# Term Project 390.4- 2019

#### R. Markdown

```
pacman::p_load(dplyr, tidyr, ggplot2, magrittr, stringr, mlr)
housing_data = read.csv("housing_data_2016_2017.csv")

##Delete features that are irrelevant to sale price
housing_data %<>%
select(-c(HITId, HITTypeId, Title, Description, Keywords, Reward, CreationTime, MaxAssignments, Requ
```

### Clean Data

```
housing_data %<>%
  mutate( zip_code = str_extract(full_address_or_zip_code, "[0-9]{5}"))
housing_data %<>%
  mutate(dogs_allowed = ifelse(substr(housing_data$dogs_allowed, 1, 3) == "yes", 1, 0)) %>%
  mutate(cats_allowed = ifelse(substr(housing_data$cats_allowed, 1, 3) == "yes", 1, 0)) %>%
  mutate( pets_allowed = ifelse( cats_allowed + dogs_allowed > 0, 1, 0)) %>%
  mutate(coop_condo = factor(tolower(coop_condo)))
housing data %<>%
  select(-c(dogs_allowed, cats_allowed, fuel_type))
d = housing_data
d %<>%
  mutate(maintenance_cost = sjmisc::rec(maintenance_cost, rec = "NA = 0; else = copy")) %<>%
  mutate(common_charges = sjmisc::rec(common_charges, rec = "NA = 0; else = copy")) ##recode from NA to
# combine maintaince cost and common charges
d %<>%
  mutate( monthly_cost = common_charges + maintenance_cost)
d %<>%
  mutate(monthly_cost = sjmisc::rec(monthly_cost, rec = "0 = NA ; else = copy"))
## convert garage_exists feature to binary
d %<>%
  mutate(garage_exists = sjmisc::rec(garage_exists, rec = "NA = 0; else = copy")) ##recode from NA to
  mutate(garage_exists = sjmisc::rec(garage_exists, rec = " eys = 1; UG = 1 ; Underground = 1; yes = 1
d %<>%
  select(-c(maintenance_cost , common_charges, model_type))
##Change variable types
d %<>%
  mutate( dining_room_type = as.factor(dining_room_type)) %>%
  mutate(garage_exists = as.character(garage_exists)) %>%
  mutate(garage_exists = as.numeric(garage_exists)) %>%
  mutate( parking_charges = as.character(parking_charges)) %>%
  mutate( parking_charges = as.numeric(parking_charges)) %>%
  mutate(sale_price = as.character(sale_price)) %>%
  mutate(sale_price = as.numeric(sale_price)) %>%
  mutate(total_taxes = as.character(total_taxes)) %>%
```

```
mutate(total_taxes = as.numeric(total_taxes)) %>%
  mutate(price_persqft = listing_price_to_nearest_1000 / sq_footage)
## Warning: NAs introduced by coercion
## Warning: NAs introduced by coercion
## Warning: NAs introduced by coercion
#Added latitude and longitude features using ggmap
#Already run and included in the data
#pacman::p load(qqmap)
#d %<>%
# mutate(lat = qeocode(full_address_or_zip_code)$lat, lon = #qeocode(full_address_or_zip_code)$lon )
#qeocoordinates for relevant LIRR stations
lirr_coord = read.csv("coord.csv")
RAD_EARTH = 3958.8
degrees_to_radians = function(angle_degrees){
  for(i in 1:length(angle_degrees))
    angle_degrees[i] = angle_degrees[i]*pi/180
  return(angle_degrees)
compute_globe_distance = function(destination, origin){
  destination rad = degrees to radians(destination)
  origin_rad = degrees_to_radians(origin)
  delta_lat = destination_rad[1] - origin_rad[1]
  delta_lon = destination_rad[2] - origin_rad[2]
 h = (sin(delta_lat/2))^2 + cos(origin_rad[1]) * cos(destination_rad[1]) * (sin(delta_lon/2))^2
  central angle = 2 * asin(sqrt(h))
 return(RAD_EARTH * central_angle)
#find the closest LIRR station and compute distance
shortest_lirr_distance = function(all_lirr_coords, house_coords){
  shortest_dist = Inf
  for (i in 1: nrow(all_lirr_coords)){
   ith_lirr = c(all_lirr_coords$lat[i], all_lirr_coords$lon[i])
   new_dist = compute_globe_distance(ith_lirr, house_coords)
   if( new_dist < shortest_dist){</pre>
      shortest_dist = new_dist
   }
  }
  return(shortest_dist)
d %<>%
 rowwise() %>%
  mutate(shortest dist = shortest lirr distance(lirr coord, c(lat, lon)) )
#makes any other addresses redundant
d %<>%
  select(-c(zip_code, full_address_or_zip_code, listing_price_to_nearest_1000))
We are trying to predict sale_price. So let's section our dataset:
####CREATE A COLUMN ID
d %<>%
```

```
ungroup(d) %>%
mutate(id = 1 : 2230)
d %<>%
mutate(total_taxes = ifelse(d$total_taxes < 1000, NA, total_taxes))
real_y = data.frame(d$id, d$sale_price)
real_d = subset(d, (!is.na(d$sale_price)))
fake_d = subset(d, (is.na(d$sale_price)))
real_d$sale_price = NULL
fake_d$sale_price = NULL</pre>
```

#Split the data that has y into train and test sets

```
train_indices = sample(1 : nrow(real_d), nrow(real_d)*4/5)
training_data = real_d[train_indices, ]
testing_data = real_d[-train_indices, ]
X = rbind(training_data, testing_data, fake_d)
```

#Let's first create a matrix with p columns that represents missingness

```
M = tbl_df(apply(is.na(X), 2, as.numeric))
colnames(M) = paste("is_missing_", colnames(X), sep = "")
```

#Some of these missing indicators are collinear because they share all the rows they are missing on. Let's filter those out:

```
M = tbl_df(t(unique(t(M))))
```

#Some featuers did not have missingness so let's remove them:

```
M %<>% select_if(function(x){sum(x) > 0})
```

#Now let's impute missing data using the package. we cannot fit RF models to the entire dataset (it's 26,000! observations) so we will sample 5 for X1 and for each of the trees and then average. That will be good enough.

```
pacman::p_load(missForest)
Ximp = missForest(data.frame(X), sampsize = rep(172, ncol(X)))$ximp
```

```
##
     missForest iteration 1 in progress...
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
##
    missForest iteration 2 in progress...
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
    missForest iteration 3 in progress...
##
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
##
    missForest iteration 4 in progress...
```

```
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
    missForest iteration 5 in progress...
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
    missForest iteration 6 in progress...
##
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
    missForest iteration 7 in progress...
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
##
    missForest iteration 8 in progress...
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
    missForest iteration 9 in progress...
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry =
## mtry, : The response has five or fewer unique values. Are you sure you want
## to do regression?
## done!
Ximp %<>%
 arrange(id)
Xnew = data.frame(cbind(Ximp, M, real y))
Xnew %<>%
 mutate(price = d.sale_price) %>%
 select(-c(id, d.id, d.sale_price))
linear_mod_impute_and_missing_dummies = lm(price ~ ., data = Xnew)
```

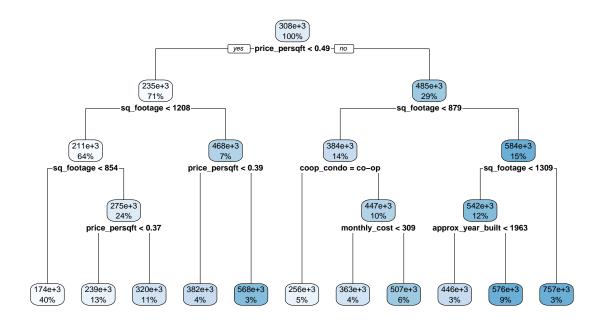
### REMOVING MISSING Y SECTION

```
Data = Xnew
### sale price is our imputed Y
Y = Data$price
Data %<>%
  filter(!is.na(price)) %>%
  select(-price)
```

```
Xtrain = Data[1:422, ]
Xtest = Data[423:528, ]
Ytrain = Y[1:422]
Ytest = Y[423:528]
dtrain = cbind(Xtrain, Ytrain) ## combine x train with y train, x test with y test
dtest = cbind(Xtest, Ytest)
```

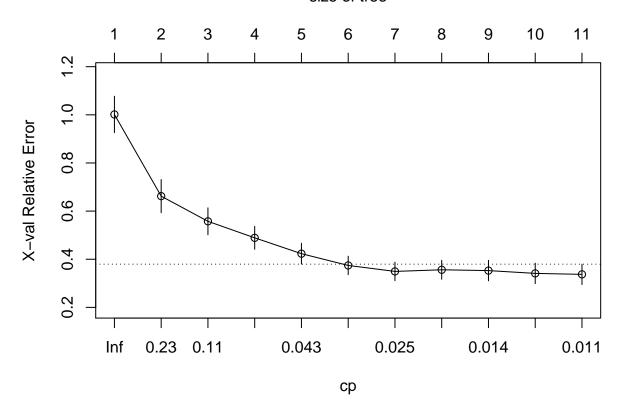
## Dropping colinear features

```
Xtrain %<>%
  select(-c(is_missing_num_total_rooms, is_missing_num_bedrooms, is_missing_price_persqft))
##Linear Regression
linear = lm(Ytrain ~ ., data = Xtrain)## simple linear model
##Linear Model Errors
yhat = predict(linear, Xtest)
e = yhat - Ytest
sqrt(sum(e^2) / nrow(Xtest))
## [1] 84480.55
#REGRESSION TREE
pacman::p_load(rsample)#data spliting
pacman::p_load(rpart) #performing reg tree
pacman::p_load(rpart.plot) #ploting reg tree
pacman::p_load(ipred) #bagging
pacman::p_load(caret) #bagging
m1 = rpart(
 formula = Ytrain ~ .,
 data = Xtrain,
 method = "anova"
rpart.plot(m1)
```



# plotcp(m1)

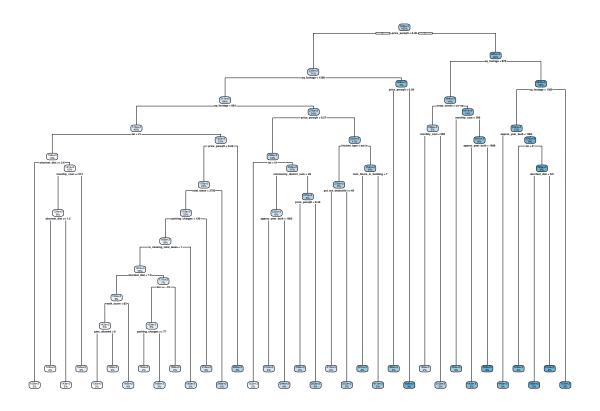
# size of tree



```
yhat = predict(m1, Xtest)
e = yhat - Ytest
sqrt(sum(e^2)/106)

## [1] 124680.6

m2 <- rpart(
    formula = Ytrain ~ .,
    data = Xtrain,
    method = "anova",
    control = list(cp = 0, xval = 10)
)
rpart.plot(m2)</pre>
```



# plotcp(m2)

## size of tree

```
1
                  3 5 7 9 11
                                          14
                                                 17
                                                       20
                                                              23
                                                                     26
                                                                           29
                                                                                  33
      1.0
X-val Relative Error
     0.8
     9.0
     0.4
     0.2
                                                0.0031
             Inf 0.068 0.017
                                     0.0089
                                                           0.0014 0.00047
                                                                                       0
                                                  ср
```

```
yhat = predict(m2, Xtest)
e = yhat - Ytest
sqrt(sum(e^2)/106)
## [1] 107729.4
jpeg(file = "save_m2.jpeg")
\#\# Tuning
m3 <- rpart(
    formula = Ytrain ~ .,
           = Xtrain,
    data
    method = "anova",
    control = list(minsplit = 10, maxdepth = 12, xval = 10)
yhat = predict(m3, Xtest)
e = yhat - Ytest
sqrt(sum(e^2)/106)
## [1] 124680.6
m3$cptable
              CP nsplit rel error
                                     xerror
## 1 0.41178122
                      0 1.0000000 1.0034543 0.07589990
## 2 0.12693151
                      1 0.5882188 0.6643831 0.06115611
                      2 0.4612873 0.5923246 0.05845056
## 3 0.09283490
```

3 0.3684524 0.4999903 0.04805964

## 4 0.04979016

```
## 5 0.03740845 4 0.3186622 0.4470845 0.04635923
## 6 0.03384382 5 0.2812538 0.3857698 0.04100711
## 7 0.01825815
                     6 0.2474099 0.3802651 0.04042839
## 8 0.01556267
                      7 0.2291518 0.3792111 0.03781015
## 9 0.01250920
                      8 0.2135891 0.3636770 0.03611701
## 10 0.01226119
                      9 0.2010799 0.3586581 0.03924726
## 11 0.01000000
                     10 0.1888187 0.3423144 0.03858264
# function to get optimal cp
get_cp <- function(x) {</pre>
        <- which.min(x$cptable[, "xerror"])</pre>
  cp <- x$cptable[min, "CP"]</pre>
# function to get minimum error
get_min_error <- function(x) {</pre>
       <- which.min(x$cptable[, "xerror"])</pre>
  xerror <- x$cptable[min, "xerror"]</pre>
}
optimal_tree <- rpart(</pre>
    formula = Ytrain ~
           = Xtrain,
    data
    method = "anova",
    control = list(minsplit = 11, maxdepth = 8, cp = 0.01)
pred <- predict(optimal_tree, newdata = Xtrain)</pre>
RMSE(pred = pred, obs = Ytrain)
## [1] 76766.17
##RANDOM FOREST
m1 <- randomForest(</pre>
  formula = Ytrain ~ .,
        = Xtrain
  data
)
m1
##
## Call:
##
   randomForest(formula = Ytrain ~ ., data = Xtrain)
##
                   Type of random forest: regression
                          Number of trees: 500
##
## No. of variables tried at each split: 11
##
##
              Mean of squared residuals: 5019464088
##
                        % Var explained: 83.92
which.min(m1$mse)
## [1] 281
# RMSE of this optimal random forest
sqrt(m1$mse[which.min(m1$mse)])
## [1] 70680.71
features <- setdiff(names(Xtrain), Ytrain)</pre>
set.seed(1989)
```

## -0.03117163 0.01 ## 0.02312618 0.01 ## 0.04316821 0.01 ## -0.002202444 0.01

