Kaggle Report Submission - Classification Data

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Introduction

For this project, we referenced a dataset that provides information about the 2020 US Presidential Election (Biden versus Trump). This dataset includes data from 3,111 counties; 75% (2,331 rows) allocated for training and the remaining 25% (780 rows) used for testing. The training set contains 126 variables, such as total population, population by gender, population by age, population by race, and population by education level. The response variable is 'winner,' which is a categorical variable that indicates the winning candidate of the county.

Looking at the voting trends from the article "Behind Biden's 2020 Victory," we see that voting preferences were influenced by demographics and education levels. For instance, Trump had high support from white men without college education, while Biden had more support from women and those with a college degree. Additionally, the ages of individuals was another factor, with younger voters tending to be Democratic and older voters leaning towards Republican. Therefore, we believe that the predictors as mentioned above are significant in predicting the 'winner' for each county.

References:

https://electionlab.mit.edu/

https://data.census.gov/

https://www.pewresearch.org/politics/2021/06/30/behind-bidens-2020-victory/

https://www.ppic.org/wp-content/uploads/jtf-immigrants-political-engagement.pdf

https://www.pewresearch.org/politics/2024/04/09/partisanship-by-gender-sexual-orientation-marital-and-parental-status/

EDA



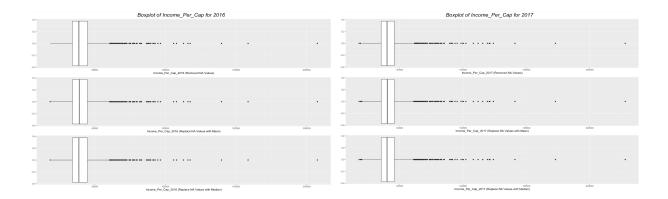
Figure 1

From this model, we can note that the Trump class has a lot more observations than the Biden class. Since there is a large number of observations overall, we should consider downsampling the data to preserve model accuracy.



Figure 2

After downsampling the data using the step_downsample() function, we can note that the proportion of observations in each class is much more balanced. Now that the data is more balanced, our models will be able to better learn from each class.



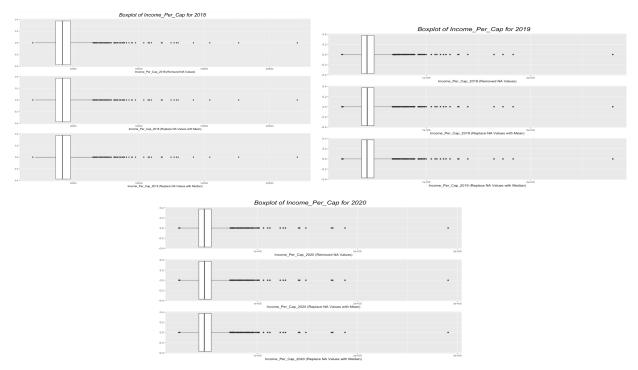


Figure 3

Because income_per_cap data had NAs, we wanted to see if filling the NAs with the median or mean of its column would drastically change the income_per_cap column's distribution. From this, we can see that either option does not change the data drastically.

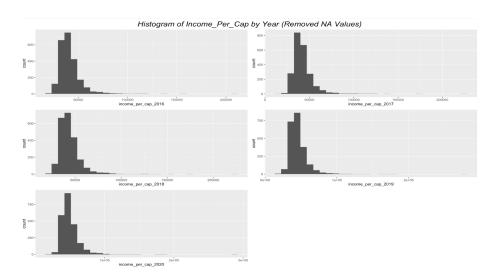


Figure 4

From this model, we can note that the income_per_cap variable tends to be more right-skewed. So, it would be a better choice to use median in order to prevent any outliers from drastically impacting our model accuracy.

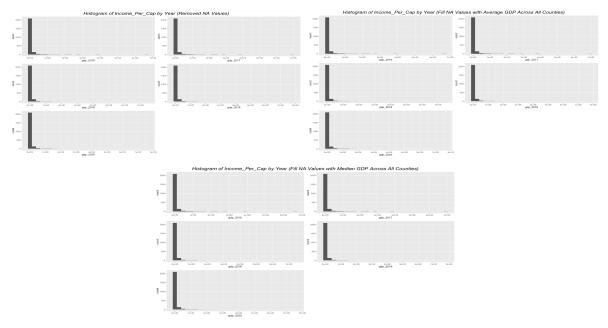


Figure 5

From this model, we noticed that imputing the mean or median for the NA values did not create a large difference in each of the GDP's variable distributions. Since the GDP variable appears to be right skewed, we ended up choosing to impute these values with the median as well.

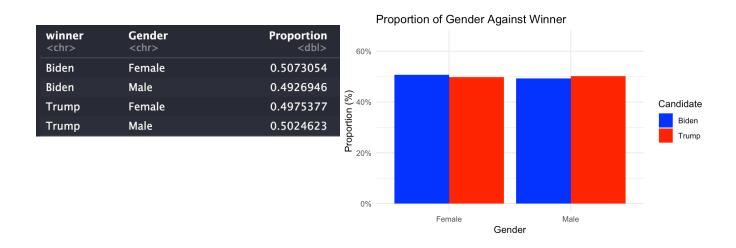


Figure 6

We plotted the proportion of gender against the winner to determine if gender impacts voting preferences. The results show that the counties favoring Biden have slightly more female voters, whereas Trump has more male voters. The difference is marginal, suggesting that gender as a predictor might not be sufficient.

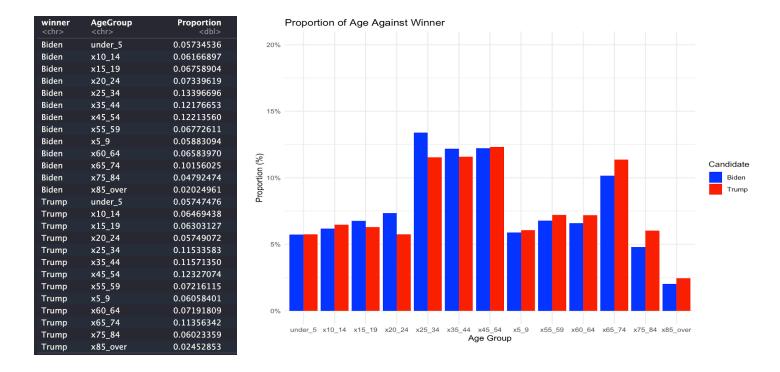


Figure 7

We considered whether age predicts the winner of a county. The bar plot shows that people under 44 tend to vote for Biden, while older populations vote for Trump. Counties with younger age groups favor Biden, indicating that age is a significant predictor.

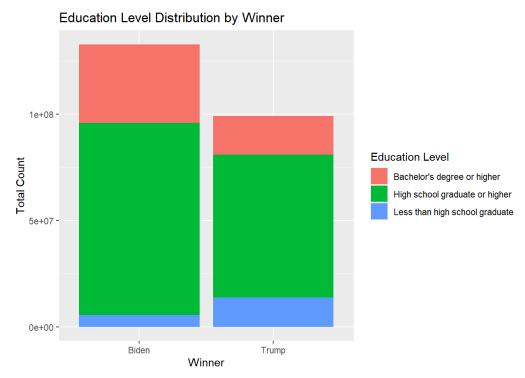


Figure 8

Another consideration was whether voters' education levels would affect their voting patterns. This stacked bar chart shows that districts in favor of Biden had more voters with Bachelor's degrees or higher whereas districts in favor of Trump had more voters whose education levels were less than high school graduates.

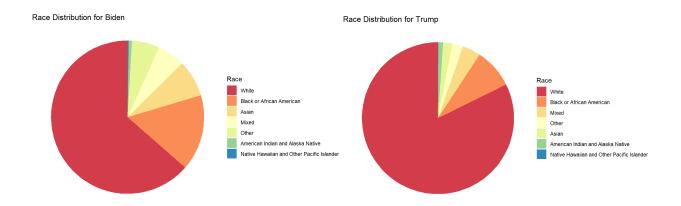


Figure 9

Amongst Biden and Trump voters, white voters make up the largest demographic, then African Americans, Asians, Mixed, Other, American Indian and Alaskan Native, and then Native Hawaiian and Other Pacific Islanders. However, Biden had more votes among minority populations than white voters.

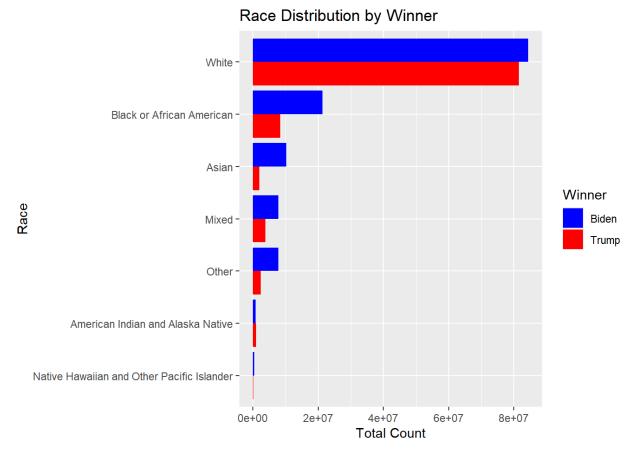


Figure 10

Expanding on figure 9, we wanted to visualize how racial demographics compare between districts. Trump and Biden had a similar amount of white voters. However, Biden had 3-4 times more voters amongst minority races than Trump. Thus, the proportion of minority races amongst voters could be an important predictor.

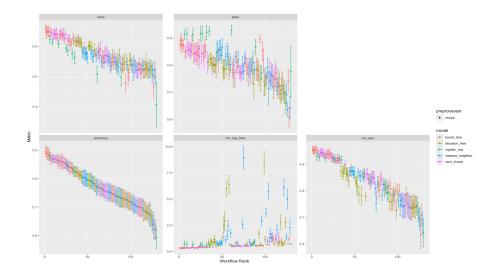
Preprocessing + Recipes

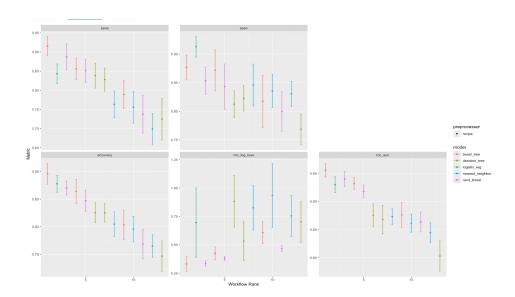
From figures 3-5, we decided to replace the NA values of the income_per_cap and gdp columns with the median of that column. Then, we removed the "x0033e", "x0036e", "x0058e" variables because they were a repeat of the "x0001e", "x0034e", and "x0035e" columns. Additionally, the "id" and "name" columns were removed since they are not predictor variables. Lastly, we noticed in figure 1 that there was an imbalance in the winner class. So, we used the step_downsample() function to create a more balanced dataset. Then, we applied the prep() and juice() recipe to prepare and apply the recipe to the training data and return the downsampled dataset, which would be used in the following recipes.

For recipes, we created a base recipe, filter recipe, and a pca recipe. For the base recipe, we set the winner variable as the response variable and the remaining variables as the predictor. We also created 7 additional variables: prop_white (proportion of people who are White), prop_minority (proportion of those who are Asian, Black or African American, American Indian and Alaska Native, Hispanic, Native Hawaiian, and/or some other race), prop_college (proportion of those who graduated with a Bachelor's degree or higher), prop_old (proportion of those who are 65+), prop_ men (proportion of men), prop_female (proportion of female), and prop_minor (proportion of those that are younger than 18). To avoid duplicates, we removed variables that correspond to the original counts of these percentages; for instance, because we have prop_white, we should not keep the variable that has the count of people who are white. Lastly, we factored the last x2013_code variable and applied step_dummy() since this variable should be treated as a categorical variable, not a numerical variable; we had to apply step_dummy() because some models would not allow for a factor predictor.

For the filter variable, we added the step_corr() function with a tunable threshold to the base recipe. For the pca recipe, we started with the base recipe. Then, we had R filter out any variables with near zero variance using the step_nzv() function to get rid of any unnecessary predictors. We also added the step_corr() filter on all numeric predictors (minus the outcomes variable) to eliminate any highly correlated predictors. We also added the step_lincomb() function to remove any linear combinations of other predictors. We then added the step_normalize() function on all numeric predictors (minus the outcome variable) because PCA assumes that the data is already normally distributed. Lastly, we applied the step_pca() function with a tunable threshold and num_comp = 5 so that R can try to reduce the number of features in this dataset.

We then created a workflow set using these 3 recipes and 5 models (boosted tree, decision trees, logistic regression, k nearest neighbor, and random forests). After removing the pca_glmnet and filter_glmnet, we then called the workflow map. After evaluating this workflow set on our 10-fold cross validation with a grid of 10, we called autoplot to visualize all the different types of models and ranked the overall results.





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mple_knn mple_knn mple_knn a_xp\text{shoots} \$1 \text{ of 65 rows} \$2 \text{ of 65 rows} \$2 \text{ of 65 rows} \$3 \text{ of 65 rows} \$4 \text{ of 65 rows} \$5 \text{ of 65 rows}	Preprocessor I, Model03 Preprocessor I, Model I Preprocessor I I Preprocessor I I Preprocessor I I I P	roc_auc sens spec accuracy .metric . <chr> .<chr> mn_log_loss roc_auc sens spec accuracy mn_log_loss roc_auc sens spec accuracy cacuracy cacuracy sens spec</chr></chr>	0.8731248 0.7634278 0.8457490 0.8032726		10 recipe 10 rec	<int> <int> 10 10 10 10 10 10 10 10 10 10 10 10 10</int></int>	nearest_neighbor nearest_neighbor nearest_neighbor boost_tree preprocessor <chr> <chr> <chr> cchr> recipe recipe recipe</chr></chr></chr>	model «chr» boost_tree boost_tree boost_tree boost_tree boost_tree nearest_neighbor nearest_neighbor nearest_neighbor nearest_neighbor nearest_neighbor nearest_neighbor nearest_neighbor rearest_neighbor		
mple_knn mple_knn mple_knn a_x.yaboost 41 of 65 rows tibble: 65 × 9 #flow_id chr> ca_xyaboost ca_xyaboost ca_xyaboost lter_knn lter_knn lter_knn lter_knn lter_knn ca_rf ca_rf	Preprocessor1_Model03 Preprocessor1_Model03 Preprocessor1_Model03 Preprocessor03_Model1 **Config** <hr/> <	roc_auc sens spec accuracy .metric .cchr> mn_log_loss roc_auc sens spec accuracy mn_log_loss roc_auc sens	0.8731248 0.7634278 0.8457490 0.8032726	0.008428719 0.021291548 0.016205609 mean <a href="</td><td>std_err
cdb>
std_err
cdb>
0.058997799
0.013696577
0.02292694
0.014114953
0.171455762
0.009769647
0.025274277
0.017651506
0.016285640
0.016033001</td><td>100 100 100 100 100 100 100 100 100 100</td><td>nearest_neighbor nearest_neighbor nearest_neighbor nearest_neighbor boost_tree preprocessor <<hr/> <hr/> chr> recipe recipe</td><td>model schr> boost_tree boost_tree boost_tree boost_tree nearest_neighbor nearest_neighbor nearest_neighbor nearest_neighbor rearest_neighbor rearest_neighbor rearest_neighbor rearest_neighbor rearest_neighbor</td><td></td></tr><tr><td>mple_knn mple_knn mple_knn a_x_pshoost 11 of 65 rows tibble: 65 x 9 fflow_id chr> ca_xgboost ca_xgboost ca_xgboost tea_kgboost tea_kgboost tex_knn lter_knn lter_knn lter_knn lter_knn ter_knn ca_rf ca_rf</td><td>Preprocessor1_Model03 Preprocessor1_Model03 Preprocessor1_Model03 Preprocessor03_Model1 </td><td>roc_auc sens spec accuracy .metric .<chr> .<chr> mn_log_loss roc_auc sens spec accuracy mn_log_loss roc_auc sens spec accuracy cacuracy cacuracy sens spec</td><td>0.8731248
0.7634278
0.8457490
0.8032726</td><td></td><td>10 recipe 10 rec</td><td>100 100 100 100 100 100 100 100 100 100</td><td>nearest_neighbor nearest_neighbor nearest_neighbor boost_tree preprocessor <chr> chr> recipe recipe</td><td>model «chr» boost_tree boos</td><td></td></tr><tr><td>mple_knn mple_knn mple_knn a_xypboost 41 of 65 rows tibble: 65 x 9 tibble: 6</td><td>Preprocessor1_Model03 Preprocessor1_Model03 Preprocessor1_Model03 Preprocessor03_Model1 **Config** <hr/> <</td><td>roc_auc sens spec accuracy .metric .cchr> mn_log_loss roc_auc sens spec accuracy mn_log_loss roc_auc sens</td><td>0.8731248
0.7634278
0.8457490
0.8032726</td><td>0.008428719 0.021291548 0.016205609 mean 						

Figure 11-14

Candidate Models/Model Evaluation/Tuning

Candidate Models:

For the first candidate model, we noticed in the workflow set (figure above) that the boosted tree model with the base recipe seemed to perform the best. So, we decided to tune the hyperparameters of the boosted tree model and see how it performs individually. However, we found that the cross validation score for this model was lower than the lightgbm engine (the fourth candidate model).

For the second candidate model, we noticed that the logistic regression, random forest, and boosted tree models with the base recipe (mentioned above) were the top performing workflows in the figure above. So, we decided to try and create an ensemble model for this as well. Ultimately, this ensemble model had the highest performing accuracy score from R amongst all the ensemble models.

Similar to the previous model, the third model is also an ensemble model. After trying out different combinations of ensemble models, we found that the ensemble model combining random forest

and the boosted tree model performed had the highest accuracy rate from cross validation. So, we decided to use this model as another potential candidate. We found that this ensemble model had a lower accuracy score from R than the second candidate model despite having the same Kaggle score. So, we decided to use the previous candidate model.

The fourth model was designed following the completion of boosted tree engine testing. When no tuning was applied, lightgbm returned higher accuracy than xgboost during cross-validation comparisons. The lightgbm engine also scored well on the public leaderboard and was a candidate for as a simple model. With this model, we found that it was less prone to overfitting. Additionally, it had the highest accuracy score from cross validation amongst all the non-ensemble models. So, we decided to use this as our second final model.

For the fifth model, we also decided to try an mlp engine. Although we did not include it in the workflow set, we were curious about its overall performance. So, we applied the same base recipe that we applied previously. Because the other mlp engines required other packages (such as tensorflow), we decided to use the "nnet" engine, which allows for a single hidden layer. This model did not perform as well as the boosted tree models mentioned above, which is shown in both its lower accuracy and Kaggle score.

For the sixth model, we tried logistic regression using glmnet engine. We chose to test this model as it was a simple and quick model to run. However, the results were average compared to our other models. Thus, we decided not to choose this as our final model. Like the fifth model, this model also had a lower accuracy and Kaggle score. So, we decided to use the lightgbm boosted tree model instead.

Recipes Used in the Candidate Models Below

Recipe Number	Recipe
Recipe 1	train_removed_downsample <- recipe(winner ~., data = train_removed) %>% step_downsample(winner) %>% prep() %>% juice() base_recipe <- recipe(winner ~., data = train_removed_downsample) %>% step_mutate(x2013_code = factor(x2013_code)) %>% step_dummy(x2013_code) %>% step_impute_median(all_numeric()) %>%
	step_mutate(prop_white = x0064e / x0001e, # black + american Indian + asian + native Hawaiian + some other race + hispanic prop_minor = (x0064e + x0065e + x0066e + x0067e + x0068e + x0069e + x0071e)/ x0001e, prop_women = x0003e / x0001e, prop_men = x0002e / x0001e, prop_minor = x0019e / x0001e, prop_college = (c01_005e + c01_013e + c01_018e + c01_021e + c01_024e + c01_027e) / x0001e, prop_old = x0024e / x0001e) %>%

```
step rm(x0064e) \% > \%
                     step rm(x0065e) \% > \%
                     step rm(x0066e) %>%
                     step rm(x0067e) %>%
                     step rm(x0068e) %>%
                     step rm(x0069e) %>%
                     step rm(x0071e) %>%
                     step rm(x0003e) %>%
                     step rm(x0002e) \% > \%
                     step rm(x0019e) %>%
                     step rm(x0020e) \% > \%
                     step rm(x0021e) %>%
                     step rm(x0022e) %>%
                     step rm(x0023e) \% > \%
                     step rm(x0024e) \% > \%
                     step rm(x0005e) %>%
                     step rm(x0006e) \% > \%
                     step rm(x0007e)
Recipe 2
                    oprop rec <-
                     recipe(winner \sim ., data = train) %>%
                     update role(id, new role = "id") %>%
                     step naomit(all predictors()) %>%
                     step mutate(
                      prop male = x0002e / x0001e,
                      prop female = x0003e / x0001e,
                      prop adult male = x0088e / x0087e,
                      prop adult female = \times 0089e / \times 0087e,
                      prop minor = x0019e / x0001e,
                      prop senior = x0024e / x0001e,
                      prop white = x0037e / x0001e,
                      prop mix combo = (x0064e + x0065e + x0066e + x0067e + x0068e + x0069e)/x0001e,
                      prop hispanic = x0071e / x0001e,
                      prop hs = (c01\ 003e + c01\ 004e + c01\ 017e + c01\ 020e + c01\ 023e + c01\ 026e) / x0001e
                      prop b = (c01\ 005e + c01\ 018e + c01\ 021e + c01\ 024e + c01\ 027e) / x0001e
                     ) %>%
                     step rm(
                      x0002e, x0003e, x0088e, x0089e,
                      x0024e,
                      x0064e:x0071e,
                      c01 003e:c01 005e, c01 017e, c01 018e, c01 020e, c01 021e,
                      c01 023e, c01 024e, c01 026e, c01 027e
                     ) %>%
                     step mutate(x2013 code = factor(x2013 code)) %>%
                     step rm(
                      x0019e:x0031e, # cumulative
                      x0033e, # dupe x0001e
                      x0036e, # dupe x0034e
                      x0058e, # dupe x0035e
                      c01 006e:c01 016e, # cumulative & dupe x0010e
```

```
c01 019e, # dupe x0011e
c01 022e, # dupe x0012e:x0014e
 c01 025e # dupe x0015e:x0017e
) %>%
step_rm(
x0040e:x0043e, # native subcategory
 x0045e:x0051e, # asian subcategory
 x0053e:x0056e, # hawaii subcategory
 x0072e:x0075e, # hispanic subcategory
 x0076e:x0085e, # non-hispanic category
) %>%
step impute median(all numeric predictors()) %>%
step normalize(all numeric predictors()) %>%
step zv(all predictors()) %>% # zero variance predictors
step nzv(all predictors()) %>% # near-zero variance predictors
step_dummy(all_nominal_predictors())
```

Candidat e Model Number	Model Identifier	Type of Model	Engine	Recipes or Listing of Other Variables in Model:	Hyperparameters
1	xgboost2024	Boosted Tree	xgboost	Recipe 1	learn_rate = 0.00425, trees = 1857, tree_depth = 12, mtry =73, min_n = 2, loss_reduction = 0.00386, sample_size = 0.987, stop_iter = 4
2	stacks	Logistic Regression Random Forest Boosted Tree	glmnet randomForest xgboost	Recipe 1	Logistic Regression: penalty = 0.000287, mixture = 0.719 Random Forest: min_n = 4, trees = 104 Boosted Tree: learn_rate = 0.0664, trees = 1309, tree_depth = 4, mtry = 253, min_n = 7, loss_reduction = 0.000105, sample_size = 0.994, stop_iter = 19

					Blend Predictions: penalty = 10^-2
3	stacks_rf_xgb oost	Random Forest Boosted Tree	randomForest xgboost	Recipe 1	Random Forest: min_n = 12, trees = 242, mtry = 15
					Boosted Tree: learn_rate = 0.0102, trees = 1500, tree_depth = 14, mtry = 43, min_n = 4, loss_reduction = 0.00000479, sample_size = 0.527, stop_iter = 20
					Blend Predictions: penalty = 10^-3
4	29_lbg	Boosted Tree	lightgbm	Recipe 2	N/A
5	mlp_nnet	MLP Model	nnet	Recipe 1	hidden_units = 7, penalty = 0, epochs = 600
6	log_reg_res	Logistic Regression	glmnet	Recipe 1	Penalty = 0.0000000127, Mixture = 0.0839

Evaluations and Tuning

Our candidate models mainly included logistic, random forest and boosted tree models. Models that include xgboost and lightgbm perform the best according to Kaggle leaderboard. We also tried MLP, however, that did not work as well with the workflow, most likely due to overfitting. In our top candidate models, we use latin hypercube sampling (LHS) to give us the most efficiency when it comes to finding the best parameters without overfitting. Some interesting things we noticed about LHS is that when comparing using LHS versus not for logistic regression is that there were many differences in accuracy or Kaggle score. LHS introduces randomness to the hyperparameter search which is more efficient when it comes to finding the best parameter combinations. One major challenge we came across with LHS is that it never produces the same exact parameters which causes our models to have different outcomes. Thus, we decided to manually input the best parameters from LHS instead of R selecting it.

To compare and evaluate the candidate models, we mainly used v-fold cross validation with 10 folds and stratified on the winner response variable. We assessed model performance based on the cross

validation accuracy score, the standard error (if applicable), and the Kaggle score. From these results, we determined that the two best models were the stacks and 29_lbg models due to their high accuracy scores, relatively low SEs, and high scores when submitted to Kaggle.

We also implemented the stacks autoplot function to gain insights into which model generally produced the most accurate predictions. As shown in the plots, the ensemble models that combine logistic regression, random forest, and xgboost achieved the highest accuracy scores. The autoplot of the stack model revealed that logistic regression had the highest stacking coefficient, indicating its importance in the ensemble.

Results

Model Identifier	Accuracy Score from Cross Validation	SE (if applicable)	Kaggle Public Leaderboard Score (if submitted)
stacks	0.939	N/A	0.96652
29_lbg	0.939	0.0040529	0.94560
stacks_rf_xgboost	0.932	N/A	0.94560
xgboost2024	0.930	0.00594	0.94560
log_reg_res	0.935	0.00356	0.93305
mlp_nnet	0.834	0.000599	N/A

Model Performance Evaluation Plots

Autoplot Comparison of stacks_rf_xgboost, log_reg_res, mlp_nnet, xgboost2024, & 29_lbg

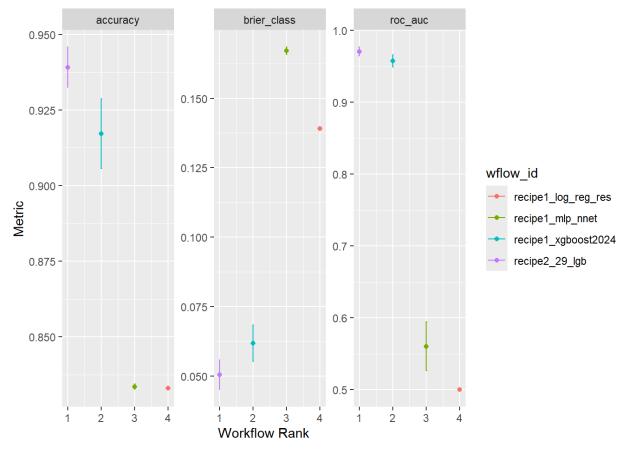
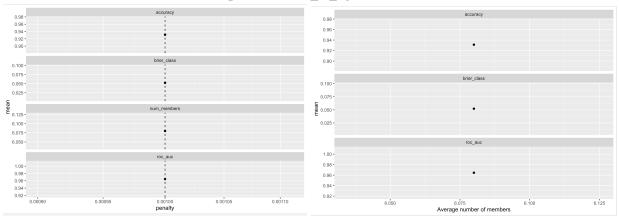


Figure 15

Autoplot of stacks_rf_xgboost Model



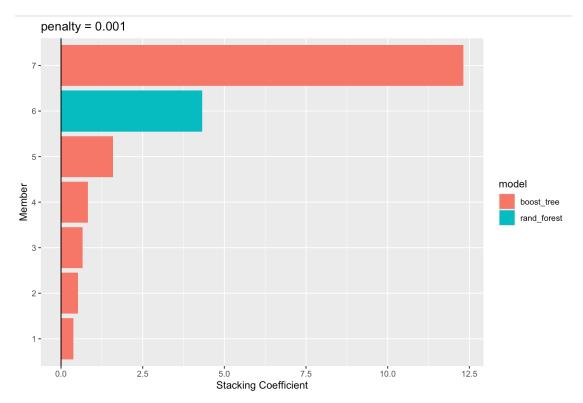
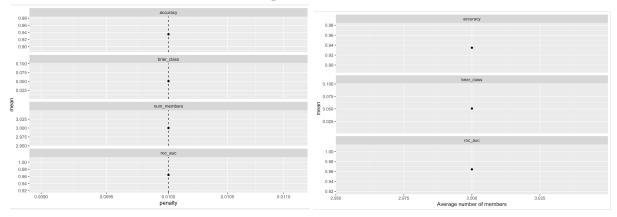


Figure 16 - 18

Autoplot of Stacks Model



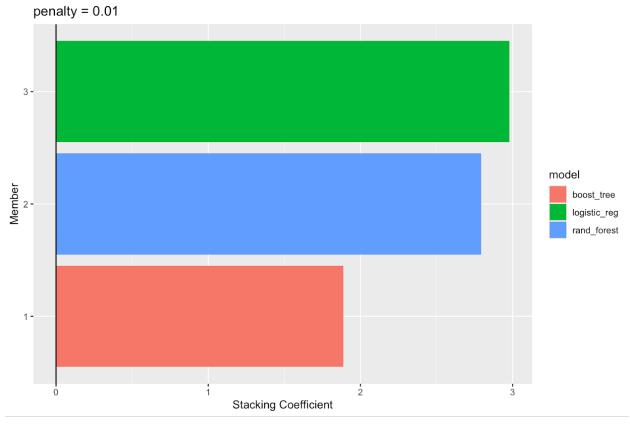


Figure 19-21

Discussion of Final Model

The two models selected for submission were 29_lbg and stacks, with 29_lbg responsible for our final private leaderboard score. These two models were selected because they produced the highest accuracy score based on cross-validation metrics. Additionally, we thought that the 29_lbg model itself would be able to compensate for the hypertuning we used in the stacks model. So, any overfitting the stacks model may have would be able to be balanced out by the 29_lbg model. After submitting both models as our final candidates, we found that our hypothesis about the 29_lbg model being able to avoid overfitting was actually correct. 29_lbg ended up being our better model according to the private leaderboard score.

The greatest strength of 29_lbg is the simplicity of the model. 29_lbg is a gradient boosting model (boosted trees) utilizing the lightgbm engine with all model parameters left at default. Ease of model application was appealing for this project since the data required complex preprocessing and recipes. Therefore, the simplicity of this setup leaves minimal points of failure for reproducibility since pseudorandomization only occurs during k-fold cross-validation. Another strength of this model is that it is able to capture nonlinear relationships, which allows it to better predict accurate results.

29_lbg has some weaknesses. As with many gradient-boosting models, lightgbm requires a sufficiently large dataset to make accurate predictions. The training set provided was on a national scale,

which provided an adequate amount of observations, but the model would likely perform worse if the scope of observations was scaled down to the state level. Another weakness was that lightgbm required a specific complex recipe to make good predictions. During engine testing, the lightgbm engine's performance differed depending on the recipe applied, so applications or alterations to the model for different data may be time-consuming. Lastly, this lightgbm model, like other boosted tree models, is more difficult to interpret than other models such as logistic regression.

There are possible improvements that may benefit the 29_lbg model. First, testing and tuning an ensemble model using lightgbm may improve performance. Engine testing was very time-consuming for this project, so the performance of the lightgbm engine was discovered late into the project. However, even with a basic implementation of the lightgbm engine, 29_lbg and an ensemble model utilizing lightgbm were our 2nd and 3rd best models on the private leaderboard respectively. Another improvement we can make with our model is experimenting more with removing correlated predictors. While we did not have enough time to extensively play around with different recipes, we believe it would be worthwhile to consider converting a lot of the population predictors into proportions because a lot of the predictors overlapped; for instance, a lot of the people in the "Race alone or in combination with one or mother other races: Total population: White" (x0064e) variable would also be in the "Total population:One race:White" (x0037e) variable. Also, we would have liked to experiment more with removing the income_per_cap and gdp variables since a lot of them would be correlated with one another; for instance, gdp_2018 should be some type of multiple of gdp_2017.

Some additional data that could help us improve the model is data on immigration status. Although there is data on the proportion of citizens that are each race, we were curious about whether being a naturalized citizen or a U.S. born citizen impacts voting results. From the Public Policy Institute of California's article "Immigrants and Political Engagement," the article writes that naturalized citizens typically vote for Democrats. However, this is already assuming that naturalized citizens will go and vote. But, what if naturalized citizens tend to abstain from voting? This is a factor that we believe the current data failed to encapsulate. We were also curious about whether married status impacts voter turnout. In the article "Changing Partisan Coalitions in a Politically Divided Nation," Pew Research Center notes that married men and women are more likely to vote for the Republican Party. With a correlation as strong as this, we felt that our model could potentially be more accurate if we were able to have metrics that quantified this factor.

Appendix

Final annotated script

Final Model Submission 29 lgb.R by Oliver Siu

Screen recording of script running

https://drive.google.com/file/d/1YSZj FL N-6FIEMeKh28CMYizZYfj35B/view?usp=sharing

```
29_lgb.R
# load libraries and set preferences
library(tidyverse)
library(tidymodels)
tidymodels prefer()
library(bonsai)
library(lightgbm)
# load data
train <- read csv("train class.csv")</pre>
test <- read_csv("test_class.csv")</pre>
# preprocessing
train <-
 train %>%
 select(!name) %>%
 mutate(winner = as.factor(winner))
# set seed
set.seed(2024)
# recipe setup
oprop_rec <-
 recipe(winner ~ ., data = train) %>%
 update role(id, new role = "id") %>%
 step naomit(all predictors()) %>%
 step mutate(
   prop male = x0002e / x0001e,
   prop female = x0003e / x0001e,
   prop adult male = x0088e / x0087e,
```

prop adult female = x0089e / x0087e,

```
prop minor = x0019e / x0001e,
   prop senior = x0024e / x0001e,
   prop white = x0037e / x0001e,
    prop mix combo = (x0064e + x0065e + x0066e + x0067e + x0068e +
x0069e)/ x0001e,
    prop hispanic = x0071e / x0001e,
    prop hs = (c01 003e + c01 004e + c01 017e + c01 020e + c01 023e +
c01 026e) / x0001e,
    prop b = (c01 005e + c01 018e + c01 021e + c01 024e + c01 027e) /
x0001e
 ) 응>응
 step rm(
    x0002e, x0003e, x0088e, x0089e,
   x0024e,
   x0064e:x0071e,
    c01 003e:c01 005e, c01 017e, c01 018e, c01 020e, c01 021e,
    c01 023e, c01 024e, c01 026e, c01 027e
  step mutate(x2013 code = factor(x2013 code)) %>%
  step rm(
   x0019e:x0031e, # cumulative
   x0033e, # dupe x0001e
   x0036e, # dupe x0034e
   x0058e, # dupe x0035e
    c01 006e:c01 016e, # cumulative & dupe x0010e
    c01 019e, # dupe x0011e
    c01 022e, # dupe x0012e:x0014e
    c01 025e # dupe x0015e:x0017e
  ) %>%
  step rm(
    x0040e:x0043e, # native subcategory
   x0045e:x0051e, # asian subcategory
   x0053e:x0056e, # hawaii subcategory
   x0072e:x0075e, # hispanic subcategory
   x0076e:x0085e, # non-hispanic category
  ) %>%
 step impute median(all numeric predictors()) %>%
  step normalize(all numeric predictors()) %>%
  step_zv(all_predictors()) %>% # zero variance predictors
  step nzv(all predictors()) %>% # near-zero variance predictors
  step dummy(all nominal predictors())
```

```
# model setup
lgb spec <-
 boost tree() %>%
 set engine("lightgbm") %>%
  set mode("classification")
# workflow setup
lgb wflow <-
 workflow() %>%
 add recipe(oprop rec) %>%
 add_model(lgb_spec)
# folds for k-fold (v-fold) cross-validation and options setup
folds <- train %>% vfold_cv(v = 10, strata = winner)
keep pred <- control resamples(save pred = TRUE, save workflow = TRUE)
# k-fold (v-fold) cross-validation
lgb res <-
 lgb wflow %>%
 fit resamples(resamples = folds, control = keep pred)
# fit model to training data
lgb fit <- lgb wflow %>% fit(data = train)
# making predictions on test data
predictions <- lgb fit %>% predict(new data = test)
# format final output
solution <-
  test %>%
  select(id) %>%
 bind cols(predictions) %>%
 rename(winner = .pred_class)
# write out final output as csv
write csv(solution, "solution.csv")
```

Team member contributions

A. Katherine Huynh

- a. Also created stacks model which we ended up submitting as one of our candidate models
- b. Created xgboost2024 and stacks rf xgboost model
 - i. Both of which tied for third place on the public leaderboard for 3 points extra credit
- c. Created Figure 1-5
- d. Created base recipe of figuring out which columns were duplicates
- e. Created workflow set of 3 recipes and 5 different models and ran a workflow map to see the models with the best predictive capability
- f. Typed the preprocessing section and the additional data section in the discussion of final model

B. Victoria Aye

- a. Tested and improved stack model, which resulted in being the highest public score out of all our submission and 2nd in overall
- b. Tested logistic regression model and wrote about it
- c. Cowrote evaluation, model tuning, and discussion of Final Model

C. Oliver Siu

- a. Designed 2 of the team's top 5 private scoring models
- b. Created Figures 8-10
- c. Created 29 lbg model
- d. Assisted with recipe research
- e. Responsible for engine research for rand forest, logist reg, & boost tree
- f. Generated comparison plots for non-ensemble models
- g. Cowrote Discussion of Final Model

D. Vivian Yee

- a. Created Figures 6 and 7
- b. Tested models to improve public score on Kaggle (logistic regression, decision tree, boosted tree)
- c. Wrote introduction
- d. Cowrote evaluation and model tuning