

# Stat 520 Project

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## Load libraries and data

```
library(tidyverse)
```

```
Warning: package 'ggplot2' was built under R version 4.3.3
```

```
Warning: package 'tidyverse' was built under R version 4.3.3
```

```
Warning: package 'purrr' was built under R version 4.3.3
```

```
Warning: package 'dplyr' was built under R version 4.3.3
```

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr     1.1.4     v readr     2.1.4
vforcats    1.0.0     v stringr   1.5.0
v ggplot2   3.5.0     v tibble    3.2.1
v lubridate 1.9.2     v tidyverse 1.3.1
v purrr     1.0.2
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()    masks stats::lag()
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to becom
```

```
library(tidymodels)
```

```
Warning: package 'tidymodels' was built under R version 4.3.3
```

```
-- Attaching packages ----- tidyverse 1.2.0 --
v broom      1.0.5     v rsample     1.2.1
v dials       1.2.1     v tune        1.2.0
v infer       1.0.7     v workflows   1.1.4
v modeldata   1.3.0     v workflowsets 1.1.0
v parsnip     1.2.1     v yardstick   1.3.1
v recipes     1.0.10

Warning: package 'dials' was built under R version 4.3.3

Warning: package 'scales' was built under R version 4.3.3

Warning: package 'infer' was built under R version 4.3.3

Warning: package 'modeldata' was built under R version 4.3.3

Warning: package 'parsnip' was built under R version 4.3.3

Warning: package 'recipes' was built under R version 4.3.3

Warning: package 'rsample' was built under R version 4.3.3

Warning: package 'tune' was built under R version 4.3.3

Warning: package 'workflows' was built under R version 4.3.3

Warning: package 'workflowsets' was built under R version 4.3.3

Warning: package 'yardstick' was built under R version 4.3.3

-- Conflicts ----- tidyverse_conflicts() --
x scales::discard() masks purrr::discard()
x dplyr::filter()   masks stats::filter()
x recipes::fixed() masks stringr::fixed()
x dplyr::lag()     masks stats::lag()
x yardstick::spec() masks readr::spec()
x recipes::step()   masks stats::step()
* Use suppressPackageStartupMessages() to eliminate package startup messages
```

```
library(corrplot)
```

Warning: package 'corrplot' was built under R version 4.3.2

corrplot 0.92 loaded

```
library(rpart)
```

Attaching package: 'rpart'

The following object is masked from 'package:dials':

prune

```
library(rpart.plot)
library(randomForest)
```

randomForest 4.7-1.1

Type rfNews() to see new features/changes/bug fixes.

Attaching package: 'randomForest'

The following object is masked from 'package:dplyr':

combine

The following object is masked from 'package:ggplot2':

margin

```
library(caret)
```

Loading required package: lattice

Attaching package: 'caret'

```
The following objects are masked from 'package:yardstick':
```

```
precision, recall, sensitivity, specificity
```

```
The following object is masked from 'package:purrr':
```

```
lift
```

```
library(car)
```

```
Warning: package 'car' was built under R version 4.3.2
```

```
Loading required package: carData
```

```
Attaching package: 'car'
```

```
The following object is masked from 'package:dplyr':
```

```
recode
```

```
The following object is masked from 'package:purrr':
```

```
some
```

```
library(stats)
library(ranger)
```

```
Warning: package 'ranger' was built under R version 4.3.3
```

```
Attaching package: 'ranger'
```

```
The following object is masked from 'package:randomForest':
```

```
importance
```

```
library(factoextra)
```

```
Warning: package 'factoextra' was built under R version 4.3.2
```

```
Welcome! Want to learn more? See two factoextra-related books at https://goo.gl/ve3WBa
```

```
realtor <- read.csv("~/Truman_Stat_520/Project/realtor-data.zip.csv")
pollution <- read.csv("~/Truman_Stat_520/Project/pollution_us_2000_2016.csv")
```

## Clean Data

```
realtor2 <- realtor %>%
  select(city, state, price) %>%
  rename(City = city, State = state, Price = price) %>%
  group_by(City, State) %>%
  summarise(
    MeanPrice = mean(Price, na.rm = TRUE),
    MedianPrice = median(Price, na.rm = TRUE),
    MaxPrice = max(Price, na.rm = FALSE),
    .groups = 'drop'
  )

pollution2 <- pollution %>%
  select(State, City, NO2.AQI, O3.AQI, SO2.AQI, CO.AQI, Date.Local)

joined_data <- inner_join(realtor2, pollution2, by = c("City", "State"))
```

## Data Exploration

```
summary(joined_data)
```

City	State	MeanPrice	MedianPrice
Length:727649	Length:727649	Min. : 42765	Min. : 45000
Class :character	Class :character	1st Qu.: 204050	1st Qu.: 149000
Mode :character	Mode :character	Median : 335658	Median : 277500
		Mean : 621899	Mean : 383565
		3rd Qu.: 614855	3rd Qu.: 430000

MaxPrice	NO2.AQI	O3.AQI	SO2.AQI
Min. : 85000	Min. : 0.00	Min. : 0.00	Min. : 0.0
1st Qu.: 1200000	1st Qu.: 14.00	1st Qu.: 25.00	1st Qu.: 1.0
Median : 3950000	Median : 23.00	Median : 32.00	Median : 4.0
Mean : 16042976	Mean : 23.59	Mean : 36.06	Mean : 10.3
3rd Qu.: 14000000	3rd Qu.: 32.00	3rd Qu.: 42.00	3rd Qu.: 13.0
Max. : 135000000	Max. : 128.00	Max. : 207.00	Max. : 200.0
NA's : 54382			NA's : 363665
CO.AQI                      Date.Local			
Min. : 0.0	Length: 727649		
1st Qu.: 2.0	Class : character		
Median : 5.0	Mode : character		
Mean : 4.9			
3rd Qu.: 7.0			
Max. : 64.0			
NA's : 363830			

```

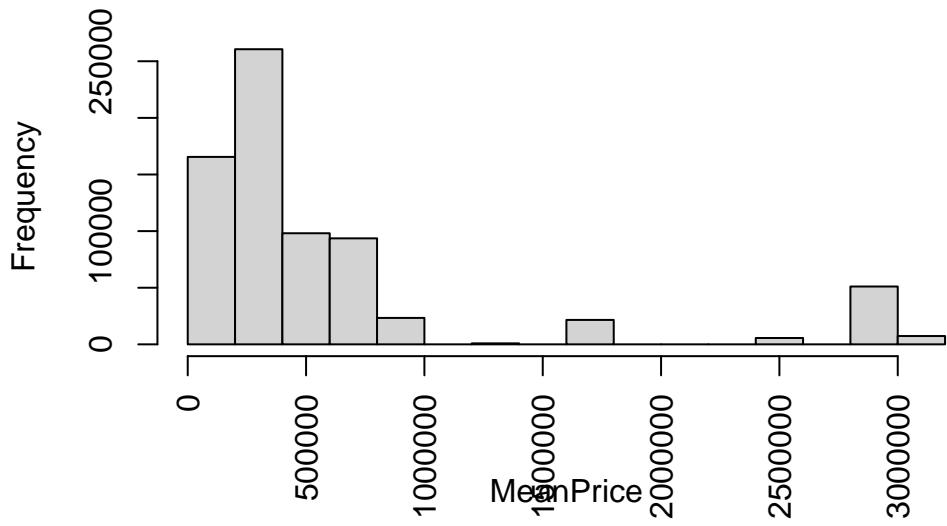
numeric_columns <- sapply(joined_data, is.numeric)

numeric_variable_names <- names(joined_data)[numeric_columns]

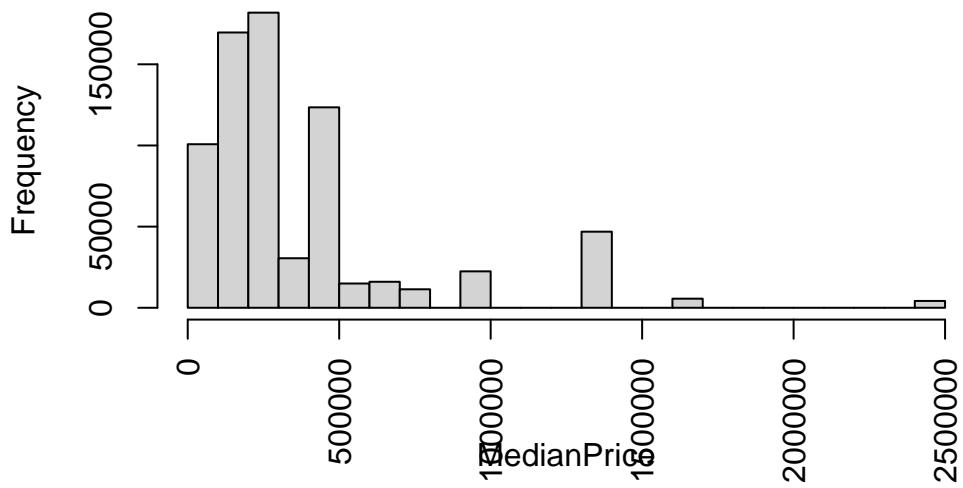
output <- lapply(numeric_variable_names, function(var_name) {
  hist(joined_data[[var_name]], main = paste("Histogram of", var_name), xlab = var_name,
       axis(1, las=2)
})

```

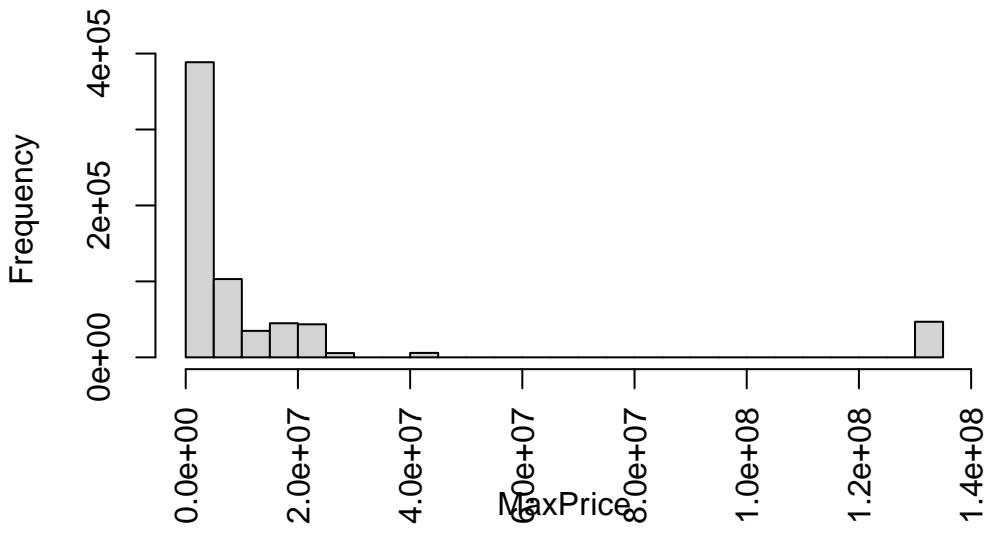
### Histogram of MeanPrice



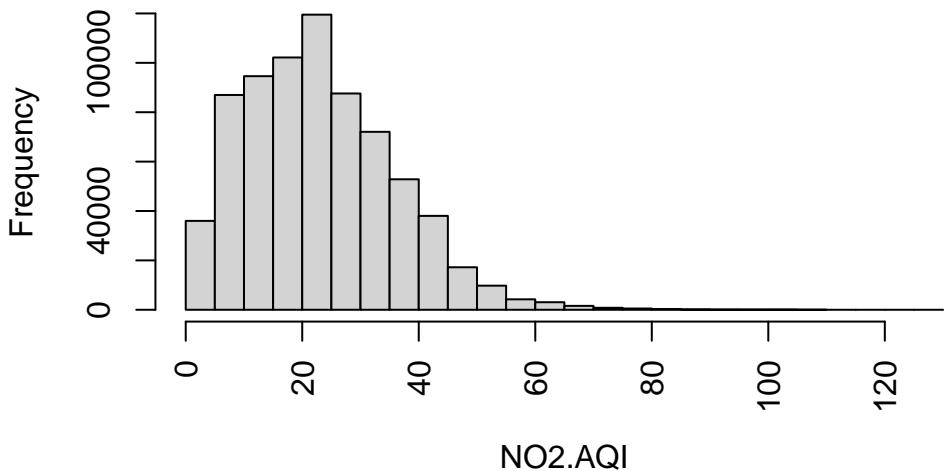
### Histogram of MedianPrice



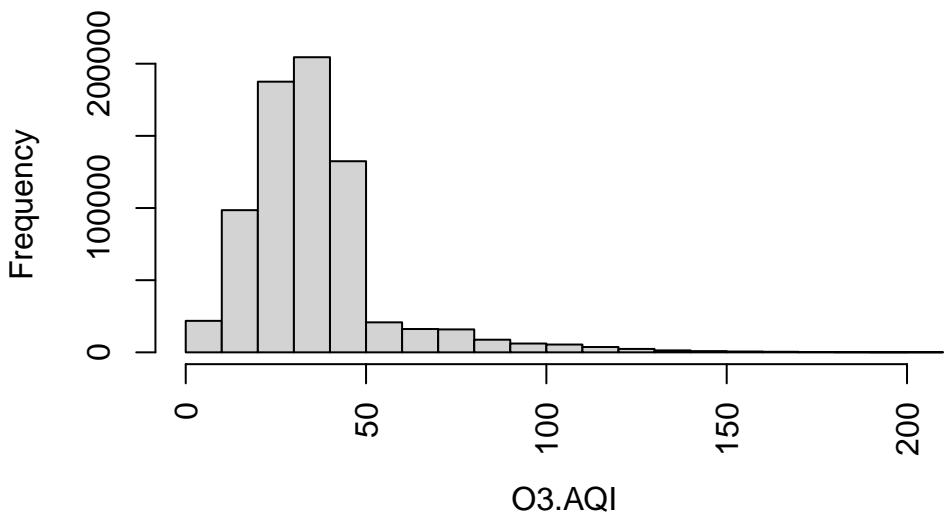
### Histogram of MaxPrice



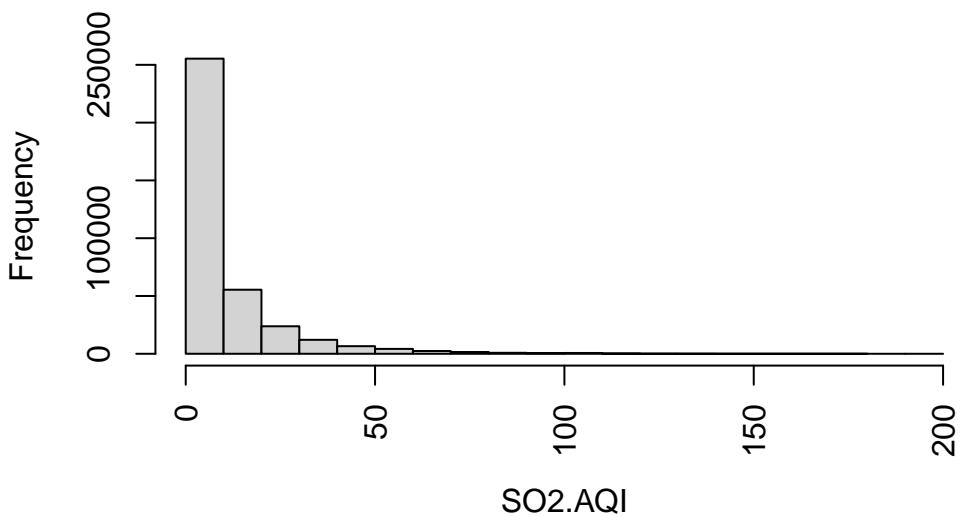
### Histogram of NO2.AQI



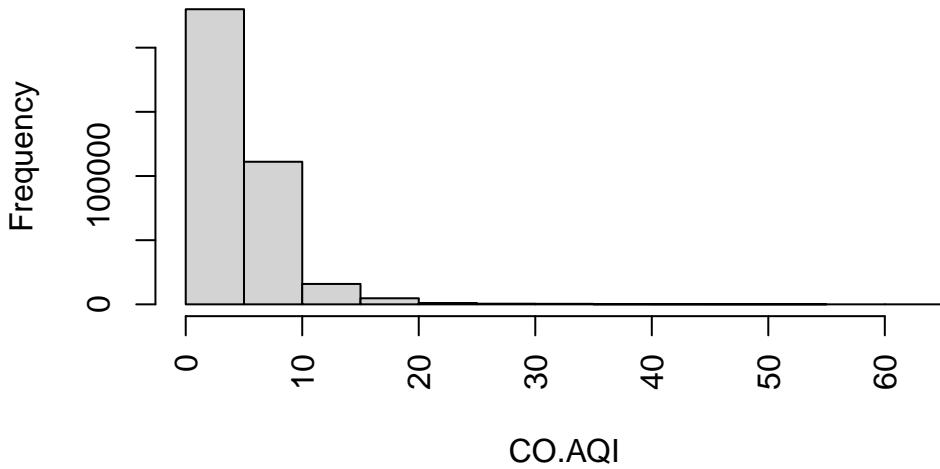
**Histogram of O3.AQI**



**Histogram of SO2.AQI**

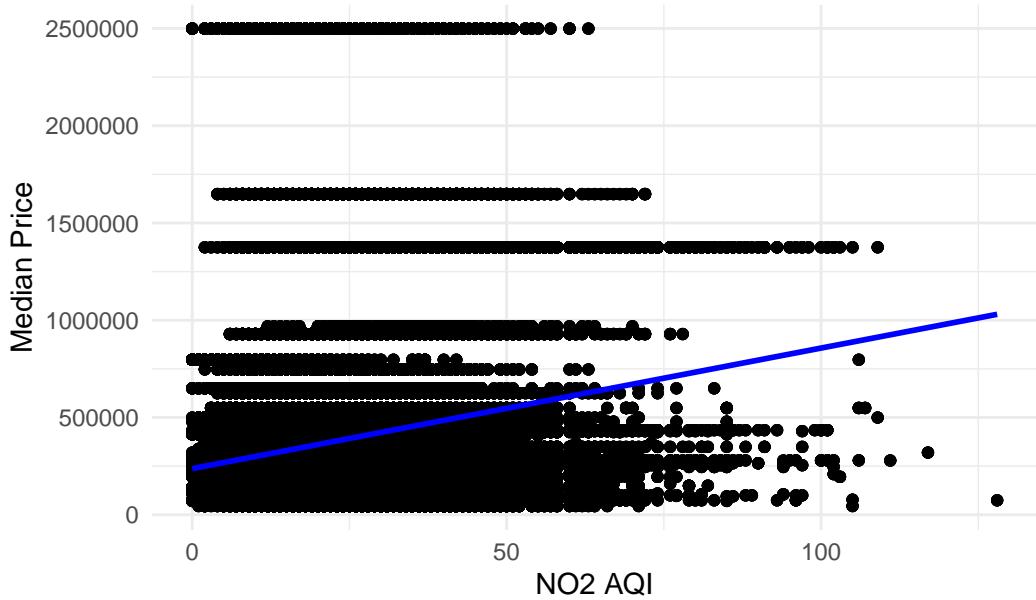


## Histogram of CO.AQI

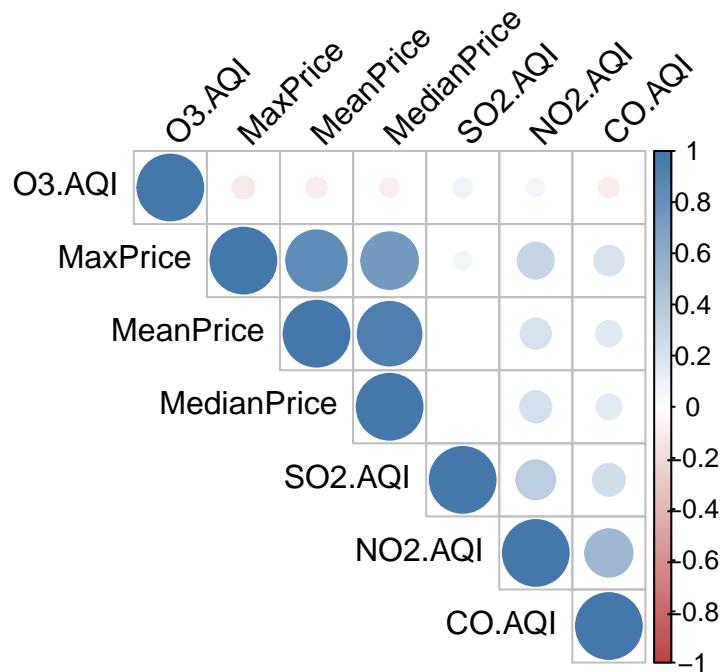


```
ggplot(joined_data, aes(x = NO2.AQI, y = MedianPrice)) +  
  geom_point() +  
  geom_smooth(method = "lm", col = "blue") +  
  theme_minimal() +  
  labs(title = "Scatter plot of Median Price vs NO2 AQI", x = "NO2 AQI", y = "Median Price")  
  
`geom_smooth()` using formula = 'y ~ x'
```

Scatter plot of Median Price vs NO2 AQI



```
cor_matrix <- cor(joined_data[, sapply(joined_data, is.numeric)], use = "complete.obs")
corrplot(cor_matrix, method = "circle", type = "upper", order = "hclust",
         tl.col = "black", tl.srt = 45,
         col = colorRampPalette(c("#BB4444", "white", "#4477AA"))(200))
```



```

joined_data$MaxPrice[is.na(joined_data$MaxPrice)] <- mean(joined_data$MaxPrice, na.rm = TRUE)

# Median imputation
joined_data$SO2.AQI[is.na(joined_data$SO2.AQI)] <- median(joined_data$SO2.AQI, na.rm = TRUE)
joined_data$CO.AQI[is.na(joined_data$CO.AQI)] <- median(joined_data$CO.AQI, na.rm = TRUE)

summary(joined_data)

```

City	State	MeanPrice	MedianPrice
Length:727649	Length:727649	Min. : 42765	Min. : 45000
Class :character	Class :character	1st Qu.: 204050	1st Qu.: 149000
Mode :character	Mode :character	Median : 335658	Median : 277500
		Mean : 621899	Mean : 383565
		3rd Qu.: 614855	3rd Qu.: 430000
		Max. : 3070828	Max. : 2499000
MaxPrice	NO2.AQI	O3.AQI	SO2.AQI
Min. : 85000	Min. : 0.00	Min. : 0.00	Min. : 0.000
1st Qu.: 1223600	1st Qu.: 14.00	1st Qu.: 25.00	1st Qu.: 4.000
Median : 4900000	Median : 23.00	Median : 32.00	Median : 4.000
Mean : 16042976	Mean : 23.59	Mean : 36.06	Mean : 7.144
3rd Qu.: 15986000	3rd Qu.: 32.00	3rd Qu.: 42.00	3rd Qu.: 4.000

```

Max.    :1350000000   Max.    :128.00   Max.    :207.00   Max.    :200.000
CO.AQI      Date.Local
Min.    : 0.000   Length:727649
1st Qu.: 5.000   Class :character
Median  : 5.000   Mode  :character
Mean    : 4.953
3rd Qu.: 5.000
Max.    :64.000

```

## Multiple Linear Regression

```

linear.model <- lm(MedianPrice ~ NO2.AQI + SO2.AQI + O3.AQI + CO.AQI, data = joined_data)
summary(linear.model)

```

Call:

```

lm(formula = MedianPrice ~ NO2.AQI + SO2.AQI + O3.AQI + CO.AQI,
  data = joined_data)

```

Residuals:

Min	1Q	Median	3Q	Max
-1010593	-221238	-111895	82665	2371699

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	292527.12	1308.63	223.54	<2e-16 ***
NO2.AQI	6642.54	37.42	177.51	<2e-16 ***
SO2.AQI	-1894.00	40.45	-46.82	<2e-16 ***
O3.AQI	-1704.19	21.72	-78.46	<2e-16 ***
CO.AQI	1880.68	174.34	10.79	<2e-16 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 371200 on 727644 degrees of freedom  
 Multiple R-squared: 0.05531, Adjusted R-squared: 0.05531  
 F-statistic: 1.065e+04 on 4 and 727644 DF, p-value: < 2.2e-16

```

# Make predictions
predictions <- predict(linear.model, newdata = joined_data)

```

```
# Actual values
actuals <- joined_data$MedianPrice

# Calculating the residuals
residuals <- predictions - actuals

# Calculating RMSE
rmse <- sqrt(mean(residuals^2))

# Calculating MAE
mae <- mean(abs(residuals))

# Print the results
cat("RMSE: ", rmse, "\n")
```

RMSE: 371212.5

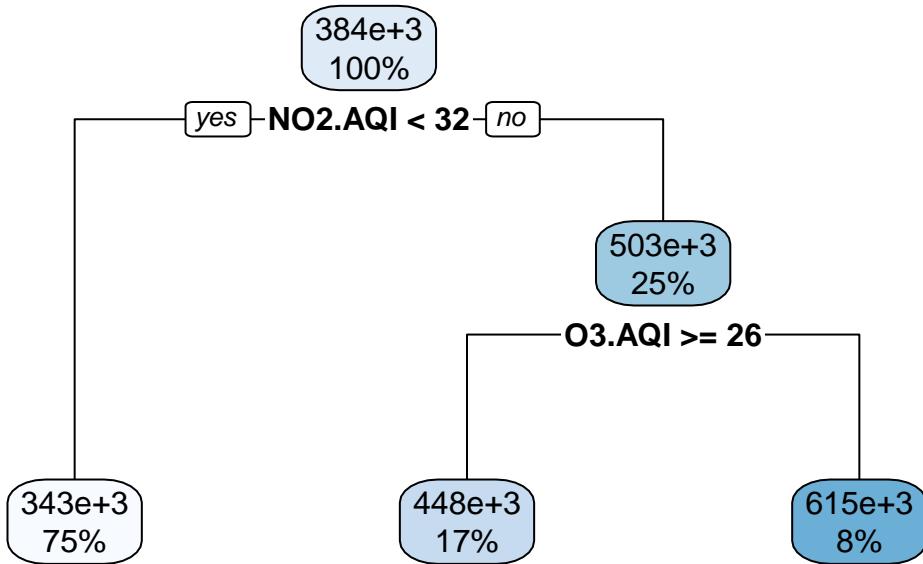
```
cat("MAE: ", mae, "\n")
```

MAE: 253927.4

## CART

```
cart.model <- rpart(MedianPrice ~ NO2.AQI + O3.AQI + SO2.AQI + CO.AQI,
                      data=joined_data,
                      method="anova")

rpart.plot(cart.model)
```



```

predictions <- predict(cart.model, newdata=joined_data)

actual_values <- joined_data$MedianPrice

# MAE
mae <- mean(abs(predictions - actual_values))
print(paste("Mean Absolute Error (MAE):", mae))

```

[1] "Mean Absolute Error (MAE): 253855.013891151"

```

# MSE
mse <- mean((predictions - actual_values)^2)
print(paste("Mean Squared Error (MSE):", mse))

```

[1] "Mean Squared Error (MSE): 139391762037.145"

```

# RMSE
rmse <- sqrt(mse)
print(paste("Root Mean Squared Error (RMSE):", rmse))

```

```
[1] "Root Mean Squared Error (RMSE): 373352.061782368"
```

```
# R-squared
rss <- sum((predictions - actual_values)^2)
tss <- sum((actual_values - mean(actual_values))^2)
rsquared <- 1 - (rss/tss)
print(paste("R-squared (R2):", rsquared))
```

```
[1] "R-squared (R2): 0.0443901148961726"
```

## Random Forest

```
rf.model <- ranger(MedianPrice ~ NO2.AQI + O3.AQI + SO2.AQI + CO.AQI,
                     data=joined_data,
                     method="anova")
```

```
Warning in ranger(MedianPrice ~ NO2.AQI + O3.AQI + SO2.AQI + CO.AQI, data =
joined_data, : Unused arguments: method
```

```
Growing trees.. Progress: 12%. Estimated remaining time: 3 minutes, 47 seconds.
Growing trees.. Progress: 25%. Estimated remaining time: 3 minutes, 8 seconds.
Growing trees.. Progress: 37%. Estimated remaining time: 2 minutes, 40 seconds.
Growing trees.. Progress: 49%. Estimated remaining time: 2 minutes, 10 seconds.
Growing trees.. Progress: 62%. Estimated remaining time: 1 minute, 37 seconds.
Growing trees.. Progress: 75%. Estimated remaining time: 1 minute, 4 seconds.
Growing trees.. Progress: 87%. Estimated remaining time: 32 seconds.
Growing trees.. Progress: 100%. Estimated remaining time: 0 seconds.
```

```
rf.predictions <- predict(rf.model, data=joined_data, type = "response")
```

```
Predicting.. Progress: 81%. Estimated remaining time: 7 seconds.
```

```
predicted_values <- rf.predictions$predictions

# Actual values
actual_values <- joined_data$MedianPrice
```

```

# MAE
mae <- mean(abs(predicted_values - actual_values))
print(paste("Mean Absolute Error (MAE):", mae))

[1] "Mean Absolute Error (MAE): 214292.857314077"

# MSE
mse <- mean((predicted_values - actual_values)^2)
print(paste("Mean Squared Error (MSE):", mse))

[1] "Mean Squared Error (MSE): 104828990677.271"

# RMSE
rmse <- sqrt(mse)
print(paste("Root Mean Squared Error (RMSE):", rmse))

[1] "Root Mean Squared Error (RMSE): 323773.054279183"

# R-squared
rss <- sum((predicted_values - actual_values)^2)
tss <- sum((actual_values - mean(actual_values))^2)
rsquared <- 1 - (rss/tss)
print(paste("R-squared (R2):", rsquared))

[1] "R-squared (R2): 0.281337589304867"

```

## PCA

```

# Extract predictor variables
predictors <- joined_data[, c("NO2.AQI", "SO2.AQI", "O3.AQI", "CO.AQI")]

# Perform PCA on the predictors
pca_results <- prcomp(predictors, center = TRUE, scale. = TRUE) # Standardizing variables
summary(pca_results)

```

```

Importance of components:
PC1      PC2      PC3      PC4
Standard deviation   1.2260  1.0208  0.9308  0.7671
Proportion of Variance 0.3758  0.2605  0.2166  0.1471
Cumulative Proportion 0.3758  0.6363  0.8529  1.0000

# Extracting the scores of the first two PCs
pc_scores <- pca_results$x[, 1:2]

# Prepare the data for regression
# Combine the PC scores with the MedianPrice
regression_data <- data.frame(cbind(pc_scores, MedianPrice = joined_data$MedianPrice))

# Linear Regression using the principal components to predict MedianPrice
model <- lm(MedianPrice ~ ., data = regression_data)
summary(model)

```

Call:  
`lm(formula = MedianPrice ~ ., data = regression_data)`

Residuals:

Min	1Q	Median	3Q	Max
-1243530	-222229	-115389	75025	2321007

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	383565.0	440.3	871.20	<2e-16 ***
PC1	47140.2	359.1	131.27	<2e-16 ***
PC2	-37676.2	431.3	-87.36	<2e-16 ***
---				
Signif. codes:	0 '***'	0.001 '**'	0.01 '*'	0.05 '.'
	0.1 '	' 1		

Residual standard error: 375600 on 727646 degrees of freedom  
Multiple R-squared: 0.03304, Adjusted R-squared: 0.03304  
F-statistic: 1.243e+04 on 2 and 727646 DF, p-value: < 2.2e-16

```

predictions <- predict(model, regression_data)

# Actual values

```

```
actuals <- regression_data$MedianPrice

# Calculate MAE
mae <- mean(abs(predictions - actuals))

# Calculate MSE
mse <- mean((predictions - actuals)^2)

# Calculate RMSE
rmse <- sqrt(mse)

# Print the metrics
cat("MAE:", mae, "\n")
```

MAE: 255133.1

```
cat("MSE:", mse, "\n")
```

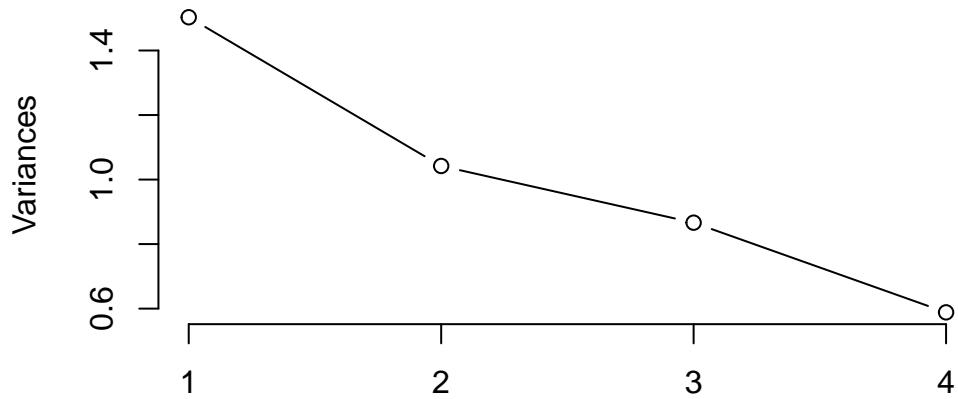
MSE: 141047372856

```
cat("RMSE:", rmse, "\n")
```

RMSE: 375562.7

```
plot(pca_results, type = "lines")
```

## pca\_results

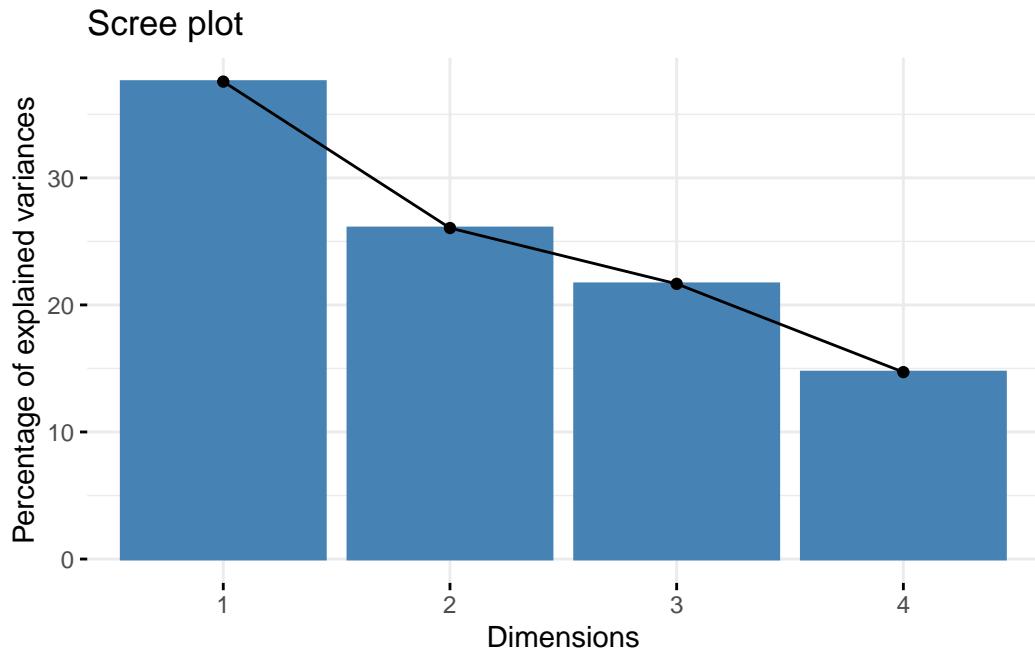


```
summary(pca_results)
```

Importance of components:

	PC1	PC2	PC3	PC4
Standard deviation	1.2260	1.0208	0.9308	0.7671
Proportion of Variance	0.3758	0.2605	0.2166	0.1471
Cumulative Proportion	0.3758	0.6363	0.8529	1.0000

```
fviz_eig(pca_results)
```



```
fviz_pca_var(pca_results,
              col.var = "contrib", # Color by contributions to the PC
              gradient.cols = c("#00AFBB", "#E7B800", "#FC4E07"),
              repel = TRUE      # Avoid text overlapping
)
```

