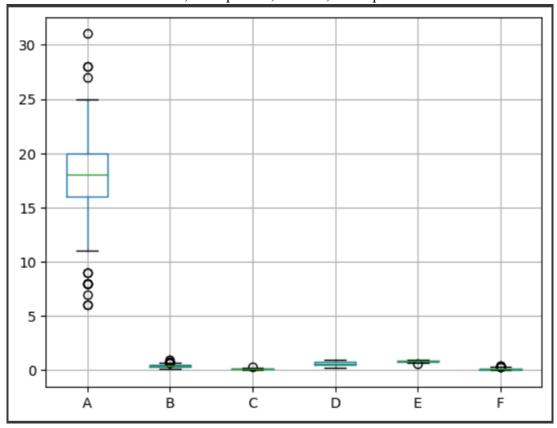
## In-Lab

### Task 1

Writing Code to visualize data using box plot:

```
from keras.models import Sequential
from keras.layers import Dense, Dropout
from sklearn.metrics import classification report, confusion matrix
from sklearn.model selection import train test split
from sklearn.metrics import mean_squared_error
import numpy as np
from sklearn import linear model
from sklearn import preprocessing
from sklearn import tree
from sklearn.ensemble import RandomForestRegressor,
GradientBoostingRegressor
import pandas as pd
import csv
import matplotlib.pyplot as plt
from google.colab import files
u = files.upload()
np.random.seed(7)
df = pd.read csv("Alumni Giving Regression
dd df 1 = df.head()
import seaborn as sns
boxplot = pd.DataFrame(df).boxplot()
```

It print the boxplot of dataset which is basically distribution of the data based on five number summaries. Minimum, first quartile, median, third quartile and maximum.

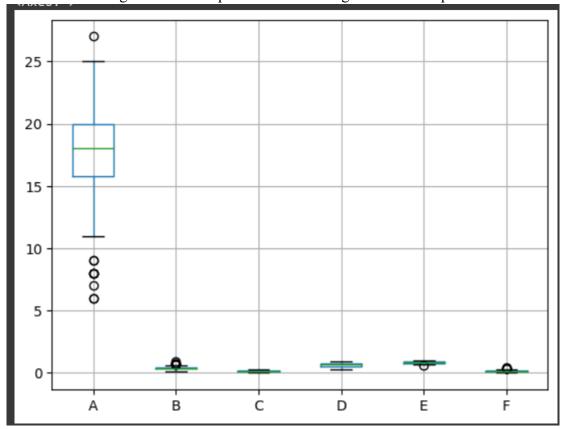


# Task 2

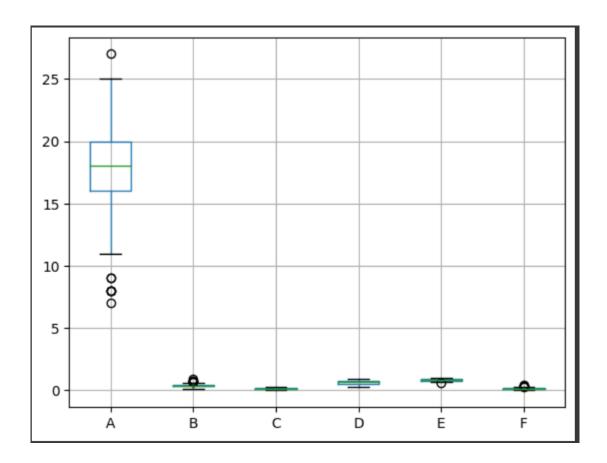
```
quantile99 = df.iloc[:,0].quantile(0.99)
df1 = df[df.iloc[:,0] < quantile99]
df1.boxplot()

quantile1 = df.iloc[:,0].quantile(0.01)
quantile99 = df.iloc[:,0].quantile(0.99)
df2 = df[(df.iloc[:,0] > quantile1) & (df.iloc[:,0] < quantile99)]
df2.boxplot()</pre>
```

Code 1 finds the 99<sup>th</sup> percentile of the first column, creates a subset df1 and store the values containing less than this percentile in it and generates a boxplot for it.



Code 2 finds the 1<sup>st</sup> and 99<sup>th</sup> percentile of the first column and select the values which are greater than the 1<sup>st</sup> percentile and less than the 99<sup>th</sup> percentile and store them in df2 subset and generates a boxplot for it.



### Task 3

```
df.dropna()
Y POSITION = 5
model 2 features = [i for i in range(0,Y POSITION)]
X = df.iloc[:,model 2 features]
Y = df.iloc[:,Y_POSITION]
X train, X test, y train, y test =
train test split(X,Y,test size=0.20,random state=2020)
model3 = RandomForestRegressor()
model3.fit(X train,y train)
RF = model3
importances = RF.feature importances
std = np.std([tree.feature_importances_ for tree in RF.estimators_],
axis=0)
indices = np.argsort(importances)[::-1]
print("Feature ranking: ")
for f in range(X.shape[1]):
 print("%d. feature (Column index) %s (%f)" % (f + 1, indices[f],
importances[indices[f]]))
```

```
Feature ranking:

1. feature (Column index) 3 (0.381108)

2. feature (Column index) 0 (0.212209)

3. feature (Column index) 1 (0.176601)

4. feature (Column index) 4 (0.137474)

5. feature (Column index) 2 (0.092607)
```

Figure 1

#### Task 4

```
indices top3 = indices[:3]
print(indices top3)
dataset = df
df = pd.DataFrame(df)
Y POSITION = 5
TOP N FEATURE = 3
X = df.iloc[:,indices top3]
Y = df.iloc[:,Y_POSITION]
X train, X test, y train, y test =
train test split(X,Y,test size=0.20,random state=2020)
model1 = linear model.LinearRegression()
model1.fit(X train,y train)
y pred train1 = model1.predict(X train)
print("Regression")
print("==========")
RMSE_train1 = mean_squared_error(y_train,y_pred_train1)
print("Regression Train set: RMSE {}".format(RMSE train1))
print("======="")
y pred1 = model1.predict(X test)
RMSE_test1 = mean_squared_error(y_test,y_pred1)
print("Regression Test set: RMSE {}".format(RMSE test1))
```

Figure 2

There is no major difference between test scores of both previous lab without selecting top 3 features and this lab by selecting top 3 features.