Project Phase2

Engel, Alec 3/6/2021

Load in data set

```
# summary(student)
# glimpse(student)
# skim(student)
```

Data cleaning to adjust for bad values, convert characters into factors and remove unnecessary variables.

Eliminate outliers

```
student = student %>%
  filter(Lot Area < 40000) %>%
  filter(BsmtFin SF 1 < 1600) %>%
  filter(BsmtFin_SF_2 < 400) %>%
  filter(Bsmt Unf SF < 2250) %>%
  filter(Total Bsmt SF < 2750) %>%
  filter(Full Bath > 0) %>%
  filter(Half Bath < 1.1) %>%
  filter(First_Flr_SF < 2750) %>%
  filter(Second_Flr_SF < 1400) %>%
  filter(Gr_Liv_Area < 3750) %>%
  filter(Fireplaces < 4) %>%
  filter(Garage Cars < 4) %>%
  filter(Garage Area < 1250) %>%
  filter(Wood_Deck_SF < 550) %>%
  filter(Open_Porch_SF < 350) %>%
  filter(Enclosed Porch < 300) %>%
```

```
filter(Three season porch < 240) %>%
  filter(Screen Porch < 400)</pre>
Train and test split
set.seed(123)
student split = initial split(student, prob = 0.75, strata = Above Median)
train = training(student split)
test = testing(student_split)
5 K Folds
set.seed(123)
folds = vfold_cv(train, v = 5)
Basic recipe
student_recipe = recipe(Above_Median ~., train) %>%
  step other(Neighborhood, threshold = .02) %>%
  step_other(MS_SubClass,threshold = .02) %>%
  step other(Overall Qual, threshold = .02) %>%
  step_other(Overall_Cond, threshold = .02) %>%
  step_other(Exterior_1st, threshold = .02) %>%
  step other(Functional, threshold = .02) %>%
  step dummy(all nominal(), -all outcomes()) %>%
  step_nzv(all_predictors())
ctrl grid = control stack grid()
ctrl_res = control_stack_resamples()
```

Student Log Regression

```
# student log model =
    logistic_reg(mode = "classification") %>%
#
    set_engine("glm")
#
# student log recipe = student recipe %>%
    step dummy(all nominal(), -all outcomes())
#
#
# Logreg_wf = workflow() %>%
#
    add recipe(student log recipe) %>%
    add model(student log model)
#
#
# set.seed(123)
# log_res =
#
  tune_grid(
#
      Logreg wf,
#
      resamples = folds,
#
      grid = 200,
#
      control = ctrl grid
      )
```

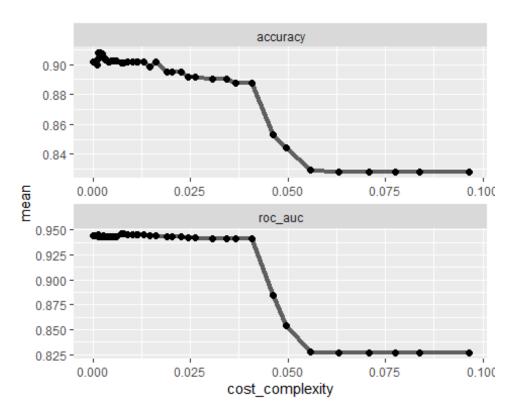
```
# saveRDS(log_res, "log_res.rds")
log_res = readRDS("log_res.rds")
```

Student Classification Tree Model

```
# tree_model = decision_tree(cost_complexity = tune()) %>%
    set_engine("rpart", model = TRUE) %>%
    set_mode("classification")
#
#
# tree_recipe = student_recipe
#
# tree_workflow = workflow() %>%
#
    add model(tree model) %>%
#
    add_recipe(tree_recipe)
#
# set.seed(123)
# tree_res =
# tree_workflow %>%
# tune_grid(
#
      resamples = folds,
#
      grid = 200,
      control = ctrl_grid
#
#
      )
# saveRDS(tree_res, "tree_res.rds")
tree_res = readRDS("tree_res.rds")
```

Classification Tree Model Accuracy Chart

```
tree_res %>%
  collect_metrics() %>%
  ggplot(aes(cost_complexity, mean)) +
  geom_line(size = 1.5, alpha = 0.6) +
  geom_point(size = 2) +
  facet_wrap(~ .metric, scales = "free", nrow = 2)
```

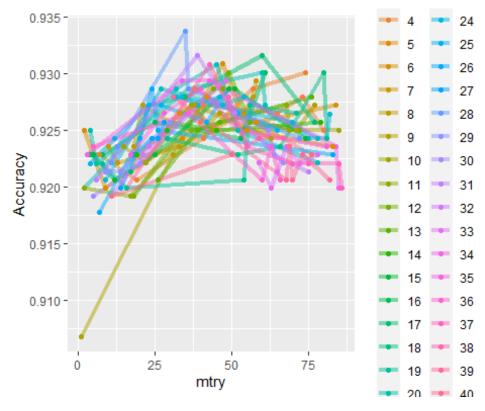


Student Random Forest Model

```
# rf_recipe = student_recipe
# rf_model = rand_forest(mtry = tune(), min_n = tune(), trees = 200) %>%
    set_engine("ranger", importance = "permutation") %>%
    set_mode("classification")
#
#
# rf_wflow =
    workflow() %>%
#
    add_model(rf_model) %>%
#
#
    add_recipe(rf_recipe)
# set.seed(123)
# rf_res = tune_grid(
    rf_wflow,
#
    resamples = folds,
#
    grid = 200,
#
    control = ctrl_grid
# saveRDS(rf_res, "rf_res.rds")
rf_res = readRDS("rf_res.rds")
```

Random Forest Model Accuracy Chart

```
rf_res %>%
  collect_metrics() %>%
  filter(.metric == "accuracy") %>%
  mutate(min_n = factor(min_n)) %>%
  ggplot(aes(mtry, mean, color = min_n)) +
  geom_line(alpha = 0.5, size = 1.5) +
  geom_point() +
  labs(y = "Accuracy")
```



Student Neural Network Model

```
# nn_recipe = student_recipe %>%
    step_normalize(all_predictors(), -all_nominal()) %>%
#
#
    step_dummy(all_nominal(), -all_outcomes())
#
# nn_model =
    mlp(hidden_units = tune(), penalty = tune(),
#
#
        epochs = tune()) %>%
#
    set_mode("classification") %>%
    set_engine("nnet", verbose = 0)
#
#
# nn_workflow <-
    workflow() %>%
#
#
    add_recipe(nn_recipe) %>%
#
    add model(nn model)
#
# set.seed(123)
```

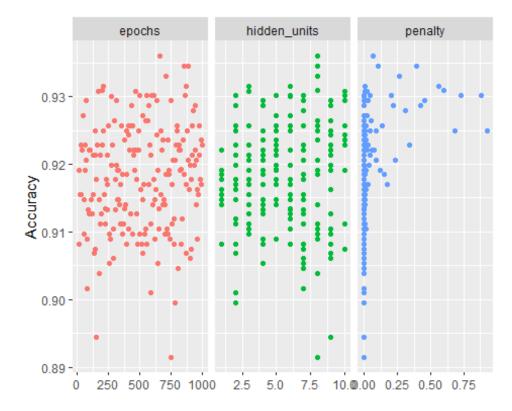
```
# neural_res <-
# tune_grid(nn_workflow,
# resamples = folds,
# grid = 200,
# control = ctrl_grid)

# saveRDS(neural_res, "neural_res.rds")

neural_res = readRDS("neural_res.rds")</pre>
```

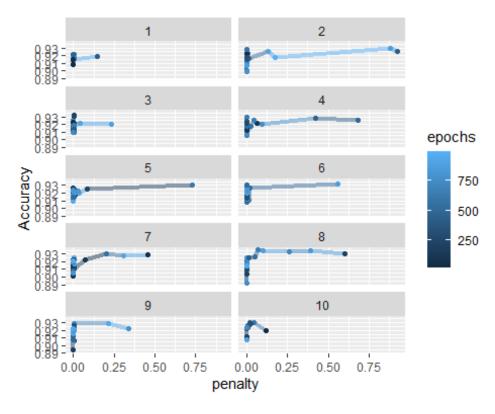
Neural Network Model Epochs/Hidden_units/Penalties Charts #1

```
neural_res %>%
   collect_metrics() %>%
   filter(.metric == "accuracy") %>%
   dplyr::select(mean, hidden_units, penalty, epochs) %>%
   pivot_longer(hidden_units:epochs,
      values_to = "value",
      names_to = "parameter"
) %>%
   ggplot(aes(value, mean, color = parameter)) +
   geom_point(show.legend = FALSE) +
   facet_wrap(~parameter, scales = "free_x") +
   labs(x = NULL, y = "Accuracy")
```



Neural Network Model Epochs/Hidden_units/Penalties Charts #2

```
neural_res %>%
  collect_metrics() %>%
  filter(.metric == "accuracy") %>%
  mutate(hidden_units = factor(hidden_units)) %>%
  ggplot(aes(penalty, mean, color = epochs)) +
  geom_line(alpha = 0.5, size = 1.5) +
  geom_point() +
  facet_wrap(~hidden_units, ncol =2 ) +
  labs(y = "Accuracy")
```



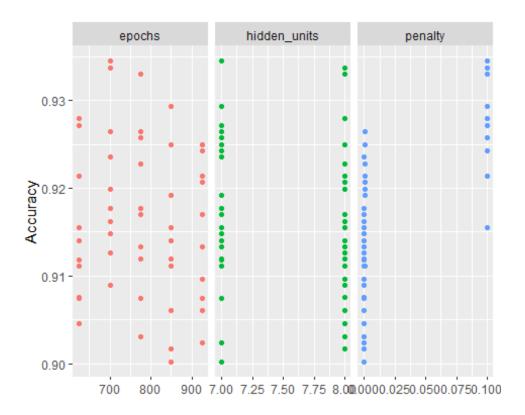
Student Neural Network Model with Parameter Tuning

```
# neural_grid = grid_regular(
    hidden units(range = c(7,8)),
#
    penalty(range = c(-10, -1)),
#
    epochs(range = c(625,925)),
    levels = 5
#
# )
#
# student nn recipe = student recipe %>%
    step_normalize(all_predictors(), -all_nominal()) %>%
#
#
    step_dummy(all_nominal(), -all_outcomes())
#
# student_nn_model =
   mlp(hidden units = tune(), penalty = tune(),
#
#
        epochs = tune()) %>%
# set_mode("classification") %>%
```

```
# set_engine("nnet", verbose = 0)
#
# student_nn_workflow <-
# workflow() %>%
# add_recipe(student_nn_recipe) %>%
# add_model(student_nn_model)
#
# set.seed(123)
# neural_tune_res <-
# tune_grid(student_nn_workflow, resamples = folds, grid = neural_grid, control = ctrl_grid)
# saveRDS(neural_tune_res, "neural_tune_res.rds")
neural_tune_res = readRDS("neural_tune_res.rds")</pre>
```

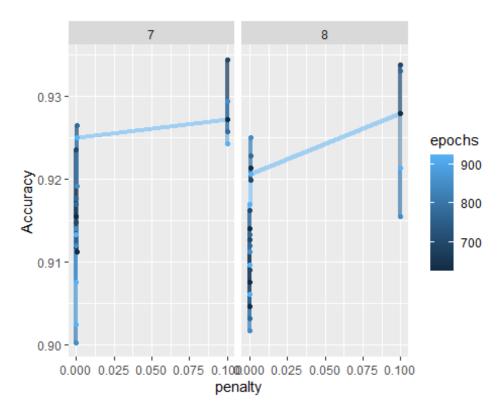
Neural Network Parameter Tuned Epochs/Hidden_units/Penalties Charts

```
neural_tune_res %>%
  collect_metrics() %>%
  filter(.metric == "accuracy") %>%
  dplyr::select(mean, hidden_units, penalty, epochs) %>%
  pivot_longer(hidden_units:epochs,
    values_to = "value",
    names_to = "parameter"
) %>%
  ggplot(aes(value, mean, color = parameter)) +
  geom_point(show.legend = FALSE) +
  facet_wrap(~parameter, scales = "free_x") +
  labs(x = NULL, y = "Accuracy")
```



Neural Network Parameter Tuned Model Epochs/Hidden_units/Penalties Charts #2

```
neural_tune_res %>%
  collect_metrics() %>%
  filter(.metric == "accuracy") %>%
  mutate(hidden_units = factor(hidden_units)) %>%
  ggplot(aes(penalty, mean, color = epochs)) +
  geom_line(alpha = 0.5, size = 1.5) +
  geom_point() +
  facet_wrap(~hidden_units, ncol =2 ) +
  labs(y = "Accuracy")
```



*Student XGBOOST Model

```
# xgboost_recipe2 <- student_recipe %>%
    #step_novel(all_nominal(), -all_outcomes()) %>%
    step_dummy(all_nominal(), -all_outcomes(), one_hot = TRUE) %>%
#
#
    step_zv(all_predictors())
# xgboost_spec2 <-</pre>
   boost_tree(trees = tune(), min_n = tune(), tree_depth = tune(),
learn_rate = tune(),
#
      loss_reduction = tune(), sample_size = tune()) %>%
#
    set_mode("classification") %>%
#
    set_engine("xgboost")
# xqboost workflow2 <-
   workflow() %>%
#
    add_recipe(xgboost_recipe2) %>%
#
    add_model(xgboost_spec2)
# set.seed(123)
# xgboost_tune_res <-</pre>
# tune_grid(xgboost_workflow2, resamples = folds, grid = 200, control =
ctrl_grid)
# saveRDS(xgboost_tune_res, "xgboost_tune_res.rds")
xgboost_tune_res = readRDS("xgboost_tune_res.rds")
```

*Student XGBOOST Model with Parameter Tuning

```
# tgrid = expand.grid(
  trees = 100,
# min n = 1,
# tree_depth = c(1,2,3,4),
#
  learn_rate = c(0.01, 0.1, 0.2, 0.3, 0.4),
# loss reduction = 0,
  sample_size = c(0.5, 0.8, 1))
#
#
# xqboost recipe <-</pre>
# student recipe %>%
# step_dummy(all_nominal(), -all_outcomes(), one_hot = TRUE) %>%
#
  step_zv(all_predictors())
# xgboost_spec <-</pre>
  boost tree(trees = tune(), min n = tune(), tree depth = tune(),
learn_rate = tune(),
    loss_reduction = tune(), sample_size = tune()) %>%
# set_mode("classification") %>%
#
  set_engine("xgboost")
#
# xqboost workflow <-
# workflow() %>%
# add_recipe(xgboost_recipe) %>%
  add model(xqboost spec)
#
# set.seed(123)
# xqb res <-
# tune_grid(xgboost_workflow,
#
             resamples = folds,
#
             grid = tgrid,
#
             control = ctrl_grid)
# saveRDS(xgb_res, "xgb_res.rds")
xgb res = readRDS("xgb res.rds")
```

Stacking Building Stack

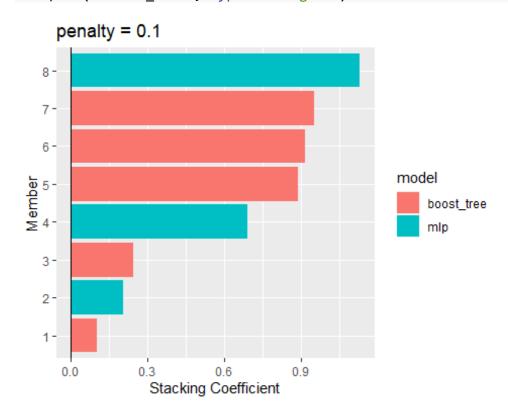
```
student_stacks = stacks() %>%
  add_candidates(tree_res) %>%
  add_candidates(rf_res) %>%
  add_candidates(xgb_res) %>%
  add_candidates(neural_tune_res) %>%
  add_candidates(log_res)
# add_candidates(neural_res)
# add_candidates(xgboost_tune_res)
```

fitting a Lasso model to the stack.

```
student_blend =
  student_stacks %>%
  blend_predictions(metric = metric_set(accuracy))
##
## Attaching package: 'rlang'
## The following objects are masked from 'package:purrr':
##
       %@%, as_function, flatten, flatten_chr, flatten_dbl, flatten_int,
##
       flatten_lgl, flatten_raw, invoke, list_along, modify, prepend,
##
##
       splice
##
## Attaching package: 'vctrs'
## The following object is masked from 'package:dplyr':
##
##
       data_frame
## The following object is masked from 'package:tibble':
##
##
       data_frame
```

Strongest Resulting Models

```
autoplot(student_blend, type = "weights")
```



Fitting the stack to training data

#student_blend <- # student_blend %>% # fit_members()

Training data predictions

```
trainpredstack = predict(student_blend, train)
head(trainpredstack)

## # A tibble: 6 x 1

## .pred_class

## <fct>
## 1 Yes

## 2 No

## 3 Yes

## 4 Yes

## 5 Yes

## 6 Yes
```

Training data confusion matrix

```
confusionMatrix(trainpredstack$.pred_class, train$Above_Median,
                positive = "Yes")
## Confusion Matrix and Statistics
##
             Reference
##
## Prediction Yes No
##
          Yes 676
                4 690
##
          No
##
##
                  Accuracy : 0.9949
                    95% CI: (0.9895, 0.9979)
##
##
       No Information Rate: 0.5047
       P-Value [Acc > NIR] : <2e-16
##
##
##
                     Kappa: 0.9898
##
##
   Mcnemar's Test P-Value : 1
##
##
               Sensitivity: 0.9941
##
               Specificity: 0.9957
##
            Pos Pred Value: 0.9956
            Neg Pred Value: 0.9942
##
##
                Prevalence: 0.4953
            Detection Rate: 0.4924
##
##
      Detection Prevalence: 0.4945
##
         Balanced Accuracy: 0.9949
##
##
          'Positive' Class : Yes
##
```

Test data predictions

```
testpredstack = predict(student_blend, test)
head(testpredstack)

## # A tibble: 6 x 1

## .pred_class

## <fct>
## 1 Yes

## 2 Yes

## 3 Yes

## 4 No

## 5 Yes

## 6 Yes
```

Test data confusion matrix

```
confusionMatrix(testpredstack$.pred class, test$Above Median,
                positive = "Yes")
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction Yes No
         Yes 204 18
##
##
          No
               22 213
##
##
                  Accuracy : 0.9125
##
                    95% CI: (0.8827, 0.9367)
##
       No Information Rate: 0.5055
##
       P-Value [Acc > NIR] : <2e-16
##
##
                     Kappa: 0.8249
##
##
   Mcnemar's Test P-Value: 0.6353
##
##
               Sensitivity: 0.9027
##
               Specificity: 0.9221
##
            Pos Pred Value : 0.9189
##
            Neg Pred Value: 0.9064
                Prevalence: 0.4945
##
##
            Detection Rate: 0.4464
##
      Detection Prevalence: 0.4858
##
         Balanced Accuracy: 0.9124
##
          'Positive' Class : Yes
##
##
```

Comparison of model performance on test set

```
test = test %>% bind_cols(predict(student_blend,.))
```

Stacked model compared to constituent models

```
member_testpreds =
 test %>%
 dplyr::select(Above Median) %>%
 bind cols(predict(student blend, test, members = TRUE))
map dfr(member testpreds, accuracy, truth = Above Median, data =
member testpreds) %>%
 mutate(member = colnames(member testpreds))
## # A tibble: 10 x 4
      .metric .estimator .estimate member
     <chr>
##
              <chr>
                             <dbl> <chr>
## 1 accuracy binary
                                   Above Median
## 2 accuracy binary
                            0.912 .pred_class
## 3 accuracy binary
                            0.906 .pred class xgb res 1 11
## 4 accuracy binary
                             0.906 .pred_class_xgb_res_1_15
## 5 accuracy binary
                            0.908 .pred_class_xgb_res_1_24
## 6 accuracy binary
                             0.899 .pred class xgb res 1 27
                             0.902 .pred_class_xgb_res_1_53
## 7 accuracy binary
## 8 accuracy binary
                             0.893 .pred_class_neural_tune_res_1_13
## 9 accuracy binary
                             0.906 .pred_class_neural_tune_res_1_27
## 10 accuracy binary
                             0.904 .pred class neural tune res 1 20
```

Implementation of stack on competition set

```
# competition = read_csv("ames_competition.csv")
# competition = competition %>%
    mutate if(is.character,as factor) %>%
    mutate(Mo_Sold = as_factor(Mo_Sold)) %>%
  mutate(Mo Sold = fct recode(Mo Sold, "Jan" = "1", "Feb" = "2", "Mar" =
"3", "Apr" = "4", "May" = "5", "Jun" = "6",
                                "Jul" = "7", "Aug" = "8", "Sep" = "9", "Oct"
= "10", "Nov" = "11", "Dec" = "12")) %>%
   mutate(BsmtFin_SF_1 = Total_Bsmt_SF - BsmtFin_SF_2 - Bsmt_Unf_SF)
# competitionpredstack = predict(student_blend, competition)
# head(competitionpredstack)
# kaggle = competition %>% dplyr::select(X1)
#
# kaggle = bind cols(kaggle, competitionpredstack)
# kaggle
# write.csv(kaggle, "kaggle_submit4.csv", row.names=FALSE)
```