

Informations

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Developing Data Products - Shiny Application and Reproducible Pitch

Overview

For the developing Data Products course project I have created a Shiny Application which will predict diamond price on the basis of chosen parameters. Diamond dataset which I have collected from the website <http://www.pricescope.com/>. Diamond price determined by several factors, such as carat, Clarity, Cut etc. In my dataset I have chosen 6 predictors - Shape, Carat, Cut, Color, Clarity, Depth.

Data Preparation

Read the dataset Diamond_price.csv which is in the current directory.

```
data <- read.csv("Diamond_price.csv", header=TRUE)
str(data)
```

```
## 'data.frame':    1000 obs. of  10 variables:
## $ Shape          : Factor w/ 10 levels "Asscher","Cushion",...: 4 4 4 4 4 4 4 4 4 4 ...
## $ Carat          : num  3.13 1.03 1.02 1.63 1.2 1.5 1.71 2.04 2.04 1.67 ...
## $ Cut            : Factor w/ 3 levels "Good","Ideal",...: 1 1 1 1 2 2 2 2 2 2 ...
## $ Color          : Factor w/ 9 levels "D","E","F","G",...: 1 5 4 8 2 2 5 3 3 6 ...
## $ Clarity        : Factor w/ 9 levels "I1","I2","IF",...: 5 1 5 5 5 5 5 5 5 5 ...
## $ Table          : num  54 51 56 63 48.4 52 51.4 52 64.9 54.5 ...
## $ Depth          : num  56.9 57.5 51.3 43 57.9 53 61.4 50.2 39.3 41.6 ...
## $ Cert           : Factor w/ 3 levels "AGS","AGSL","GIA": 1 1 1 1 1 1 1 1 1 1 ...
## $ Measurements: Factor w/ 792 levels "", "0 x 0 x 0",...: 750 450 475 649 614 681 699 728 787 651 ...
## $ Price          : Factor w/ 868 levels "$1,010","$1,036",...: 428 462 459 565 665 795 827 176 151 756
```

```
data$Price <- gsub('\\$', '', data$Price)
data$Price <- gsub(',', '', data$Price)
mydata <- data[,c(1,2,3,4,5,7,10)]
mydata$Price <- as.numeric(as.character(mydata$Price))
mydata <- mydata[mydata$Price < 15000,] # remove outliers
head(mydata)
```

```
##   Shape Carat   Cut Color Clarity Depth Price
## 2 Heart  1.03  Good    H     I1  57.5  3188
```

```
## 3 Heart 1.02 Good G SI2 51.3 3158
## 4 Heart 1.63 Good K SI2 43.0 4009
## 5 Heart 1.20 Ideal E SI2 57.9 5256
## 6 Heart 1.50 Ideal E SI2 53.0 7860
## 7 Heart 1.71 Ideal H SI2 61.4 8557
```

Build Model

```
library(caret)
library(randomForest)
inTrain <- createDataPartition(mydata$Price, p=0.7,list = FALSE)
traindata <- mydata[inTrain,]
testdata <- mydata[-inTrain,]
model.forest <- train(Price~., traindata, method = "rf", trControl = trainControl(method = "cv", number = 10))
testdata$pred <- predict(model.forest, newdata = testdata)
ggplot(aes(x=actual, y=prediction),data=data.frame(actual=testdata$Price, prediction=predict(model.forest, testdata)))
  geom_point() +geom_abline(color="red") +ggtitle("RandomForest Regression in R" )
```

The shiny application I developed has been published in shiny server at <https://bdalmaso.shinyapps.io/diamondsapp/>.

To reproduce the shiny application on your local system, you need to install the relevent packages (caret and randomForest) and download diamond dataset, server.R and ui.R from github repository.

How to Run the Application

After downloading the above mentioned files you have to keep all the files in a folder and run **runApp()** function. Instantly application will be open locally in default browser. In the html page you will see at left side there are several input parameters you have to select by drop down or by increasing/decreasing the values. After selection you have to press the Submit button, the diamond price will be shown at right side. The predictors are :

1. Shape - Diamond shapes are Heart ,Round, Princess, Cushion,Pear,Marquise, Emerald, Radiant, Oval, Asscher;
2. Carat - The weight or size of the diamond (in this project diamond weight can be from .32 carat to 4.0 carat);
3. Cut - The proportions and relative angles of the facets. 3 type of cuts : Good ,Ideal, Very Good;
4. Color - Color has several values, such as D, E, F, G, H, I, J, K, L;
5. Clarity - The absence of internal imperfections. Clarity has following values: 'I1', 'I2', 'IF', 'SI1', 'SI2', 'VS1', 'VS2', 'VVS1', 'VVS2';
6. Depth - Diamond depth can be very from 40 to 80.

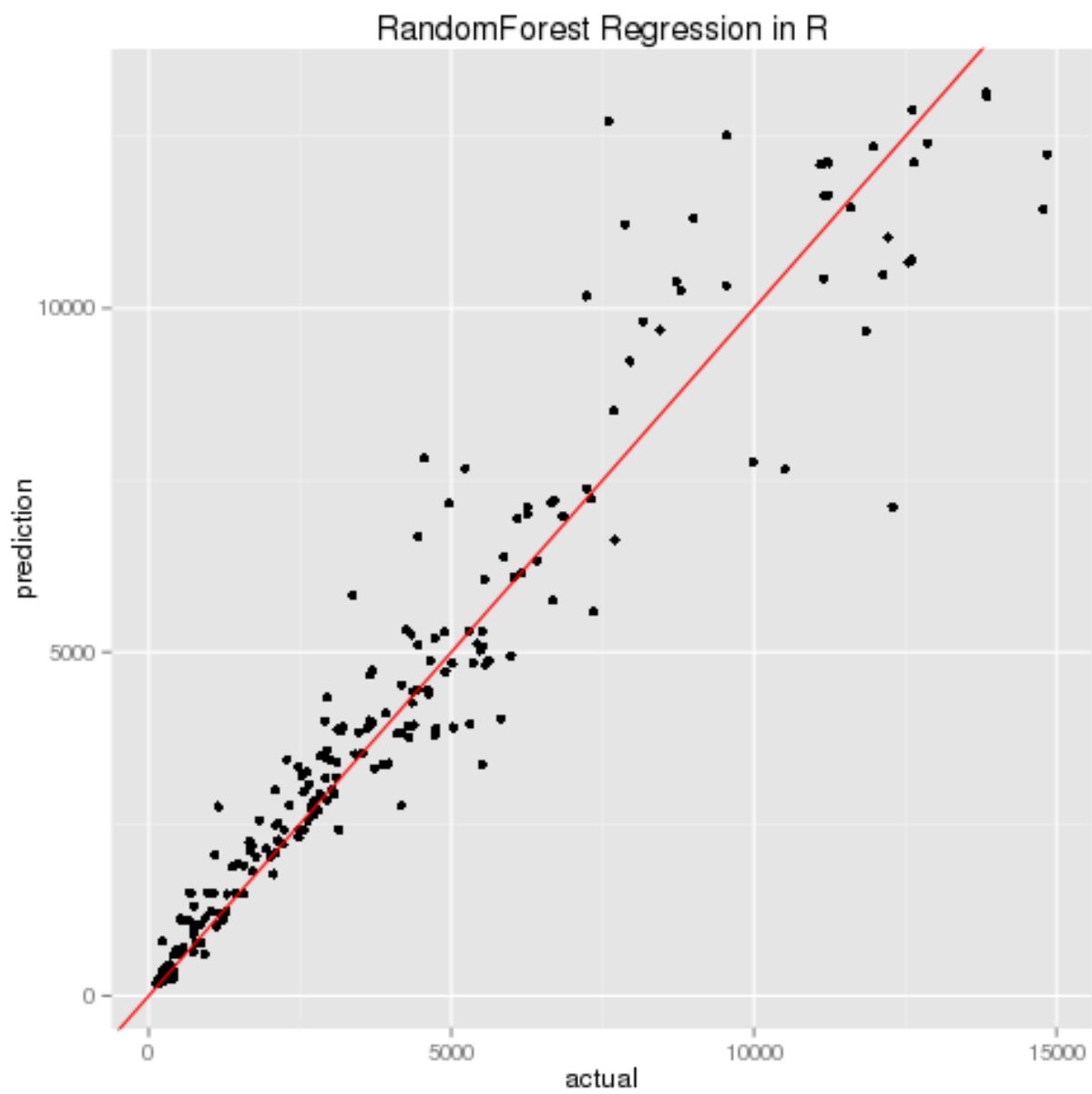


Figure 1: plot of chunk unnamed-chunk-2