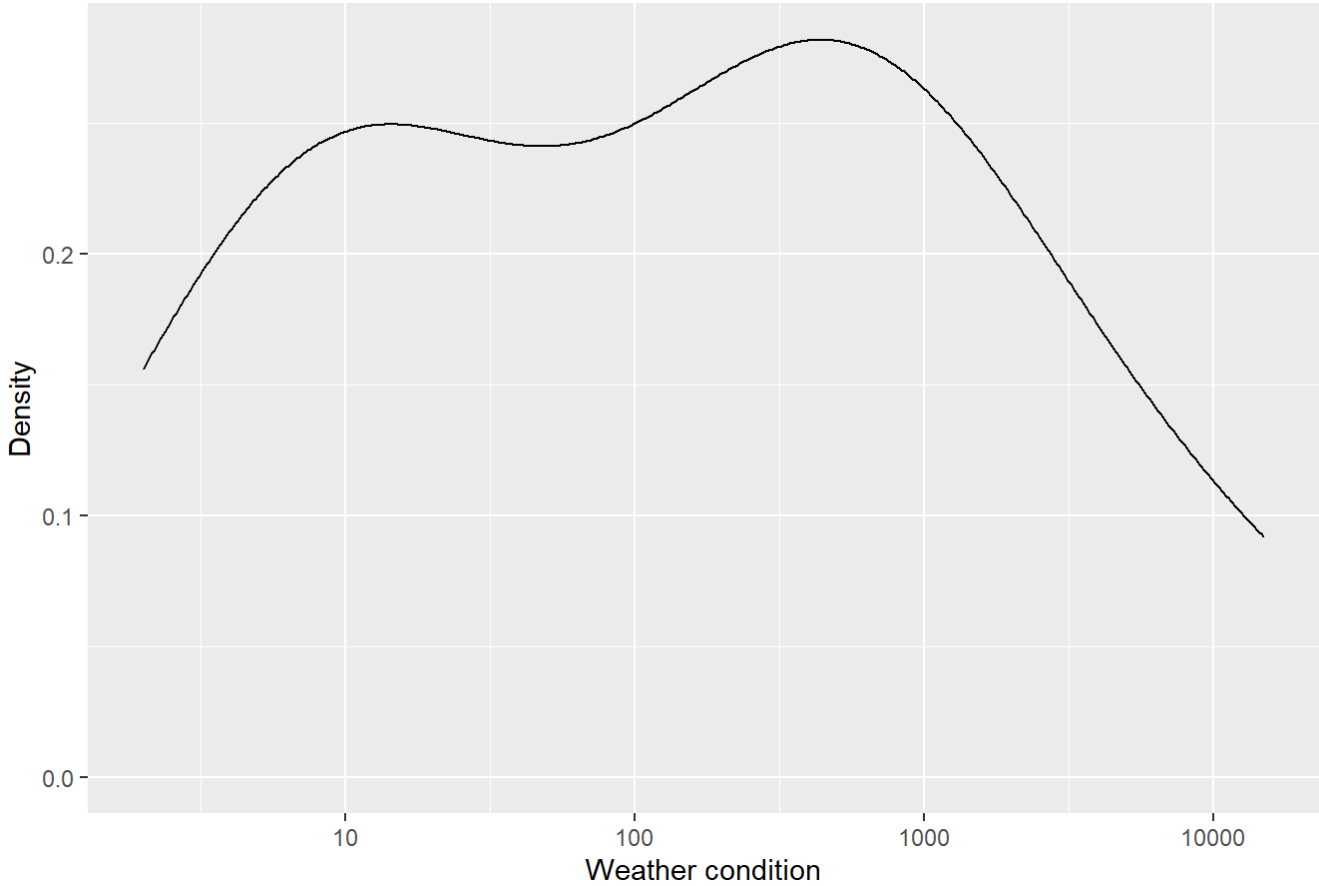


This graph helps display what's considered to be the most abundant weather conditions at the time of the accident. This is important to display because abundances/large disparities between the appearance of certain weather conditions might highly skew the models ability to predict weather condition as valuable predictors. Additionally, large disparities in certain data values can mean it wont yield an accurate representation of what's actually true, so therefore printing this is valuable because we should take our models conclusion with a grain of salt until we can get more data on low counts of certain conditions, such as mist, t-storm, snow, heavy t-storm, mist, and heavy snow.

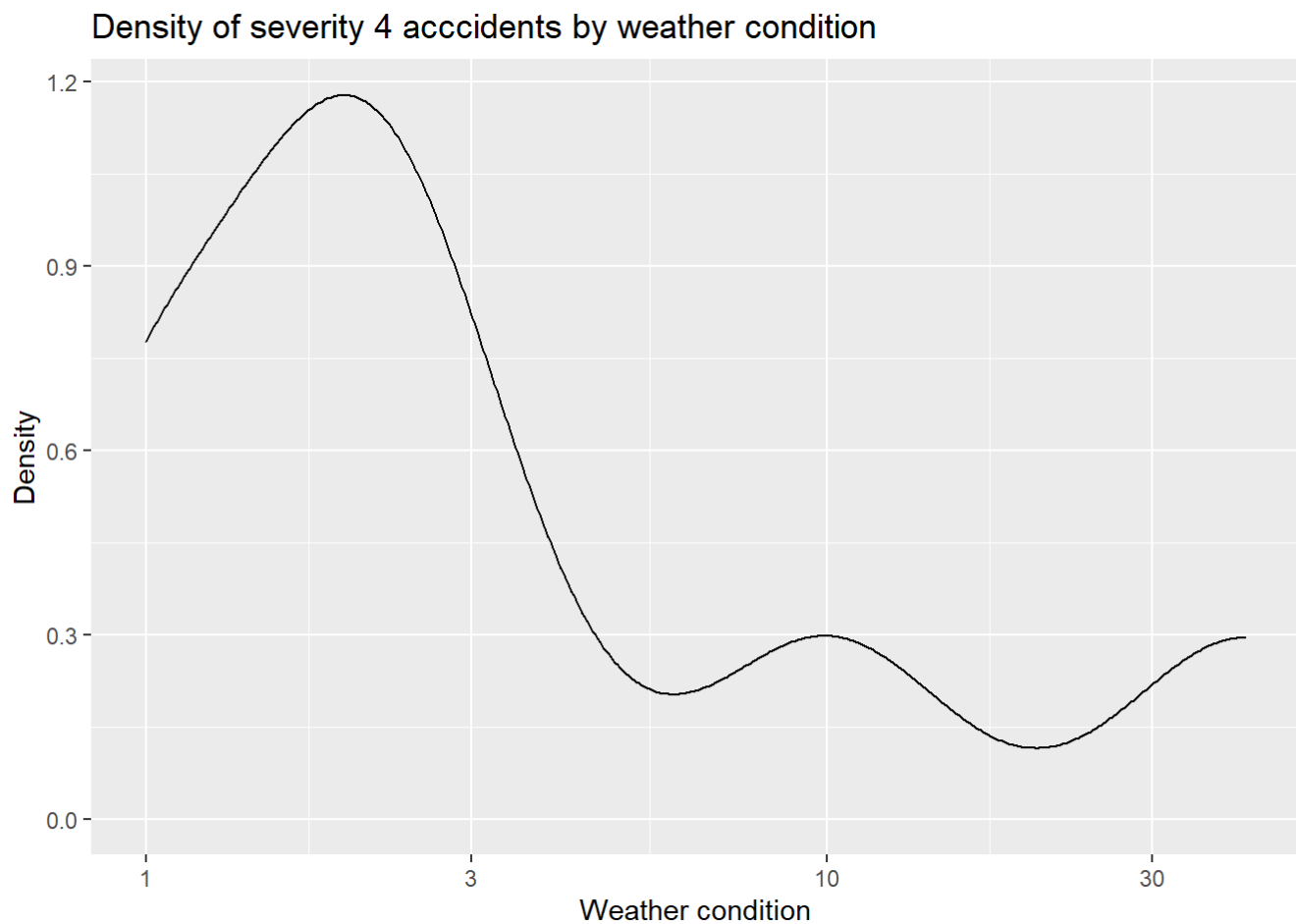
## Severity and Weather Condition

[Code](#)

Density of severity 2 acccidents by weather condition



Code



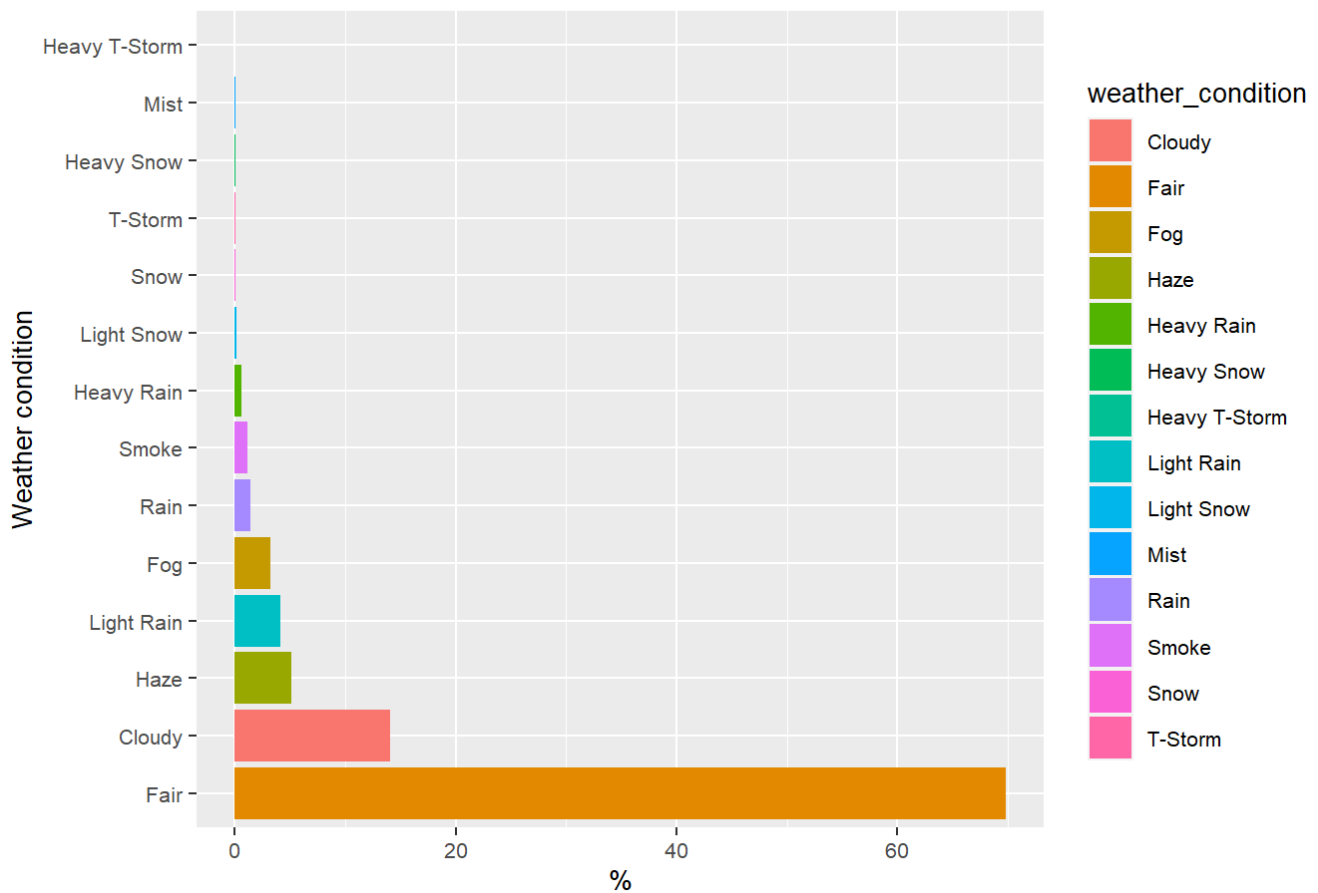
Essentially, what this graph is telling is that as the amount of different weather conditions increase in our data set, there are fewer and fewer severity 4 accidents. Basically, there are only a few weather conditions that actually account for severity 4 accidents. But on the other hand, there are quite a few weather conditions that account for our severity 2 accidents. Meaning that severity 4 has little diversity in weather conditions that describe the incidents, but there are a lot more weather condition diversity in severity 2 accidents which may be a good in using weather conditions to determine what the accident was. We can visual this better with the graphs below.

[Code](#)

```
## # A tibble: 14 × 2
##   weather_condition    count
##   <fct>              <dbl>
## 1 Fair                69.8
## 2 Cloudy              14.0
## 3 Haze                5.16
## 4 Light Rain          4.12
## 5 Fog                 3.25
## 6 Rain                1.41
## 7 Smoke              1.18
## 8 Heavy Rain          0.655
## 9 Light Snow          0.192
## 10 Snow               0.0796
## 11 T-Storm            0.0421
## 12 Heavy Snow         0.0375
## 13 Mist               0.0281
## 14 Heavy T-Storm      0.00936
```

[Code](#)

Severity 2 accidents by weather condition

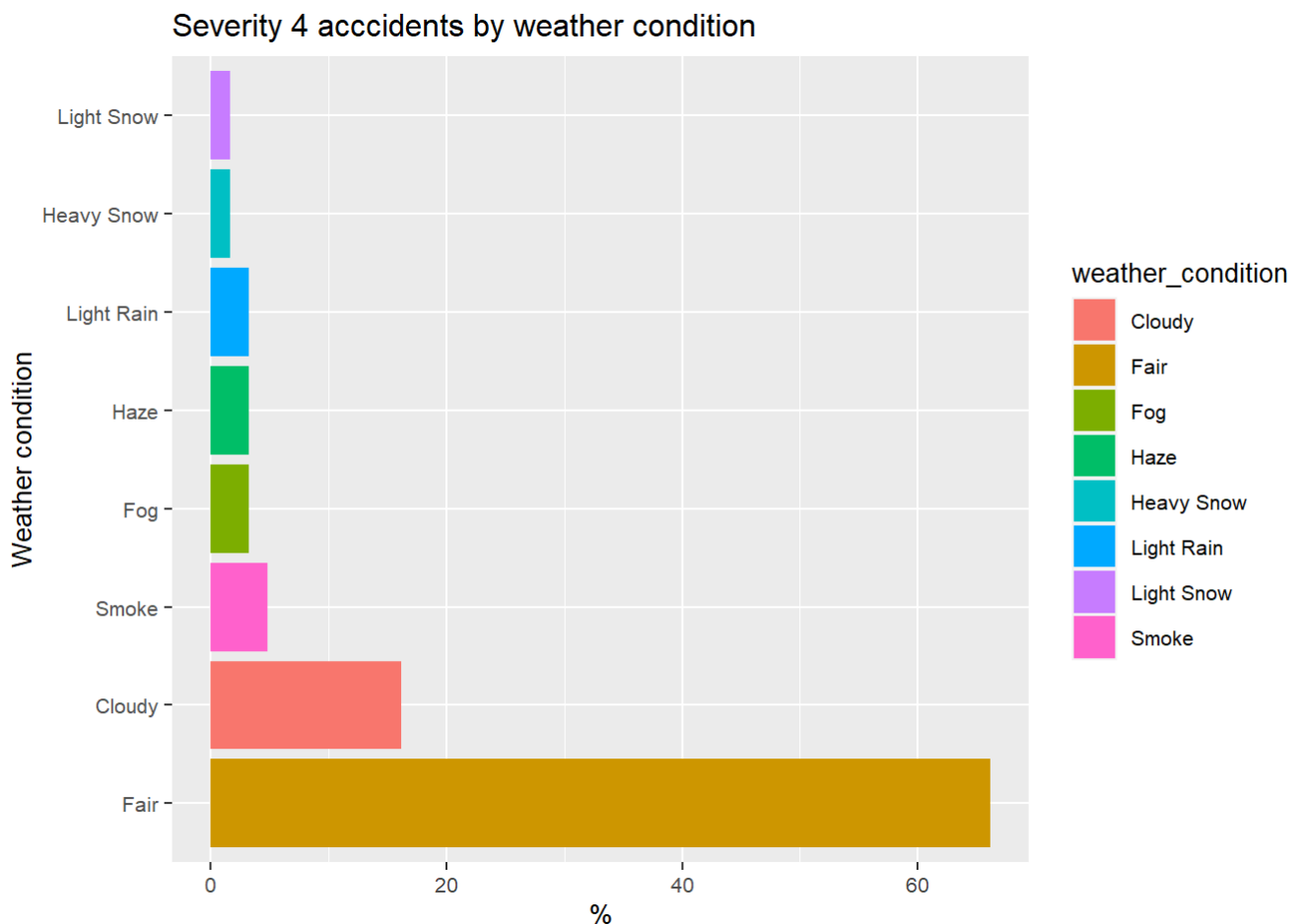


In this graphic, we numerically graph the types and amount of weather conditions in severity 2 incidents. Notice that there are 14 weather conditions that happened in severity 2 accidents, which is essentially all of the accident types.

[Code](#)

```
## # A tibble: 8 × 2
##   weather_condition count
##   <fct>             <dbl>
## 1 Fair              66.1
## 2 Cloudy            16.1
## 3 Smoke             4.84
## 4 Fog               3.23
## 5 Haze              3.23
## 6 Light Rain        3.23
## 7 Heavy Snow        1.61
## 8 Light Snow        1.61
```

Code



We build a similar model for our severity 4 accidents and you can already notice a difference. First, it's important to note that the weather condition "fair" is most abundant in both severities which make sense since this is California only data, and California typically doesn't experience an abundance of weather conditions. But what's unique of the two severities is that severity 4 accidents only cover 7 different weather conditions. Again, this can be due to a lack of severity 4 representation in the data which is worth nothing, but nonetheless, it's something that is very apparent between the two.

## Model Building

Now that we have taken some idea of the distributions of our data, it is time to build our recipe's and models to hopefully achieve some predictive powers in determining "predictive" variables in predicting severity of California accidents, and to potentially find what's a good telling sign of what might be valuable in predicting what finds an accident's severity.

## Recipe, Folds, and Saving the Data

[Hide](#)

```
accidents_folds <- vfold_cv(accidents_train, v=5, repeats = 4, strata=severity)

accidents_recipe <- recipe(severity ~ ., data = accidents_train) %>%
  step_dummy(side) %>%
  step_dummy(weather_condition) %>%
  step_dummy(bump) %>%
  step_dummy(crossing) %>%
  step_dummy(railway) %>%
  step_dummy(stop) %>%
  step_dummy(traffic_signal) %>%
  step_dummy(sunrise_sunset) %>%
  step_normalize(all_predictors()) %>%
  step_upsample(severity, over_ratio = 0.50)

save(accidents_folds, accidents_recipe, accidents_train, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/model_setup.rda")
```

- Firstly, I create only 5 folds since we have such an abundance of observations, but to compensate for the lower value of folds, I will increase repeats to 4. And again, due to our class unbalance, I must make sure to stratify our data by severity.
- Next, I will dummy encode all of our factor variables
- Normalize the predictor variables
- Finally, as I noted before, a valuable way to deal with the class imbalance is to upsample. This means I will randomly sample with replacement our under-represented class until it matches about 50% (in my case) of the more abundance class. The reason for doing this is because if we don't upsample, then our model will continually predict severity 2 because that's the only data that the model is trained to predict since there's almost no appearances of severity 4 accidents, meaning that the models predictive power will be extremely low.

## Preparing the Models

The models I'll be using are:

- logistic regression
- single decision tree
- boosted tree
- random forests
- k-nearest neighbors

Below, all my models were tuned to 4 levels as too many more levels would greatly increase run-time and computing power as 4 levels already took over ~4 hours to run due to my high number of observations, so 4 levels should be enough especially since I have 4 repeats.

## logistic regression

[Hide](#)

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

log_wkflow <- workflow() %>%
  add_model(log_reg) %>%
  add_recipe(accidents_recipe)

save(log_wkflow, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/log_model.rda")
```

## single decision tree

[Hide](#)

```
tree_spec <- decision_tree() %>%
  set_engine("rpart")

class_tree_spec <- tree_spec %>%
  set_mode("classification")

class_tree_wf <- workflow() %>%
  add_model(class_tree_spec) %>%
  set_args(cost_complexity = tune()) %>%
  add_recipe(accidents_recipe)

class_tree_param_grid <- grid_regular(cost_complexity(range = c(0,1), trans = identity_trans()), levels = 4)

class_tree_tune_res <- tune_grid(
  class_tree_wf,
  resamples = accidents_folds,
  grid = class_tree_param_grid)

save(class_tree_wf, class_tree_tune_res, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/class_tree_model.rda")
```

## random forests

[Hide](#)



```
spec <- rand_forest() %>%
  set_engine("ranger", importance = "impurity")

rf_spec <- spec %>%
  set_mode("classification")

rf_spec_wf <- workflow() %>%
  add_model(rf_spec %>% set_args(mtry = tune(), trees = tune(), min_n = tune())) %>%
  add_recipe(accidents_recipe)

rf_param_grid <- grid_regular(mtry(range = c(1,16)), trees(range = c(100,500)), min_n(range
= c(1,16)), levels = c(4,4,4))

rf_tune_res <- tune_grid(
  rf_spec_wf,
  resamples = accidents_folds,
  grid = rf_param_grid)

save(rf_spec_wf, rf_tune_res, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fittin
g/rf_model.rda")
```

## boosted tree

Hide

```
spec <- boost_tree() %>%
  set_engine("xgboost")

boost_spec <- spec %>%
  set_mode("classification")

boost_spec_wf <- workflow() %>%
  add_model(boost_spec %>% set_args(mtry = tune(), trees = tune(), min_n = tune())) %>%
  add_recipe(accidents_recipe)

boost_param_grid <- grid_regular(mtry(range = c(1,16)), trees(range = c(100,500)), min_n(ran
ge = c(1,16)), levels = c(4,4,4))

boost_tune_res <- tune_grid(
  boost_spec_wf,
  resamples = accidents_folds,
  grid = boost_param_grid)

save(boost_spec_wf, boost_tune_res, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_f
itting/boost_model.rda")
```

Again, I tuned both my boosted trees and random forest to similar hyperparameter ranges because the trees are relatively similar.

## K-nearest Neighbors

[Hide](#)

```
spec <- nearest_neighbor() %>%
  set_engine("kknn")

knn_spec <- spec %>%
  set_mode("classification")

knn_wkflow <- workflow() %>%
  add_model(knn_spec %>% set_args(neighbors = tune())) %>%
  add_recipe(accidents_recipe)

knn_param_grid <- grid_regular(neighbors(range = c(1,1000)),levels = 4)

knn_tune_res <- tune_grid(
  knn_wkflow,
  resamples = accidents_folds,
  grid = knn_param_grid)

save(knn_tune_res, knn_wkflow, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/knn_model.rda")
```

The only parameters I tuned were neighbors, and it's generally safe that neighbors is tuned to the square root of the number of observations (in my case, 30k). Therefore, a range from 1-1000 is pretty safe. I set the engine to kknn to denote k-nearest neighbors and set the mode to classification. And I saved my workflow

## Running the Models

### Loading Data

[Hide](#)

```
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/model_setup.rda")
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/log_model.rda")
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/class_tree_model.rda")
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/rf_model.rda")
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/boost_model.rda")
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/knn_model.rda")
```

### Logistic

[Code](#)

```
## # A tibble: 29 × 5
##   term                estimate std.error statistic  p.value
##   <chr>              <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)        -1.86      2.20     -0.848 3.97e- 1
## 2 distance_mi        -0.200    0.0195   -10.2   1.70e-24
## 3 temperature_f       0.905    0.216     4.18  2.86e- 5
## 4 wind_chill_f        -1.08    0.213    -5.08  3.78e- 7
## 5 humidity_percent   -0.0784   0.0263   -2.98  2.88e- 3
## 6 pressure_in         0.0255    0.0164    1.56  1.19e- 1
## 7 visibility_mi       -0.144    0.0274   -5.27  1.40e- 7
## 8 wind_speed_mph      -0.117    0.0177   -6.62  3.60e-11
## 9 precipitation_in    -0.284    0.0522   -5.44  5.40e- 8
## 10 side_R             0.0527    0.0150    3.50  4.58e- 4
## # ... with 19 more rows
```

From initial view, we see that distance\_mi, temp\_f, wind\_chill, humidity, visibility, wind\_speed, precipitation, side, haze, heavy and light snow, crossing, railway, stop, traffic\_signal, and sunrise\_sunset all appear to be significant predictors with a p-value < 0.05. Again, but our data is extremely skewed as we noted in our EDA, so this might lead to some problems of predictions and it might draw conclusions due to lack of data even despite upsampling. Therefore, let's find the accuracy of our model and create a confusion matrix below to better/simply visual our data.

Code

```
## # A tibble: 1 × 3
##   .metric .estimator .estimate
##   <chr>   <chr>      <dbl>
## 1 accuracy binary      0.982
```

We see that the our logistic model is extremely accurate, but how true is this? Let's create the confusion matrix.

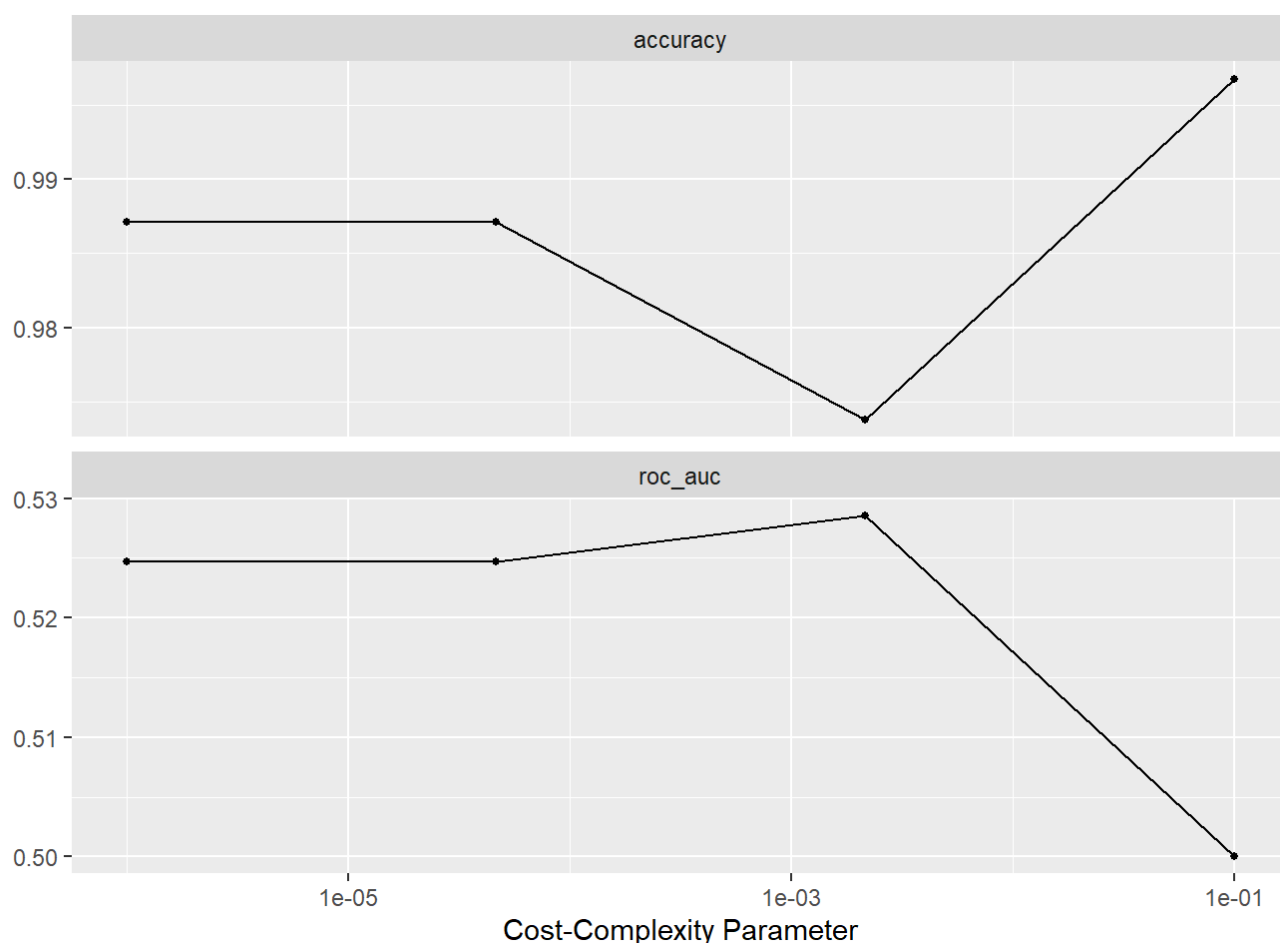
Code

```
##           Truth
## Prediction    2    4
##           2 24047    74
##           4   358    5
```

Ahhh, now we know why our logistic model is accurate. Since logistic accuracy the model's ability to predict the right thing it'll obviously be extremely accurate since there are so many severity 2's to predict. But from our confusion matrix we actually see that the model is extremely bad at predicting severity 4's and more often than not, it'll wrongly predict or never predict severity 4's. This means logistic might be a poor model to run despite it's high accuracy, and although it might be a linear model, our data is over-represented and the simplicity of logistic models might not be able to capture the differences due to lack of data. So we should probably perform different models below.

## Single Decision Tree

Plot of accuracy and roc\_auc

[Code](#)

The first thing we should note is that our roc\_auc is extremely poor no matter what our cost-complexity is. Even though our accuracy is increasing, it might be for the same reason that we are continually predicting severity 2's and not 4's so therefore our accuracy increase although that's not too helpful. Therefore, we can't accurately predict severity 4's and ultimately, our roc\_auc decreases as a result. A hover roc\_auc means that our model is as good as coin flipping if it's severity 2 or 4, so therefore, using the model wouldn't be quite useful.

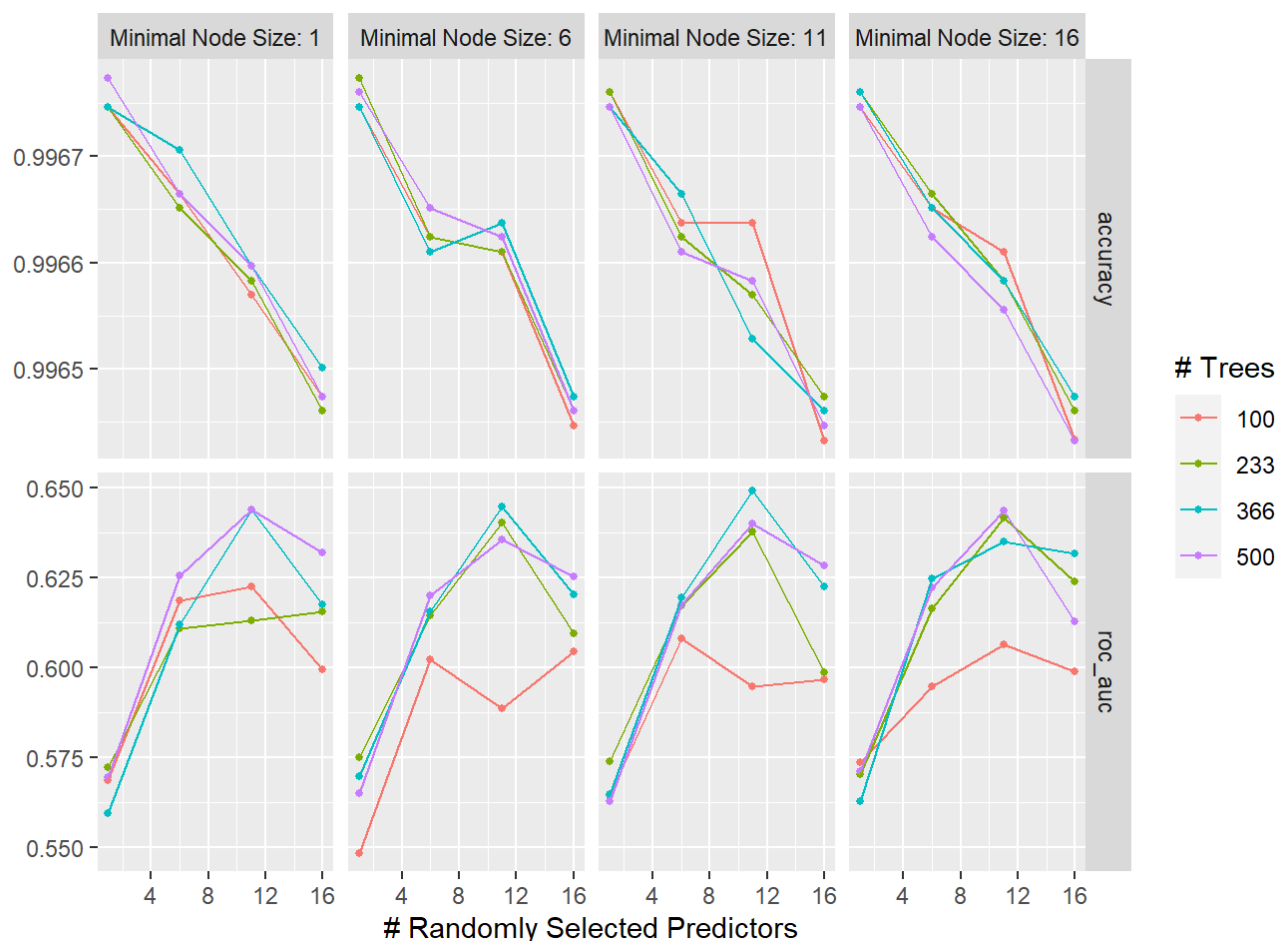
### Best Performing Model

[Code](#)

```
## # A tibble: 4 × 7
##   cost_complexity .metric .estimator  mean    n std_err .config
##         <dbl> <chr>    <chr>    <dbl> <int>  <dbl> <chr>
## 1      0.00215  roc_auc  binary    0.529   30 0.0108 Preprocessor1_Model3
## 2      0.000001  roc_auc  binary    0.525   30 0.00974 Preprocessor1_Model11
## 3      0.0000464  roc_auc  binary    0.525   30 0.00974 Preprocessor1_Model12
## 4       0.1      roc_auc  binary    0.5     30 0       Preprocessor1_Model14
```

## Random Forest

Maybe performing more in-depth models such as random forest and boosted trees might end up being more valuable and predictive, so let's see the results of both below.

[Code](#)

We can see that depending on our hyper-parameters, accuracy is not quite an issue because although it seems to dip heavily, there really isn't much difference in accuracy as we get a >99% accuracy every single time. Now, let's analyze the roc\_auc. Already we see that our roc\_auc for random forest models is doing MUCH better than our logistic and single decision tree, which might be expected due to their limitations and flexibility. Our random forest generally yields an roc\_auc of around ~0.63-0.65, meaning that our models are not too bad and this might be a noteworthy model when testing our test data.

For the most part, we see that there seems to be a declining success after around 11 predictors, meaning that there seems to be 5 predictors that are pretty invaluable for our model. Additionally, having our tree's above 233 generally don't increase our roc\_auc by all too much so any amount of trees after this point don't seem to produce any groundbreaking changes.

But what I find interesting is that I only limited the minimum node size from 1-16, which relative to having a data set with 30k observations is quite low. That means that we create trees with 1-16 minimum observations before moving onto the next node. A larger minimum node amount means that a smaller tree is built, and we see that 1 node doesn't seem to differ much when compared with larger minimum node values so our trees are overall pretty flexible. But we will explore this further in a later section.

Best Performing model

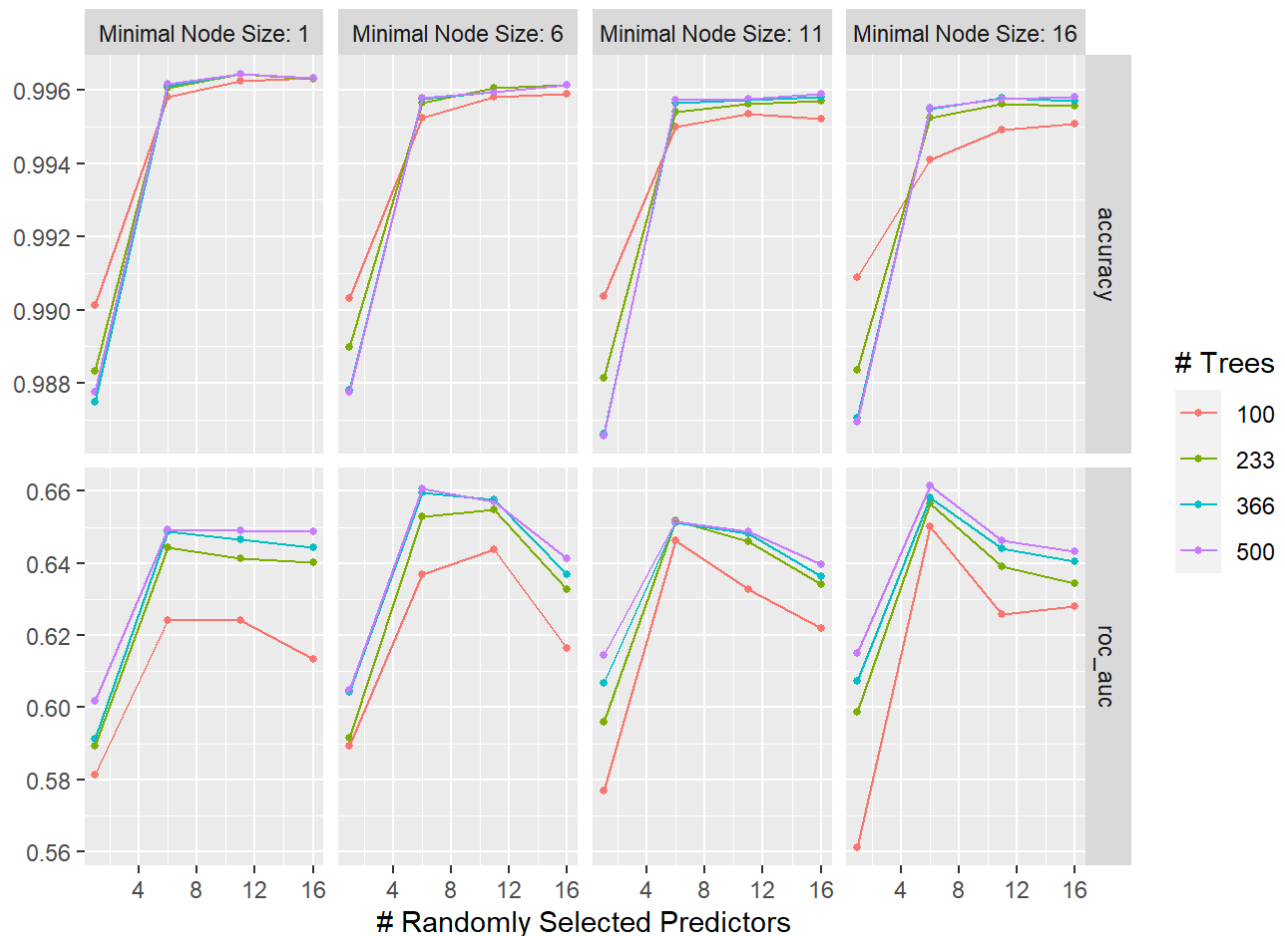
[Code](#)

```
## # A tibble: 5 × 9
##   mtry trees min_n .metric .estimator mean      n std_err .config
##   <int> <int> <int> <chr>   <chr>   <dbl> <int>   <dbl> <chr>
## 1    11    366     11 roc_auc binary    0.649    30  0.0219 Preprocessor1_Model143
## 2    11    366      6 roc_auc binary    0.645    30  0.0173 Preprocessor1_Model127
## 3    11    366      1 roc_auc binary    0.644    30  0.0182 Preprocessor1_Model111
## 4    11    500      1 roc_auc binary    0.644    30  0.0196 Preprocessor1_Model115
## 5    11    500     16 roc_auc binary    0.643    30  0.0223 Preprocessor1_Model163
```

As I noted, since there doesn't seem to be a big difference in increase node size, and a flexible model is possible, then perhaps we can try lowering our node range in a different mode using our best hyper-parameters in terms of trees and mtry. We see that a minimum node of 1 also produces pretty good roc\_auc, therefore we can create a range that's around 1 and maybe see if a min\_n around there might find better roc\_auc/accuracy.

## Boosted Tree

Code



Compared to random forest, we already see that generally our model is performing better in terms of roc\_auc, even if by a little. Also compared to random forest, our sweet spots for predictors are around 6 predictors while the amount of trees seem to produce a pretty negligible difference in roc\_auc. And what's even more interesting, while random forest accuracy seemed to continually decrease when adding more

predictors, it seems that after 6 predictors variables, our accuracy actually increase while roc\_auc decreases slightly. Again, this can maybe be attributed to over-predictions of severity 2's, but we can draw a confusion matrix later to see if that's the case.

Best Performing model

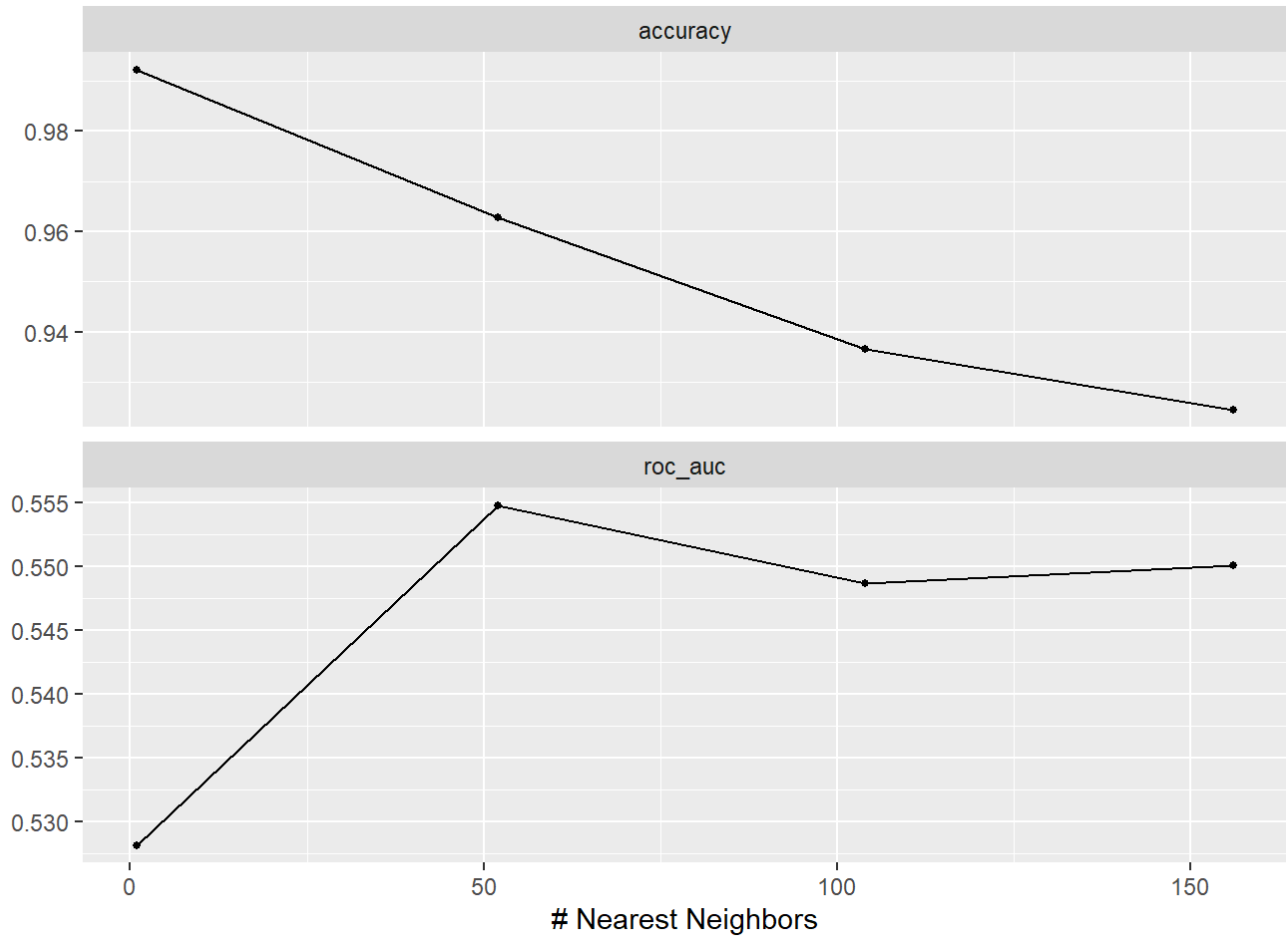
Code

```
## # A tibble: 5 x 9
##   mtry trees min_n .metric .estimator  mean      n std_err .config
##   <int> <int> <int> <chr>   <chr>    <dbl> <int>   <dbl> <chr>
## 1     6   500    16 roc_auc binary    0.661    30  0.0201 Preprocessor1_Model132
## 2     6   500     6 roc_auc binary    0.661    30  0.0165 Preprocessor1_Model124
## 3     6   366     6 roc_auc binary    0.660    30  0.0162 Preprocessor1_Model123
## 4     6   366    16 roc_auc binary    0.658    30  0.0200 Preprocessor1_Model131
## 5    11   366     6 roc_auc binary    0.658    30  0.0201 Preprocessor1_Model139
```

Since boosted trees are similar to random forest, we will do the model fitting by seeing if a change in the range of min\_n around the best min\_n will produce a better roc\_auc. Especially because we see a min\_n of 6 and 16 generally produce the best roc\_auc.

K-nearest Neighbors

Code



One thing to note is that k-nearest neighbors actually perform poorly when it comes to large datasets, high number of dimensions, and many outliers, and with a class imbalanced, large data set, and from our eda, we found distance\_mi had many outliers, and finally with over 17 predictors, we can already assume this model won't be too significant.

And follow our assumptions, we see that our roc\_auc is quite low, with almost being similar to the coinflip probabilities of our decision tree model. It also has a pretty notable continuous decrease in accuracy as the amount of neighbors increase.

Since this is a model that isn't particularly impressive, there's no need to show the best performing tuned parameters since we won't be using it to fit our testing set.

## Finding New Models

Above we ran some models with tuned parameters in the forest and boosted models. Notice we had a pretty large minimum node range, this can mean that we actually couldn't capture what's exactly happening at certain node levels since we can't see the minute difference that might have happened at lower ranges.

Also, we see the models actually performed pretty well at lower minimum nodes too, meaning that our model is actually quite flexible as it can take in one observation's minimum for each node and still perform well in terms of roc\_auc. Therefore, maybe we can decrease the range of our min\_n around those runner-up values in our best performing mtry and trees values to see how this might produce a more precise story within those ranges and if it at all will maybe show us if those ranges are better since we can see more details with a lower range. So we will change the grid and hypertune the parameters differently to see if any changes could be made.

## Setting up the grid of hyperparameters

Hide



```

# Random Forest

spec <- rand_forest() %>%
  set_engine("ranger", importance = "impurity")

rf_spec <- spec %>%
  set_mode("classification")

min_n_rf_spec_wf <- workflow() %>%
  add_model(rf_spec %>% set_args(mtry = 11, trees = 366, min_n = tune())) %>%
  add_recipe(accidents_recipe)

min_n_rf_param_grid <- grid_regular(min_n(range = c(1,3)), levels = 4)

min_n_rf_tune_res <- tune_grid(
  min_n_rf_spec_wf,
  resamples = accidents_folds,
  grid = min_n_rf_param_grid)

save(rf_spec_wf, min_n_rf_tune_res, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/min_n_rf_model.rda")

# Boosted Tree

spec <- boost_tree() %>%
  set_engine("xgboost")

boost_spec <- spec %>%
  set_mode("classification")

min_n_boost_spec_wf <- workflow() %>%
  add_model(boost_spec %>% set_args(mtry = 6, trees = 500, min_n = tune())) %>%
  add_recipe(accidents_recipe)

min_n_boost_param_grid <- grid_regular(min_n(range = c(5,7)), levels = 4)

min_n_boost_tune_res <- tune_grid(
  min_n_boost_spec_wf,
  resamples = accidents_folds,
  grid = min_n_boost_param_grid)

save(boost_spec_wf, min_n_boost_tune_res, file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/min_n_boost_model.rda")

```

As I noted before, I want to fit models around a `min_n` that performed almost as good as our best model so we can maybe see a more detailed story of what might be happening at those closer ranges. For the random forest, this will be around 1-3, and for boosted trees, around 5-7. Let's print the results below.

## Printing out the Graphs

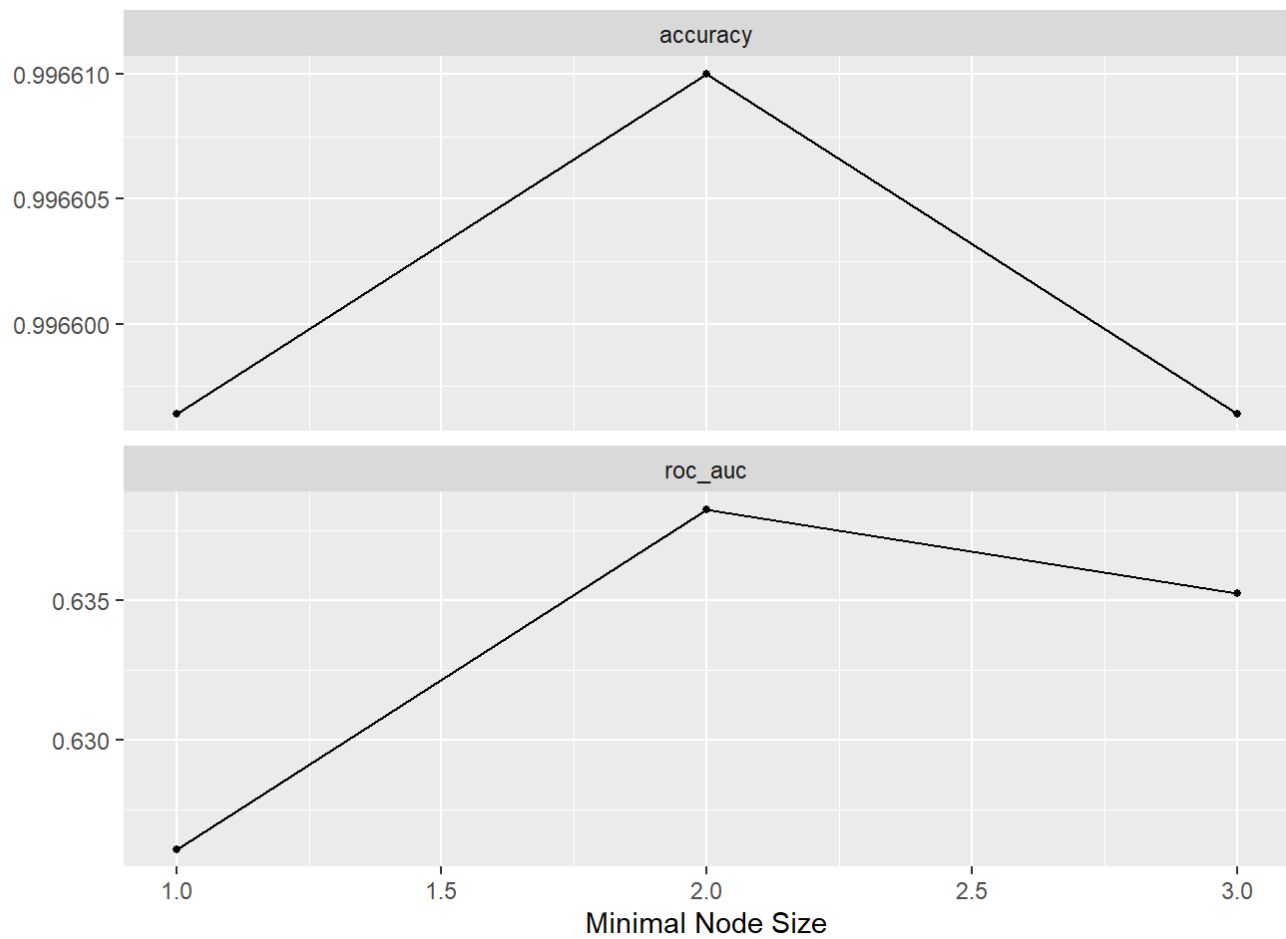
Loading data

Hide

```
# Loading the data  
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/min_n_rf_model.rda")  
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/min_n_boost_model.rda")
```

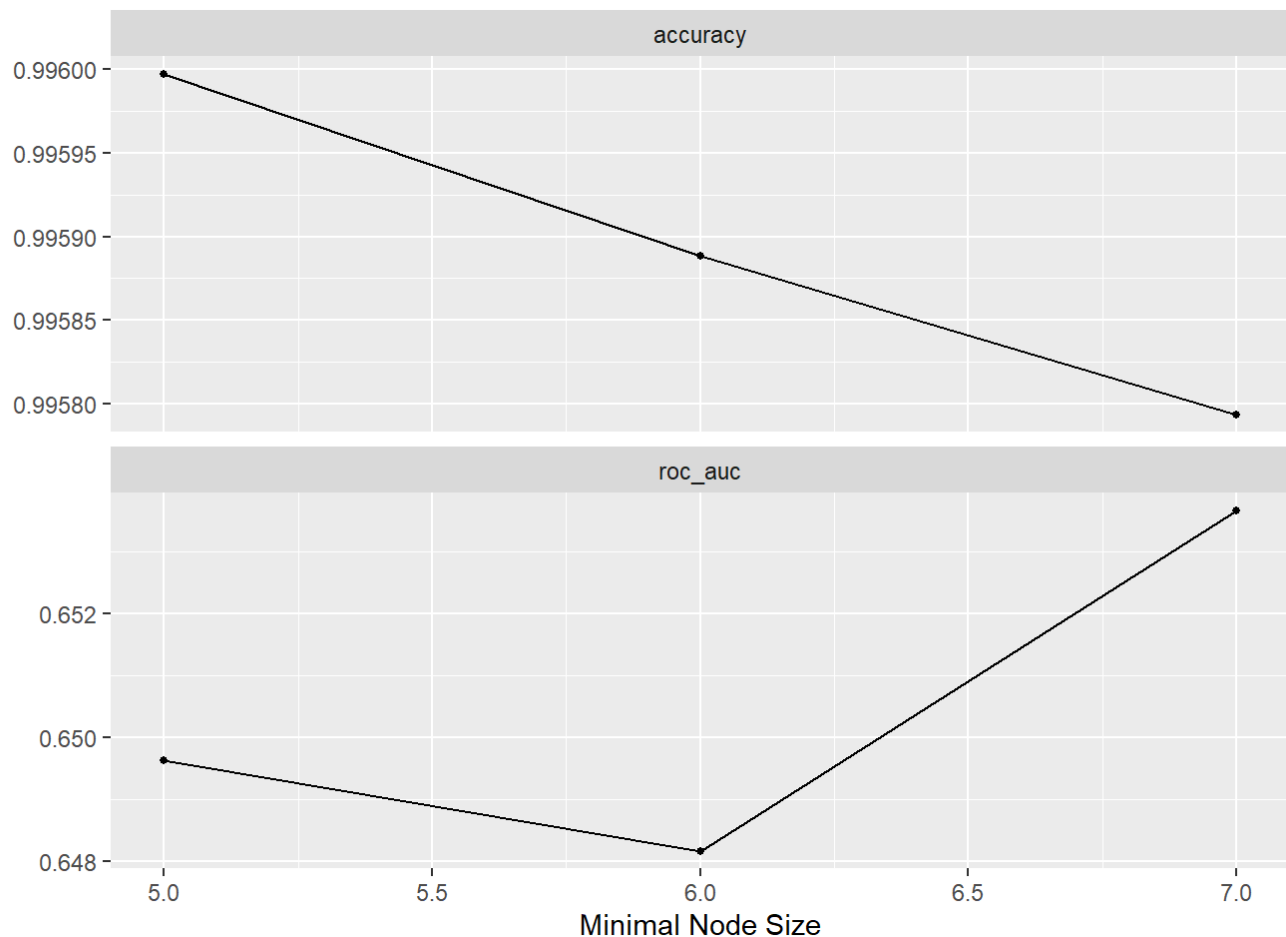
New min\_n range Random Forest

Code



New min\_n range Boosted Tree

Code



In conclusion, we see that although there might be some increase in roc\_auc near ~7 and beyond for min\_n for our boosted tree models, our accuracy continuously decreases and roc\_auc is still lower than our best performing model so it might be in our best interest to just maintain our best hyper parameters previously of mtry = 5, trees = 500, and min\_n = 16 to maintain an equal balance of accuracy and roc\_auc. As for the random forest model, although a min\_n of two isn't too bad, notice that 11 actually yields our best results of accuracy combined with roc\_auc. So for the most part, it might just be the safest option to also revert back to our original best hyper parameters as well since we have confirmed it might be the best working model within our original and tighter min\_n range doesn't seem to produce much difference.

## Fitting the Testing Set

Ultimately, we determined that it's in our best interest to just use our previously best tuned hyperparameters found initially, therefore we will just stick with those when fitting the testing set.

## Setting up the best final workflows

Since we determined that the boosted trees overall yields the best roc\_auc while maximizing accuracy, we will fit our testing data with this model.

Hide

```
boost_final <- finalize_workflow(boost_spec_wf,select_best(boost_tune_res,metric = "roc_auc"))

boost_final_result <- fit(boost_final, data = accidents_train)

save(boost_final_result,file = "D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/best_model.rda")
```

## Displaying Graphics

loading data

Hide

```
load("D:/UCSB/PSTAT131/Final Project/R_Scripts/model_fitting/best_model.rda")
```

roc\_auc value of best boosted tree Model on test data

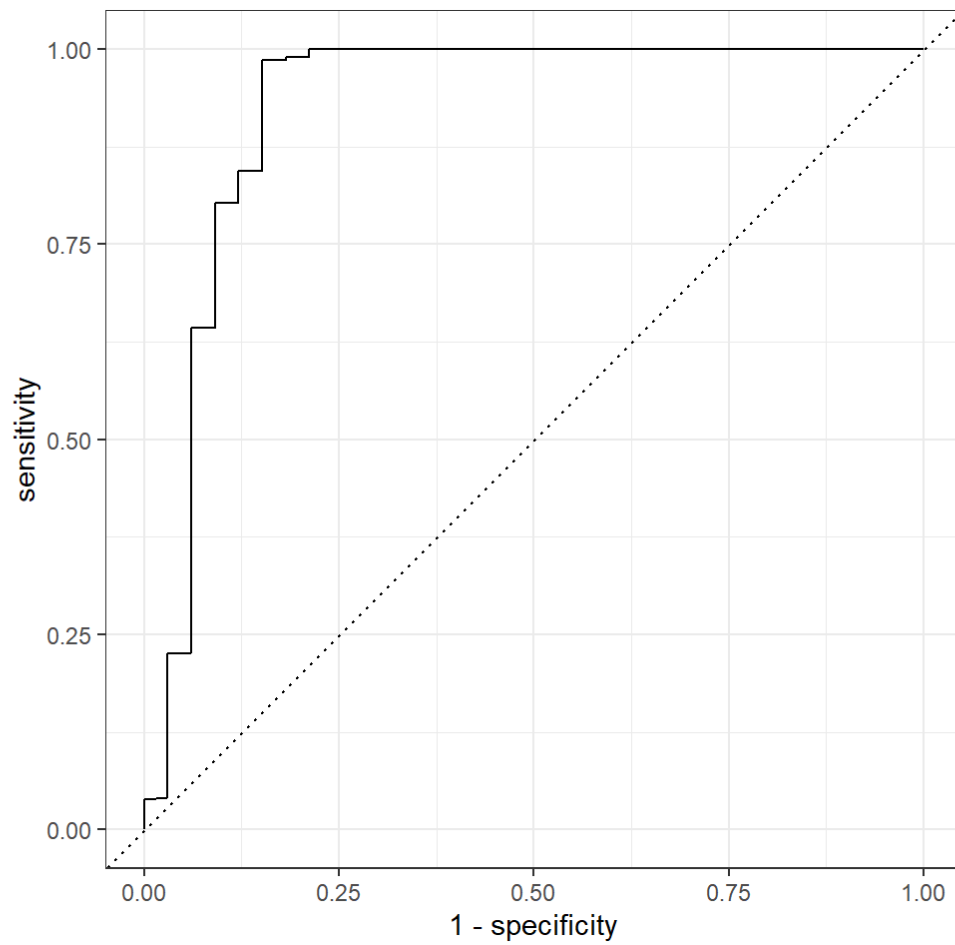
Code

```
## # A tibble: 1 × 3
##   .metric .estimator .estimate
##   <chr>   <chr>       <dbl>
## 1 roc_auc binary      0.925
```

Notice that overall, when it comes to predicting our severity 2 accidents, we actually have an extremely high roc\_auc. This is predicted since we have an abundance of severity 2 values, so lets see the roc\_auc for severity 4 accidents. Wow! Our roc\_auc is very high and initially it may seem like our model is excellent at predicting severity 4. But let's not take the number for granted as it could've just been overfitted to predict severity 2, so it'll be in our best interest to see how well our model did based on looking at a heat matrix.

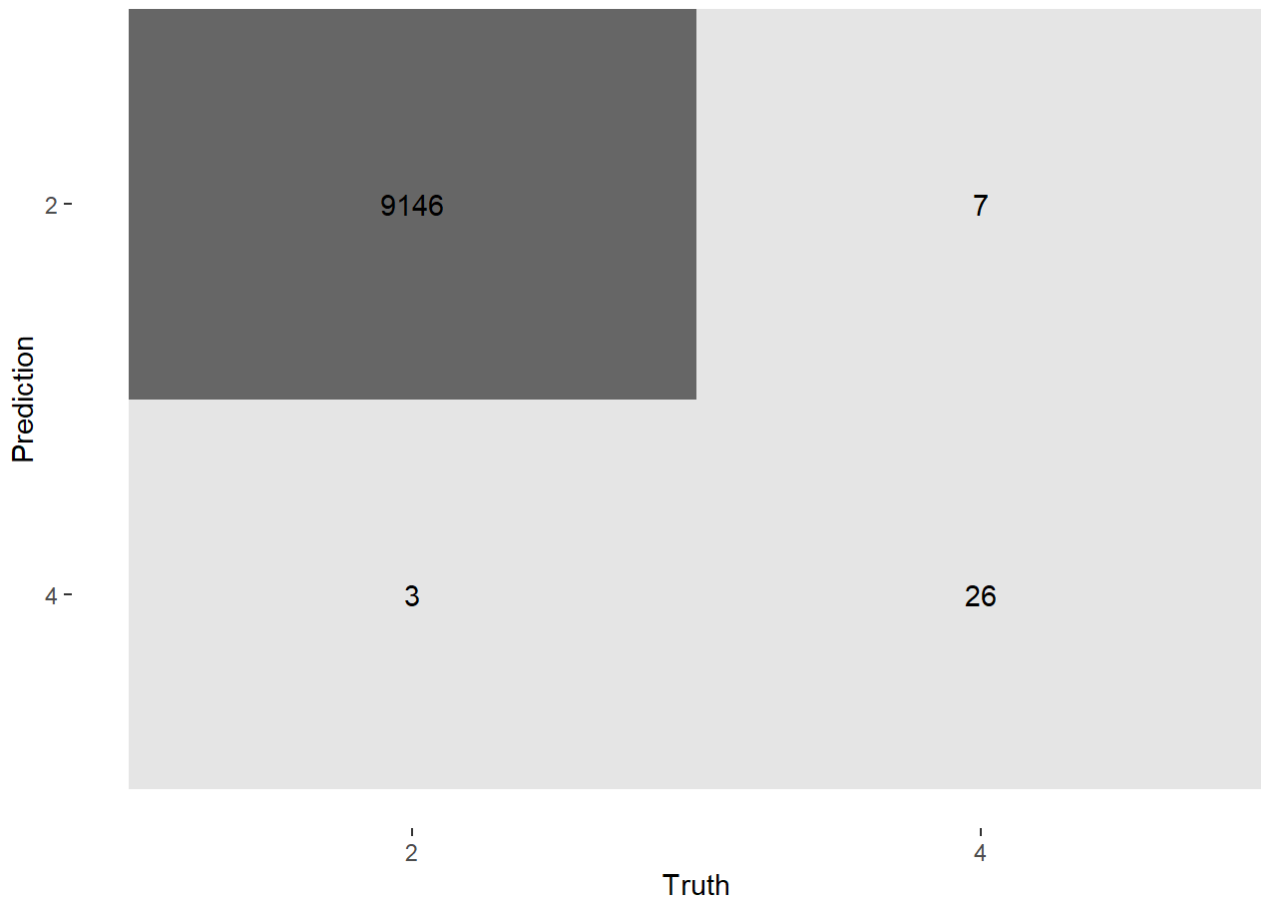
roc\_auc curve of best boosted tree Model on test data

Code



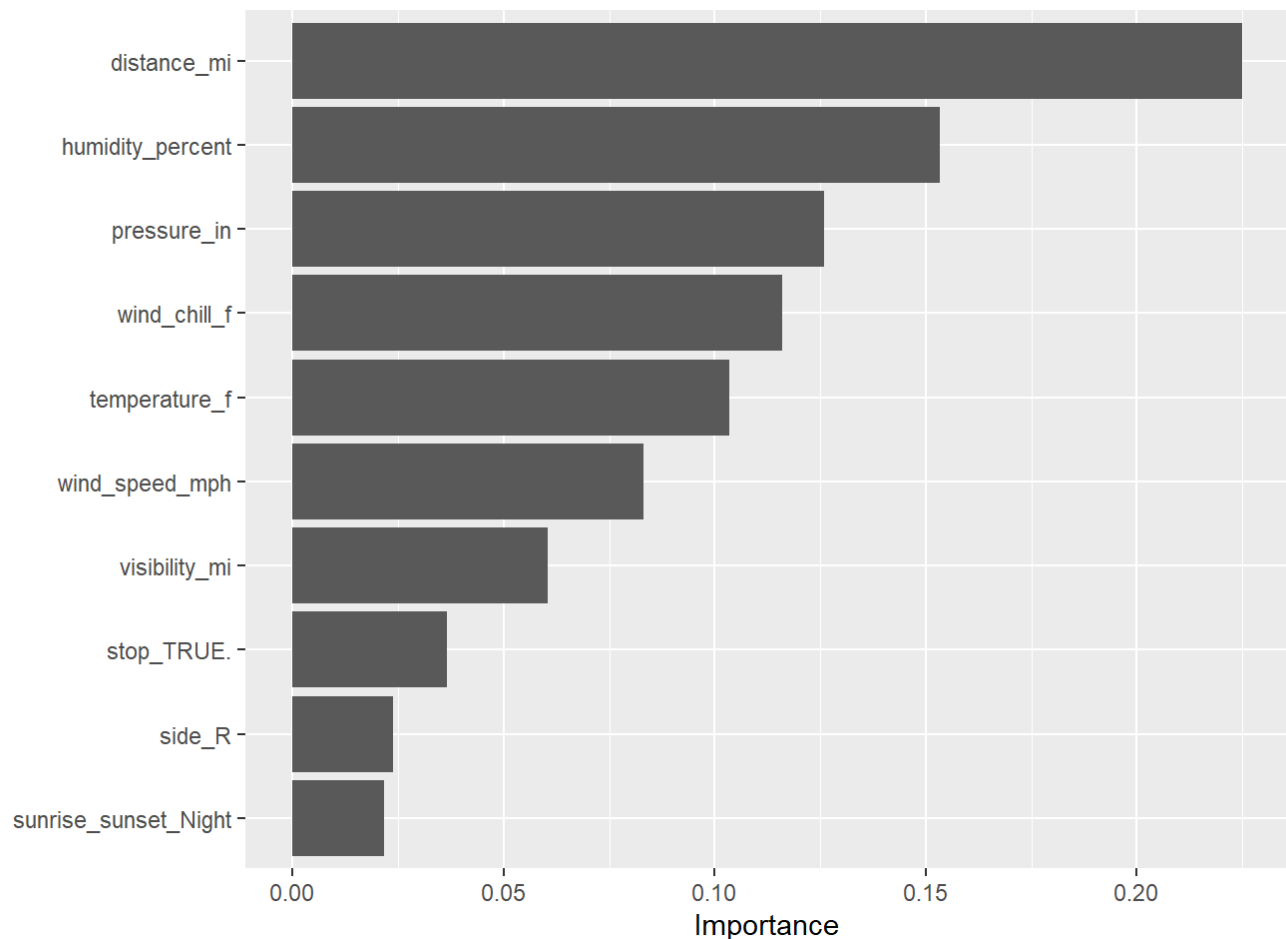
The graphic above is just the roc\_auc curve for visualization of how well our model fitted the the testing data.

Code



Looking at our heat matrix, we actually see our model did quite well in predicting severity 4 despite the major disparity in severity 4 accidents. It only predicted severity 4 wrong 3 times, while accurately predicting it right 26 times. And as expected, it did a great job in predicting severity 2, which may be due to the obvious fact that there's an abundance of severity 2 observations as I've continually noted.

[Code](#)



It also might be in our best interest to see what the most valuable predictors were in our boosted tree model. Our variable importance chart tells us that distance\_mi was by far the best predictor of severity which was expected since we'd expect a more severe accident to block more of traffic of course. After we see that humidity\_percent, pressure\_in, wind\_chill\_f, temperature\_f were all pretty significant predictors which is actually extremely surprising. All of them being important also may be due to the fact that they were all already initially significantly correlated so potentially running interactions in the recipes would be worth modeling in the future for further accuracy perhaps. Then we see that stop signs, sidedness, and sunrise\_sunset had decently/low importance. Stop signs might be significant because those are typically where intersections exist so more traffic would be held up as a result of this, and potentially more severe accidents would happen in the day time since more traffic exists during those hours, but what was surprising is road sidedness which I'd think wouldn't be too important.

## Conclusion and Final Remarks

### Regarding the Training data models