### Lab Report 2

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```
# Insert necessary packages
library('tidyverse')
library('gridExtra')
library('ISLR')
library('plotly')
library('caret')
library('MASS')
library('glmnet')
library('gam')
library('splines')
```

#### **Question 1: Nonlinear Regression**

#### 1.1. Process your data

```
# Read the data
diamonds <- read.csv('data/diamonds.csv')

# Remove all rows that contain NA
diamonds <- na.omit(diamonds)

# Downsample data
diamonds <- diamonds[sample(5000),]

# Select columns
diamonds <- diamonds[, c('X', 'carat', 'cut', 'depth', 'table', 'price', 'x', 'y', 'z')]

# Convert string column to categorical column

# Convert diamond string variables to categorical
diamonds$cut <- as.factor(diamonds$cut)

dim(diamonds)

## [1] 5000 9

head(diamonds)</pre>
```

```
##
           X carat
                          cut depth table price
                                                     Х
          62
              0.91
                                62.7
                                        57
## 62
                        Ideal
                                            3557 6.15 6.19 3.87
## 1697 1697
              0.90 Very Good
                                62.5
                                        57
                                            3209 6.12 6.20 3.85
  3478 3478
              1.00
                      Premium
                                61.3
                                            3584 6.45 6.40 3.94
  4502 4502
              0.72
                        Ideal
                                62.4
                                            3084 5.76 5.72 3.58
## 408
         408
              0.74
                        Ideal
                                62.0
                                        54
                                            3694 5.85 5.81 3.62
## 213
         213
              0.91
                               64.3
                                        57
                                            3632 6.00 6.04 3.87
                         Good
```

#### 1.3. Visualize the data

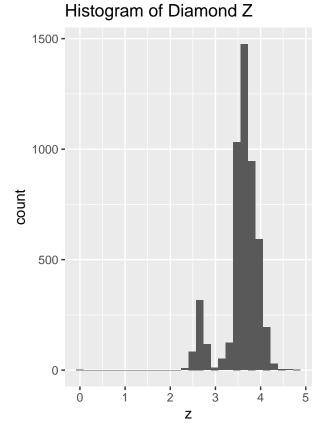
It was decided to do step 1.3. before step 1.2 because before splitting the input data into train and test, it makes more sense to apply the necessary transformations first.

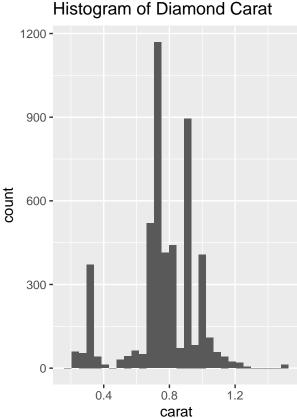
```
histo1 <- ggplot(data = diamonds) +
  geom_histogram(aes(x = z)) +
  ggtitle("Histogram of Diamond Z")

histo2 <- ggplot(data = diamonds) +
  geom_histogram(aes(x = carat)) +
  ggtitle("Histogram of Diamond Carat")

grid.arrange(histo1, histo2, ncol=2)</pre>
```

```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```





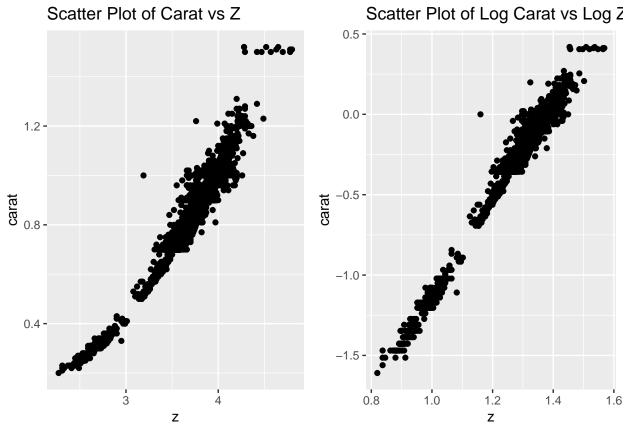
```
# Remove outliers
diamonds <- diamonds[(diamonds$z>1),]

# Apply logarithmic transform
numeric.cols <- summarize_all(diamonds, is.numeric) %>% unlist()
logDiamonds <- diamonds
logDiamonds[,numeric.cols] <- log(diamonds[,numeric.cols])

scatter1 <- ggplot(data = diamonds) +
    geom_point(aes(x = z, y=carat)) +
    ggtitle("Scatter Plot of Carat vs Z")

scatter2 <- ggplot(data = logDiamonds) +
    geom_point(aes(x = z, y=carat)) +
    geom_point(aes(x = z, y=carat)) +
    ggtitle("Scatter Plot of Log Carat vs Log Z")

grid.arrange(scatter1, scatter2, ncol=2)</pre>
```



removing some outliers, it can be clearly seen that as z is increased, the carat is also increased. In terms of linearity, it is slightly non-linear and showing exponential form. When applying log transform to all the carat and x values, then the relationship is clearly linear.

After

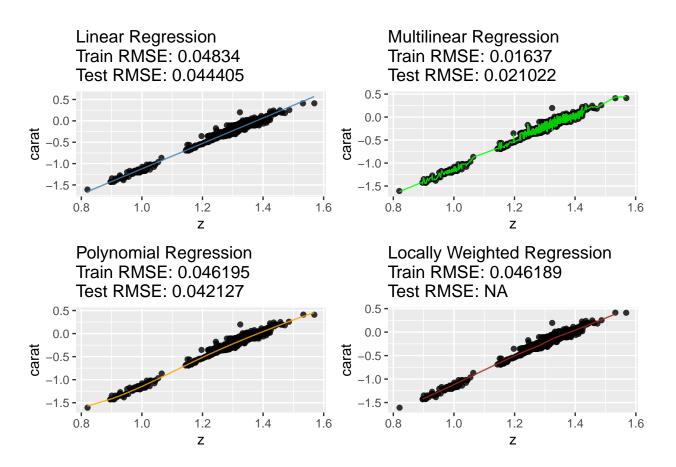
#### 1.2. Train / Test Split

```
set.seed(156)
train_inds <- sample(1:nrow(logDiamonds), floor(nrow(logDiamonds)*0.8))
train <- logDiamonds[ train_inds, ]
test <- logDiamonds[-train_inds, ]
cat('train: ', nrow(train), ', test: ', nrow(test))
## train: 3997 , test: 1000</pre>
```

#### 1.4. Fit 4 Models

```
# Linear Regression
fit.lm <- lm(carat ~ z, data = train)</pre>
# Predictions
preds.lm_train <- predict(fit.lm, train)</pre>
preds.lm_test <- predict(fit.lm, test)</pre>
# RMSE
rmse.lm_train <- RMSE(preds.lm_train, train$carat)</pre>
rmse.lm_test <- RMSE(preds.lm_test, test$carat)</pre>
# Multilinear Regression
fit.mlm <- lm(carat ~ ., data = train)</pre>
# Predictions
preds.mlm_train <- predict(fit.mlm, train)</pre>
preds.mlm_test <- predict(fit.mlm, test)</pre>
# RMSE
rmse.mlm_train <- RMSE(preds.mlm_train, train$carat)</pre>
rmse.mlm_test <- RMSE(preds.mlm_test, test$carat)</pre>
# Polynomial Regression
fit.poly \leftarrow lm(carat \sim poly(z,6), data = train)
# Predictions
preds.poly_train <- predict(fit.poly, train)</pre>
preds.poly_test <- predict(fit.poly, test)</pre>
# RMSE
rmse.poly_train <- RMSE(preds.poly_train, train$carat)</pre>
rmse.poly_test <- RMSE(preds.poly_test, test$carat)</pre>
# Locally Weighted Regression
```

```
fit.wlm <- loess(carat ~ z, data=train)</pre>
# Predictions
preds.wlm_train <- predict(fit.wlm, train)</pre>
preds.wlm_test <- predict(fit.wlm, test)</pre>
# RMSE
rmse.wlm_train <- RMSE(preds.wlm_train, train$carat)</pre>
rmse.wlm_test <- RMSE(preds.wlm_test, test$carat)</pre>
# Plots
linear_plot <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.lm_test), colour = 'steelblue', alpha=.8) +
  ggtitle(paste0('Linear Regression \nTrain RMSE: ',round(rmse.lm_train, 6), "\nTest RMSE: ", round
multilinear_plot <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.mlm_test), colour = 'green', alpha=.8) +
  ggtitle(paste0('Multilinear Regression \nTrain RMSE: ',round(rmse.mlm_train, 6), "\nTest RMSE: '
poly_plot <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.poly_test), colour = 'orange', alpha=.8) +
  ggtitle(paste0('Polynomial Regression \nTrain RMSE: ',round(rmse.poly_train, 6), "\nTest RMSE: '
weighted_plot <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.wlm_test), colour = 'firebrick', alpha=.8) +
  ggtitle(paste0('Locally Weighted Regression \nTrain RMSE: ',round(rmse.wlm_train, 6), "\nTest RI
grid.arrange(linear_plot, multilinear_plot, poly_plot, weighted_plot, ncol=2)
## Warning: Removed 2 row(s) containing missing values (geom_path).
```



Based off of Train RMSE, the order of models from best (lowest RMSE) to worst (highest RMSE) is: 1. Multilinear Regression 2. Polynomial Regression 3. Locally Weighted Regression 4. Linear Regression

Based off of the Test RMSE, the order of models from best (lowest RMSE) to worst (highest RMSE) is:
1. Multilinear Regression 2. Polynomial Regression 3. Locally Weighted Regression 4. Linear Regression
The order of models from best to worst did not change when ordering by train RMSE or test RMSE.

#### 1.5. Cross Validation

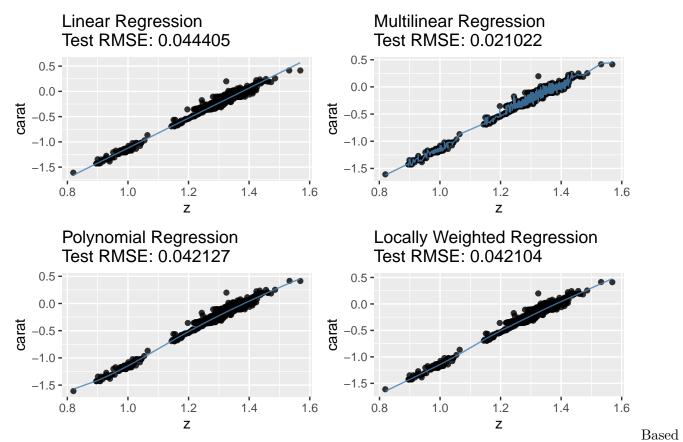
```
ctrl <- trainControl(method = "repeatedcv", number = 10, repeats=5)

# Linear regression
cv_fit.lm <- train(
  form = carat ~ z,
  data = train,
  method = "lm",
  trControl = ctrl
)

preds.cv_lm <- predict(cv_fit.lm,test)
rmse.cv_lm <- RMSE(preds.cv_lm,test$carat)

# Multilinear regression
cv_fit.mlm <- train(</pre>
```

```
form = carat ~ .,
  data = train,
 method = "lm",
 trControl = ctrl
preds.cv_mlm <- predict(cv_fit.mlm,test)</pre>
rmse.cv_mlm <- RMSE(preds.cv_mlm,test$carat)</pre>
# Polynomial regression
cv_fit.poly <- train(</pre>
 form = carat \sim poly(z, 6),
 data = train,
 method = "lm",
 trControl = ctrl
)
preds.cv_poly <- predict(cv_fit.poly,test)</pre>
rmse.cv_poly <- RMSE(preds.cv_poly,test$carat)</pre>
# Locally weighted regression
cv_fit.wlm <- train(</pre>
 form = carat \sim z,
 data = train,
 method = "gamLoess",
 trControl = ctrl
)
preds.cv_wlm <- predict(cv_fit.wlm,test)</pre>
rmse.cv_wlm <- RMSE(preds.cv_wlm,test$carat)</pre>
cv_linear_plot <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.cv_lm), colour = 'steelblue', alpha=.8) +
  ggtitle(paste0('Linear Regression \nTest RMSE: ',round(rmse.cv_lm, 6)))
cv_multilinear_plot <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.cv_mlm), colour = 'steelblue', alpha=.8) +
  ggtitle(paste0('Multilinear Regression \nTest RMSE: ',round(rmse.cv_mlm, 6)))
cv_poly_plot <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.cv_poly), colour = 'steelblue', alpha=.8) +
  ggtitle(paste0('Polynomial Regression \nTest RMSE: ',round(rmse.cv_poly, 6)))
cv_wlm_plot <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.cv_wlm), colour = 'steelblue', alpha=.8) +
```



off of the Test RMSE, the order of models from best (lowest RMSE) to worst (highest RMSE) is: 1. Multilinear Regression 2. Locally Weighted Regression 3. Polynomial Regression 4. Linear Regression

The order of models have changed when adding k fold cross validation. The Locally Weighted Regression is now second best and moved in front of Polynomial Regression. Therefore, by applying cross validation, more precise performance metrics have been measured for the models.

#### 1.6. Shrinkage

```
x_train <- model.matrix(carat ~ z,train)
x_train_multi <- model.matrix(carat ~ .,train)
x_train_poly <- model.matrix(carat ~ poly(z,6),train)
y_train <- train$carat

x_test <- model.matrix(carat ~ z,test)
x_test_multi <- model.matrix(carat ~ .,test)
x_test_poly <- model.matrix(carat ~ .,test)

# Ridge
fit.ridge_lm <- cv.glmnet(x_train,y_train,alpha=0, nfolds = 10)
fit.ridge_lm <- glmnet(x_train,y_train,alpha=0, lambda=fit.ridge_lm$lambda.min)</pre>
```

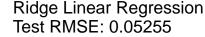
```
fit.ridge_mlm <- cv.glmnet(x_train_multi,y_train,alpha=0, nfolds = 10)</pre>
fit.ridge_mlm <- glmnet(x_train_multi,y_train,alpha=0, lambda=fit.ridge_mlm$lambda.min)</pre>
fit.ridge_poly <- cv.glmnet(x_train_poly,y_train,alpha=0, nfolds = 10)</pre>
fit.ridge_poly <- glmnet(x_train_poly,y_train,alpha=0, lambda=fit.ridge_poly$lambda.min)</pre>
preds.ridge_lm <- predict(fit.ridge_lm, x_test)</pre>
preds.ridge_mlm <- predict(fit.ridge_mlm, x_test_multi)</pre>
preds.ridge_poly <- predict(fit.ridge_poly, x_test_poly)</pre>
rmse.ridge_lm <- RMSE(preds.ridge_lm, test$carat)</pre>
rmse.ridge_mlm <- RMSE(preds.ridge_mlm, test$carat)</pre>
rmse.ridge_poly <- RMSE(preds.ridge_poly, test$carat)</pre>
# Lasso
fit.lasso_lm <- cv.glmnet(x_train,y_train,alpha=1, nfolds = 10)</pre>
fit.lasso_lm <- glmnet(x_train,y_train,alpha=1, lambda=fit.lasso_lm$lambda.min)</pre>
fit.lasso_mlm <- cv.glmnet(x_train_multi,y_train,alpha=1, nfolds = 10)</pre>
fit.lasso_mlm <- glmnet(x_train_multi,y_train,alpha=1, lambda=fit.lasso_mlm$lambda.min)</pre>
fit.lasso_poly <- cv.glmnet(x_train_poly,y_train,alpha=1, nfolds = 10)</pre>
fit.lasso_poly <- glmnet(x_train_poly,y_train,alpha=1, lambda=fit.lasso_poly$lambda.min)
preds.lasso_lm <- predict(fit.lasso_lm, x_test)</pre>
preds.lasso_mlm <- predict(fit.lasso_mlm, x_test_multi)</pre>
preds.lasso_poly <- predict(fit.lasso_poly, x_test_poly)</pre>
rmse.lasso_lm <- RMSE(preds.lasso_lm, test$carat)</pre>
rmse.lasso_mlm <- RMSE(preds.lasso_mlm, test$carat)</pre>
rmse.lasso_poly <- RMSE(preds.lasso_poly, test$carat)</pre>
# Ridge
ridge_lm <-ggplot(test, aes(y=carat, x=z)) +</pre>
  geom point(alpha=.8, position = position jitter()) +
  geom_line(aes(y=preds.ridge_lm), colour = 'steelblue', alpha=.8) +
  ggtitle(paste0('Ridge Linear Regression \nTest RMSE: ',round(rmse.ridge_lm, 6)))
ridge_mlm <- ggplot(test, aes(y=carat, x=z)) +</pre>
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.ridge_mlm), colour = 'steelblue', alpha=.8) +
  ggtitle(paste0('Ridge MultiLinear Regression \nTest RMSE: ',round(rmse.ridge_mlm, 6)))
ridge_poly <- ggplot(test, aes(y=carat, x=z)) +
  geom_point(alpha=.8, position = position_jitter()) +
  geom_line(aes(y=preds.ridge_poly), colour = 'steelblue', alpha=.8) +
  ggtitle(paste0('Ridge Polynomial Regression \nTest RMSE: ',round(rmse.ridge_poly, 6)))
```

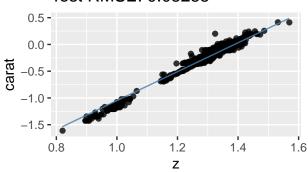
```
# Lasso
lasso_lm <-ggplot(test, aes(y=carat, x=z)) +
    geom_point(alpha=.8, position = position_jitter()) +
    geom_line(aes(y=preds.lasso_lm), colour = 'steelblue', alpha=.8) +
    ggtitle(paste0('Lasso Linear Regression \nTest RMSE: ',round(rmse.lasso_lm, 6)))

lasso_mlm <- ggplot(test, aes(y=carat, x=z)) +
    geom_point(alpha=.8, position = position_jitter()) +
    geom_line(aes(y=preds.lasso_mlm), colour = 'steelblue', alpha=.8) +
    ggtitle(paste0('Lasso MultiLinear Regression \nTest RMSE: ',round(rmse.lasso_mlm, 6)))

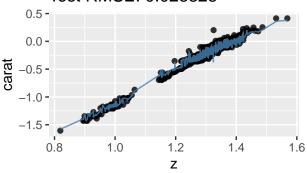
lasso_poly <- ggplot(test, aes(y=carat, x=z)) +
    geom_point(alpha=.8, position = position_jitter()) +
    geom_line(aes(y=preds.lasso_poly), colour = 'steelblue', alpha=.8) +
    ggtitle(paste0('Lasso Polynomial Regression \nTest RMSE: ',round(rmse.lasso_poly, 6)))

grid.arrange(ridge_lm, ridge_mlm, ridge_poly, ncol=2)</pre>
```

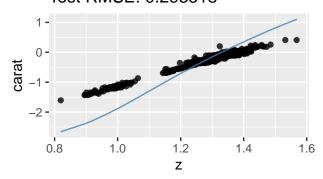




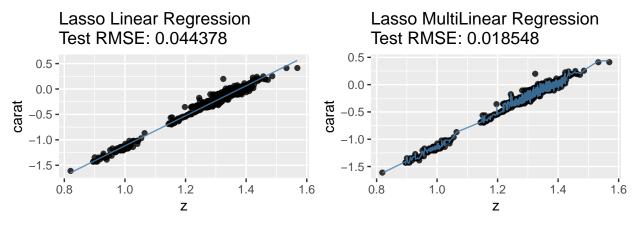
## Ridge MultiLinear Regression Test RMSE: 0.028328



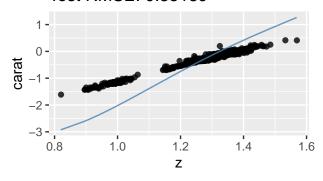
Ridge Polynomial Regression Test RMSE: 0.293518



grid.arrange(lasso\_lm, lasso\_mlm, lasso\_poly, ncol=2)



# Lasso Polynomial Regression Test RMSE: 0.35189

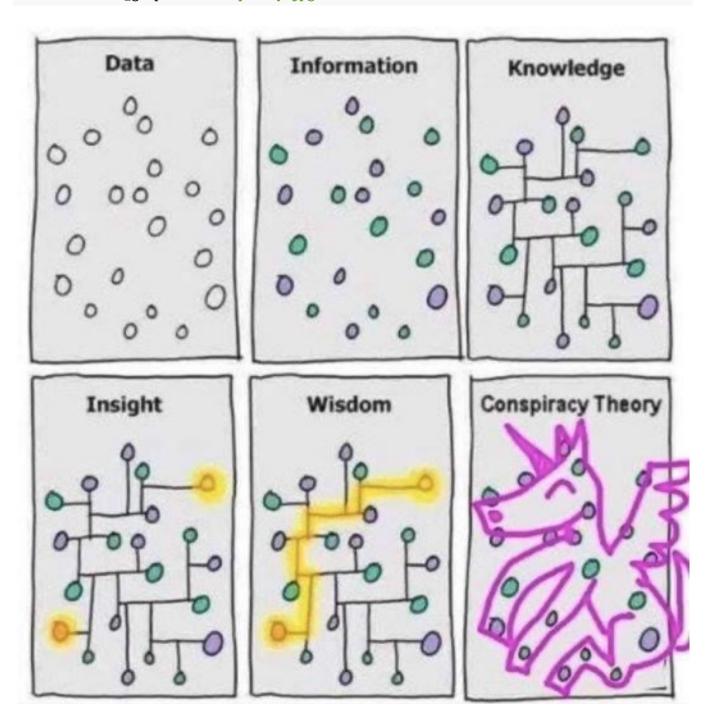


The

model that yielded the lowest RMSE was Lasso Multilinear Regression. Based off of the Test RMSE, the order of models from best (lowest RMSE) to worst (highest RMSE) is: 1. Lasso Multilinear Regression 2. Ridge Multilinear Regression 3. Lasso Linear Regression 4. Ridge Linear Regression 5. Ridge Polynomial Regression 6. Lasso Polynomial Regression

### Question 2

knitr::include\_graphics('conspiracy.jpg')



## Question 3

$$\theta := \theta + \frac{\alpha}{N} X^T (Y - \theta X)$$