Model Training: XGBOOST truncated regressor

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```
# Environment Cleaning: Clearing workspace
rm(list = ls())
# load packages
library(rpart)
library(dplyr)
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
       intersect, setdiff, setequal, union
##
library(caret)
## Loading required package: ggplot2
## Loading required package: lattice
library(data.table)
## Attaching package: 'data.table'
## The following objects are masked from 'package:dplyr':
##
##
       between, first, last
library(mlflow)
library(reticulate)
library(Metrics)
## Attaching package: 'Metrics'
```

```
## The following objects are masked from 'package:caret':
##
##
       precision, recall
library(themis)
## Loading required package: recipes
## Attaching package: 'recipes'
## The following object is masked from 'package:stats':
##
##
       step
library(doMC)
## Loading required package: foreach
## Loading required package: iterators
## Loading required package: parallel
library(here)
## here() starts at /Users/masinde/Projects/causal_fairness_Ph_IbF
```

Inputs

```
# Recipe inputs
truncated_train <- read.csv(here("data", "truncated_train.csv"))
truncated_validation <- read.csv(here("data", "truncated_validation.csv"))
truncated_test <- read.csv(here("data", "truncated_test.csv"))

# Combining train and validation datasets to one
# Because we are going to use CV to train the models later
# naming it df_base_train2 to remain consistent with df naming
df_trunc_train2 <- rbind(truncated_train, truncated_validation)

cat("number of rows in combined train data:", nrow(df_trunc_train2), sep = " ")</pre>
```

Model training

number of rows in combined train data: 396

```
# Set up train control with custom seeds
n folds <- 10
n_models <- 25  # adjust depending on search space size, affects seeds length
# Reproducibility: Defining seeds (a little bit complicated because of parallel processing)
# Generate a reproducible list of seeds
set.seed(1234)
seeds_list <- vector(mode = "list", length = n_folds + 1)</pre>
for (i in 1:n folds) {
  seeds_list[[i]] <- sample.int(1000000, n_models) # one seed per model per fold</pre>
seeds_list[[n_folds + 1]] <- sample.int(1000000, 1) # for final model</pre>
# Set up train control with 10-fold cross-validation
train_control <- trainControl(</pre>
  method = "cv",
  number = n_folds,
  summaryFunction = defaultSummary,
  search = "random", # random selection of the expanded grid
  seeds = seeds_list
)
# Detect and register the number of available cores (use all but one)
num_cores <- parallel::detectCores() - 2</pre>
registerDoMC(cores = num_cores) # Enable parallel processing
# Measure the time for a code block to run
system.time({
# Train the model using grid search with 3-fold CV
trunc_xgb_reg_model <- train(</pre>
  damage_perc ~ track_min_dist +
    wind max +
    rain total +
    roof_strong_wall_strong +
    roof_strong_wall_light +
    roof_strong_wall_salv +
    roof_light_wall_strong +
    roof_light_wall_light +
    roof_light_wall_salv +
    roof_salv_wall_strong +
    roof_salv_wall_light +
    roof_salv_wall_salv +
    blue_ss_frac +
    yellow_ss_frac +
    orange_ss_frac +
    red_ss_frac +
    blue_ls_frac +
    yellow_ls_frac +
    orange_ls_frac +
    red_ls_frac,
```

```
data = df_trunc_train2,
  method = "xgbTree",
 trControl = train control,
 tuneLength = n_models, # this replaces tuneGrid
 metric = "RMSE" # Optimize based on RMSE
Sys.sleep(2) # This is just an example to simulate a delay
})
      user system elapsed
## 43.600 0.850 7.614
# Print best parameters
print(trunc_xgb_reg_model$bestTune)
                                    gamma colsample_bytree min_child_weight
##
    nrounds max_depth
                              eta
## 3
                    7 0.07421103 3.36614
                                                0.6218847
## subsample
## 3 0.9780272
# now train again using best parameters
# ----- best parameters -----
best_params <- trunc_xgb_reg_model$bestTune</pre>
trunc_damage_fit_reg <- train(damage_perc ~ track_min_dist +</pre>
                              wind max +
                              rain_total +
                              roof strong wall strong +
                              roof_strong_wall_light +
                              roof_strong_wall_salv +
                              roof_light_wall_strong +
                              roof_light_wall_light +
                              roof_light_wall_salv +
                              roof_salv_wall_strong +
                              roof_salv_wall_light +
                              roof_salv_wall_salv +
                              blue_ss_frac +
                              yellow_ss_frac +
                              orange_ss_frac +
                              red_ss_frac +
                              blue ls frac +
                              yellow_ls_frac +
                              orange_ls_frac +
                              red_ls_frac,
                              method = "xgbTree",
                              trControl = trainControl(method = "none"),
                              tuneGrid = best_params, # Use the best parameters here
                              metric = "RMSE",
                              data = df_trunc_train2
                         )
```

```
# obtain predicted values
train_predictions <- predict(trunc_damage_fit_reg, newdata = df_trunc_train2)</pre>
# Define bin edges
# Define bin edges
bins \leftarrow c(0.00009, 1, 10, 50, 100)
# Assign data to bins
bin_labels <- cut(df_trunc_train2$damage_perc, breaks = bins, include.lowest = TRUE, right = TRUE)
# Create a data frame with actual, predicted, and bin labels
data <- data.frame(</pre>
 actual = df_trunc_train2$damage_perc,
 predicted = train_predictions,
 bin = bin_labels
# Calculate RMSE per bin
unique_bins <- levels(data$bin) # Get unique bin labels</pre>
rmse_by_bin <- data.frame(bin = unique_bins, rmse = NA, count = NA) # Initialize results data frame
for (i in seq_along(unique_bins)) {
 bin_data <- data[data$bin == unique_bins[i], ] # Filter data for the current bin
  rmse_by_bin$rmse[i] <- sqrt(mean((bin_data$actual - bin_data$predicted)^2, na.rm = TRUE)) # Calculate
 rmse_by_bin$count[i] <- nrow(bin_data) # Count observations in the bin</pre>
# Display RMSE by bin
print(rmse_by_bin)
##
           bin
                    rmse count
## 1 [9e-05,1]
                     {\tt NaN}
                              0
## 2
        (1,10]
                              0
                     NaN
      (10,50] 4.759353
                            322
## 4 (50,100] 10.302785
# Testing the model on out of sample dataset
# obtain predicted values
test_predictions <- predict(trunc_damage_fit_reg, newdata = truncated_test)</pre>
# Create a data frame with actual, predicted, and bin labels
test_data <- data.frame(</pre>
 actual = truncated_test$damage_perc,
 predicted = test_predictions,
  bin = bin_labels
# Calculate RMSE per bin
unique_bins <- levels(data$bin) # Get unique bin labels</pre>
rmse_by_bin <- data.frame(bin = unique_bins, rmse = NA, count = NA) # Initialize results data frame
for (i in seq_along(unique_bins)) {
```

```
test_bin_data <- test_data[test_data$bin == unique_bins[i], ] # Filter data for the current bin
  rmse_by_bin$rmse[i] <- sqrt(mean((test_bin_data$actual - test_bin_data$predicted)^2, na.rm = TRUE)) #</pre>
  rmse_by_bin$count[i] <- nrow(test_bin_data) # Count observations in the bin</pre>
# Display RMSE by bin
print(rmse_by_bin)
           bin
                   rmse count
## 1 [9e-05,1]
                    {\tt NaN}
                            0
## 2
       (1,10]
                    {\tt NaN}
                            0
      (10,50] 15.45866
## 3
                          322
## 4 (50,100] 16.46157
                          74
saveRDS(trunc_damage_fit_reg, here("associational XGBOOST", "trunc_damage_fit_reg.rds"))
# set.seed(1234)
# tune_grid <- expand.grid(</pre>
   nrounds = c(200, 250, 275, 300, 350),
#
    max_depth = c(3, 6),
#
   eta = c(0.01, 0.05, 0.1, 0.2, 0.3),
#
   gamma = c(0, 1, 5, 10),
   colsample_bytree = c(0.5, 0.7, 0.8, 1.0),
#
   min\_child\_weight = c(1, 3, 5, 10),
   subsample = c(0.5, 0.7, 0.8, 1.0)
```