```
In [8]:
# PREDICTING PRICE OF PRE-OWNED CARS
import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean squared error
import statsmodels.api as sm
# Importing model_selection library for using cross_validation
from sklearn import model_selection
# Importing the library for PCA
from sklearn.decomposition import PCA
In [9]:
# Setting dimensions for plot
# -----
sns.set(rc={'figure.figsize':(11.7,8.27)})
In [80]:
# Reading CSV file
data = pd.read_csv('Toyota.csv',index_col=0)
```

In [81]:

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 1436 entries, 0 to 1435
Data columns (total 10 columns):
Price
             1436 non-null int64
             1436 non-null float64
Age
             1436 non-null float64
KM
FuelType
             1436 non-null object
             1436 non-null float64
MetColor
             1436 non-null float64
Automatic
             1436 non-null int64
             1436 non-null int64
CC
Doors
             1436 non-null int64
Weight
             1436 non-null int64
dtypes: float64(4), int64(5), object(1)
memory usage: 123.4+ KB
None
```

In [83]:

```
####### CHECK FOR THE MISSING VALUES
data.isnull().sum()
Out[83]:
Price
             0
Age
             0
             0
ΚM
FuelType
ΗP
MetColor
Automatic
             0
CC
Doors
             0
Weight
dtype: int64
In [84]:
data=pd.get_dummies(data,drop_first=True)
In [85]:
# Storing the column names in variables
features = list(set(data.columns)-set(['Price']))
target = list(['Price'])
print(features)
print(target)
['FuelType_Diesel', 'HP', 'Automatic', 'CC', 'Age', 'FuelType_Petrol', 'K
M', 'Weight', 'MetColor', 'Doors']
['Price']
In [123]:
x = data.loc[:, features]
y = data.loc[:,target]
# Sklearn - package to split data into train & test
from sklearn.model_selection import train_test_split
```

test_size=0.3,
random_state=40)

Splitting test & train as 30% and 70%

train_x, test_x, train_y, test_y = train_test_split(x,y,

In [103]:

In [104]:

```
# finding the mean for test data value
base_pred = np.mean(test_y)
print(base_pred)

# Repeating same value till length of test data
base_pred = np.repeat(base_pred, len(test_y))
```

Price 10521.624 dtype: float64

In [105]:

```
# finding the RMSE
from sklearn.metrics import mean_squared_error
base_RMSE =(mean_squared_error(test_y,base_pred))**0.5
print(base_RMSE)
```

3518.8221654138106

In [106]:

```
# RMSE of the linear model
lr_rmse = (mean_squared_error(test_y,predictions))**0.5
print("RMSE corresponding to Linear Regression model between X and Y: ",lr_rmse)
```

1345.6596407279985

Below is the script for Principal Component Regression

- 1. Response variable is Price (train y, test y)
- 2. Independent variables are 'FuelType_Diesel', 'HP', 'Automatic', 'CC', 'Age', 'FuelType_Petrol', 'KM', 'Weight', 'MetColor', 'Doors' (train_x and test_x)
- 3. Choose number of PCs starting from 1 till number of features in the dataset (which is 10 in this case). i. Run PCA among independent variables of train_x and get the PCs in pc_train, pc_test for each fold in the cross validation. ii. pc_train, pc_test are linear combinations of train_x and test_x iii. Regress train_y and test_y on pc_train and pc_test respectively. iv. Predict value of test_y based on value of pc_test
- 4. Draw a plot between the number of PCs Vs. RMSE
- 5. Choose the number of PCs corresponding to lowest RMSE

In [124]:

```
Function to linearly regress the training samples of X against Y and to return the list of rmse for each variable in the output (y) of the testing samples Arguments: X_train (numpy.ndarray),y_train (numpy.ndarray),X_test (numpy.ndarray),y_test (numpy.ndarray)

Returns: rmse (list of arrays)

...

def linreg(X_train,X_test,y_train,y_test):
    ols = LinearRegression()
    ols.fit(X_train, y_train)
    y_pred = ols.predict(X_test)
    rmse=np.sqrt(((y_test-y_pred)**2).mean()) #square root (mean square error)
    return rmse
```

In [125]:

```
. . .
Function to perform linear regression using leave one out cross validation between Y
and reduced set of X iteratively from 1 principal component to maximum number of princi
pal coomponents specified
Arguments: X (numpy.ndarray), y (numpy.ndarray), nfact (int)
Returns: RMSE_pcr_arr (numpy.ndarray of size nfact,number of variables of y)
def pca kfold ols(X,y,nfact):
RMSE_pcr_lst=list()
for pc in range(1,nfact,1): # Iterating over number of principal components
     pca = PCA(n_components=pc) # Instantiating pca instance for each number of princip
al components in the iteration process
     X_red = pca.fit_transform(X)
     rmse pcr lst=list()
     k = model_selection.KFold(5) # Instantiating a kfold cv instance
     for tr_idx, tst_idx in k.split(X_red): # interating through multiple folds
         pc_train, pc_test = X_red[tr_idx], X_red[tst_idx] # input X for both train and
test
         y_train, y_test = y[tr_idx], y[tst_idx] # output Y for both train and test
         rmse_pcr_lst.append(linreg(pc_train,pc_test,y_train,y_test)) # Appending rmse
for each fold into a list
     RMSE_pcr_lst.append(np.array(rmse_pcr_lst).mean()) # Averaging RMSE of all the fol
ds per
 return np.array(RMSE pcr lst) #
```

In [126]:

```
train_x.head()
```

Out[126]:

| | FuelType_Diesel | HP | Automatic | CC | Age | FuelType_Petrol | KM | Weight |
|------|-----------------|---------|-----------|------|--------|-----------------|-----------|--------|
| 978 | 0 | 110.000 | 0 | 1600 | 65.000 | 1 | 45681.000 | 1050 |
| 1206 | 0 | 110.000 | 0 | 1600 | 73.000 | 1 | 87358.000 | 1050 |
| 1168 | 0 | 86.000 | 0 | 1300 | 78.000 | 1 | 96000.000 | 1015 |
| 1226 | 0 | 110.000 | 0 | 1600 | 55.672 | 1 | 84000.000 | 1075 |
| 1010 | 0 | 110.000 | 0 | 1600 | 60.000 | 1 | 36943.000 | 1070 |
| 4 | | | | | | | | • |

In [127]:

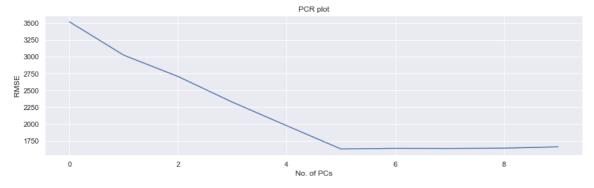
```
pcs=10
train_x=train_x.to_numpy()
train_y=train_y.to_numpy()
RMSE_pcr=np.append(base_RMSE,pca_kfold_ols(train_x,train_y,pcs))
RMSE_pcr.shape
```

Out[127]:

(10,)

In [128]:

```
fig = plt.figure(figsize=(15,4))
plt.plot(range(pcs),RMSE_pcr)
plt.xlabel('No. of PCs')
plt.ylabel('RMSE')
plt.title('PCR plot')
plt.grid(True)
```



First five PCs capture maximum variance in the given data.

In [129]:

```
print("The RMSE corresponding to linreg model between Z(5 PCs) and Y is: ", RMSE_pcr[5
])
```

The RMSE corresponding to linreg model between Z(5 PCs) and Y is: 1628.83 5612295244

In [130]:

END OF SCRIPT