## BUISNESS REPORT – Linear Regression:

The data is taken from the comp-activ databases which is a collection of computer systems activity measures. The data was collected from a Sun Sparcstation 20/712 with 128 Mbytes of memory running in a multi-user university department. The data was collected continuously every 5 seconds. Using the various system programs which are running in the background for every task being performed by the user, Predict the percentage portion of time (out of 100), that cpu runs in user mode, and how does each system program affect the same.

## Tasks to be performed:

- 1. Load data and describe data
- a. Import necessary libraries & packages
- b. Load dataset
- c. Check necessary details about data like shape, data types of the variable, missing values etc.

```
import pandas as pd
import numpy as np
import seaborn as sns
import seaborn as sb
import matplotlib.pyplot as plt
import statsmodels.api as sm
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_squared_error
from statsmodels.stats.outliers_influence import variance_inflation_factor
import math
from sklearn.preprocessing import MinMaxScaler
import warnings
warnings.filterwarnings("ignore")
```

```
Loading the dataset

df = pd.read_csv('compactiv.csv')

df.head()
```

From the table we can see all the required variables that we require excel sheet

	0	1	2	3	4
Iread	1	0	15	0	5
lwrite	0	0	3	0	1
scall	2147	170	2162	160	330
sread	79	18	159	12	39
swrite	68	21	119	16	38
fork	0.2	0.2	2.0	0.2	0.4
exec	0.2	0.2	2.4	0.2	0.4
rchar	40671.0	448.0	NaN	NaN	NaN
wchar	53995.0	8385.0	31950.0	8670.0	12185.0
pgout	0.0	0.0	0.0	0.0	0.0
ppgout	0.0	0.0	0.0	0.0	0.0
pgfree	0.0	0.0	0.0	0.0	0.0
pgscan	0.0	0.0	0.0	0.0	0.0
atch	0.0	0.0	1.2	0.0	0.0
pgin	1.6	0.0	6.0	0.2	1.0
ppgin	2.6	0.0	9.4	0.2	1.2
pflt	16.0	15.63	150.2	15.6	37.8
vflt	26.4	16.83	220.2	16.8	47.6
runqsz	CPU_Bound	Not_CPU_Bound	Not_CPU_Bound	Not_CPU_Bound	Not_CPU_Bound
freemem	4670	7278	702	7248	633
freeswap	1730946	1869002	1021237	1863704	1760253
usr	95	97	87	98	90

Check the necessary detail about data is given by its info, shape, data types:

```
df.shape
(8192, 22)
df.info()
<class 'pandas.core.frame.DataFrame'>
Int64Index: 8192 entries, 0 to 8191
Data columns (total 23 columns):
                              Non-Null Count Dtype
#
    Column
    lread
                              8192 non-null
                                               float64
0
 1
     lwrite
                              8192 non-null
                                               float64
                              8192 non-null
                                               float64
 2
     scall
     sread
                              8192 non-null
                                               float64
     swrite
                              8192 non-null
                                               float64
 5
     fork
                              8192 non-null
                                               float64
                                               float64
 6
     exec
                              8192 non-null
                              8192 non-null
                                               float64
 8
                              8192 non-null
                                               float64
     wchar
 9
     pgout
                              8192 non-null
                                               float64
 10
     ppgout
                              8192 non-null
                                               float64
     pgfree
                                               float64
 11
                              8192 non-null
                              8192 non-null
                                               float64
 12
     pgscan
 13
     atch
                              8192 non-null
                                               float64
                              8192 non-null
                                               float64
 14
     pgin
                                               float64
 15
     ppgin
                              8192 non-null
 16
     pflt
                              8192 non-null
                                               float64
 17
     vflt
                              8192 non-null
                                               float64
 18
     freemem
                              8192 non-null
                                               float64
                                               float64
 19
     freeswap
                              8192 non-null
 20 usr
                              8192 non-null
                                               float64
 21 CPU_Bound_CPU_Bound
                              8192 non-null
                                               uint8
 22 CPU_Bound_Not_CPU_Bound
                              8192 non-null
                                               uint8
dtypes: float64(21), uint8(2)
memory usage: 1.4 MB
```

# df.dtypes

lread	int64
lwrite	int64
scall	int64
sread	int64
swrite	int64
fork	float64
exec	float64
rchar	float64
wchar	float64
pgout	float64
ppgout	float64
pgfree	float64
pgscan	float64
atch	float64
pgin	float64
ppgin	float64
pflt	float64
vflt	float64
runqsz	object
freemem	int64
freeswap	int64
usr	int64
dtype: objec	ct

lread	0
lwrite	0
scall	0
sread	0
swrite	0
fork	0
exec	0
rchar	104
wchar	15
pgout	0
ppgout	0
pgfree	0
pgscan	0
atch	0
pgin	0
ppgin	0
pflt	0
vflt	0
runqsz	0
freemem	0
freeswap	0
usr	0

From the above code and result we can figure out if there any null values, missing values, datatype of all the variables. Rchar has 104 missing values and wchar has 15 missing values.

## 2. Perform EDA and data cleaning

a. Generate the summary statistics for each of the variables and write comments on your observations

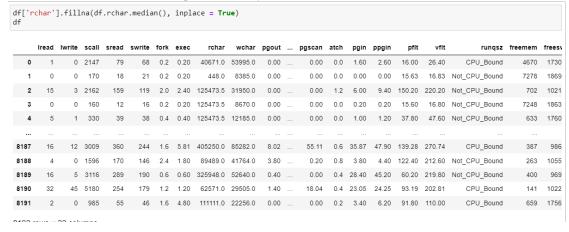
df.describe().T								
	count	mean	std	min	25%	50%	75%	max
Iread	8192.0	1.955969e+01	53.353799	0.0	2.0	7.0	20.000	1845.0
lwrite	8192.0	1.310620e+01	29.891726	0.0	0.0	1.0	10.000	575.0
scall	8192.0	2.306318e+03	1633.617322	109.0	1012.0	2051.5	3317.250	12493.0
sread	8192.0	2.104800e+02	198.980146	6.0	86.0	166.0	279.000	5318.0
swrite	8192.0	1.500582e+02	160.478980	7.0	63.0	117.0	185.000	5456.0
fork	8192.0	1.884554e+00	2.479493	0.0	0.4	0.8	2.200	20.1
exec	8192.0	2.791998e+00	5.212456	0.0	0.2	1.2	2.800	59.5
rchar	8088.0	1.973857e+05	239837.493526	278.0	34091.5	125473.5	267828.750	2526649.0
wchar	8177.0	9.590299e+04	140841.707911	1498.0	22916.0	46619.0	106101.000	1801623.0
pgout	8192.0	2.285317e+00	5.307038	0.0	0.0	0.0	2.400	81.4
ppgout	8192.0	5.977229e+00	15.214590	0.0	0.0	0.0	4.200	184.2
pgfree	8192.0	1.191971e+01	32.363520	0.0	0.0	0.0	5.000	523.0
pgscan	8192.0	2.152685e+01	71.141340	0.0	0.0	0.0	0.000	1237.0
atch	8192.0	1.127505e+00	5.708347	0.0	0.0	0.0	0.600	211.5
pgin	8192.0	8.277960e+00	13.874978	0.0	0.6	2.8	9.765	141.2
ppgin	8192.0	1.238859e+01	22.281318	0.0	0.6	3.8	13.800	292.6
pflt	8192.0	1.097938e+02	114.419221	0.0	25.0	63.8	159.600	899.8
vflt	8192.0	1.853158e+02	191.000603	0.2	45.4	120.4	251.800	1365.0
freemem	8192.0	1.763456e+03	2482.104511	55.0	231.0	579.0	2002.250	12027.0
reeswap	8192.0	1.328126e+06	422019.426957	2.0	1042623.5	1289289.5	1730379.500	2243187.0
usr	8192.0	8.396887e+01	18.401905	0.0	81.0	89.0	94.000	99.0

We can see from the above summary statistics the mean of - Number of characters transferred per second by system write calls is the maximum. Most of the variables have minimum values from 0. The maximum Number of characters transferred per second by system read calls is 2526649 which is the highest frequency reached among all other variables.

## b. Working with Null/Missing values

i. Check for Missing values and perform the necessary steps for data imputation and provide reasoning for your approach.

As we saw in the previous question there are missing values that must be replaced. Since both are numerical in nature we must replace them by the median.





ii. Check for the Zero values and understand the importance of that data point. Please provide your comments on if we need to change them (impute) or drop them.

df.isin([0	]).Sum(		
lread	675		
lwrite	2684		
scall	0		
sread	0		
swrite	0		
fork	21		
exec	21		
rchar	0		
wchar	0		
pgout	4878		
ppgout	4878		
pgfree	4869		
pgscan	6448		
atch	4575		
pgin	1220		
ppgin	1220		
pflt	3		
vflt	0		
runqsz	0		
freemem	0		
freeswap	0		
usr	283		

We can see from the above result we can see how many of the variables have the 0 values.

- c. Working with Outliers
- i. Check for outliers and provide comments

Check Outliers in the HTML file.

On performing our code we can see that almost every variable has an outlier(refer from HTML file.)

'lread', 'sread', 'swrite', rchar, wchar, exec, atch 'pgout', 'ppgout', 'pgfree', 'pgin', 'ppgin', 'pflt', 'vflt', freemem, freeswap are all right skwed and have the mode is often less than the median, which is less than the mean.

Pg scan has no distribution hence no skewness determined.

Freeswap and usr have a left skewed.

```
ii. Perform outlier treatment (only if required)
def remove_outlier(column):
sorted(column)
  q1=df[column].quantile(0.25)
  q3=df[column].quantile(0.75)
  iqr=q3-q1
  lower=q1-1.5*iqr
  upper=q3+1.5*iqr
  return lower,upper
or i in nums:
  lower,upper=remove_outlier(i)
  df[i]=np.where(df[i]>upper,upper,df[i])
  df[i]=np.where(df[i]<lower,lower,df[i])
for i in nums:
  sns.boxplot(df[i],showmeans=True)
  plt.show()
```

We create a function to remove the outlier and use quartiles to treat the inconsistency if any. From the above code we can treat the outlier.(Can see the corrected outlier treatment in the HTML File)

## d. Scaling the data

i. Only if scaling is necessary, please perform the same and provide your reasoning

```
scaler = MinMaxScaler()
columns=['Iread','Iwrite','scall','sread','swrite','fork','exec','rchar','wchar','pgout','ppgout','pgscan','pg
free','atch','pgin','ppgin','pflt','vflt','freemem','freeswap','usr']
```

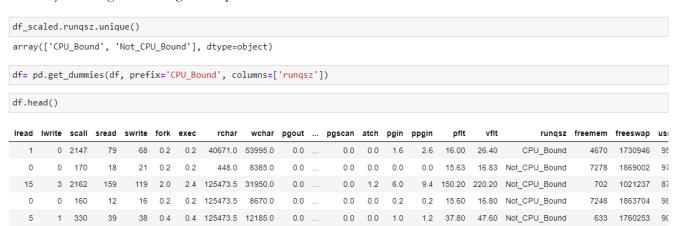
```
df_scaled = scaler.fit_transform(df[columns].to_numpy())
df_scaled = pd.DataFrame(df_scaled,columns=columns)
print("Scaled Dataset Using MinMaxScaler")
df_scaled['runqsz']=df['runqsz']
df_scaled.head().T
```

runqsz	CPU_Bound	Not_CPU_Bound	Not_CPU_Bound	Not_CPU_Bound	Not_CPU_Bound
usr	0.893333	0.946667	0.68	0.973333	0.76
freeswap	0.770522	0.832369	0.45258	0.829996	0.783651
freemem	1.0	1.0	0.140526	1.0	0.12554
vfit	0.046686	0.029633	0.392017	0.029579	0.084462
pfit	0.04426	0.043237	0.415491	0.043154	0.104564
ppgin	0.077381	0.0	0.279762	0.005952	0.035714
pgin	0.068049	0.0	0.255183	0.008506	0.042531
atch	0.0	0.0	0.8	0.0	0.0
pgfree	0.0	0.0	0.0	0.0	0.0
pgscan	0.0	0.0	0.0	0.0	0.0
ppgout	0.0	0.0	0.0	0.0	0.0
pgout	0.0	0.0	0.0	0.0	0.0
wchar	0.229117	0.030057	0.132904	0.031301	0.046642
rchar	0.066119	0.000278	0.20493	0.20493	0.20493
exec	0.029851	0.029851	0.358209	0.029851	0.059701
fork	0.040816	0.040816	0.408163	0.040816	0.081633
swrite	0.168975	0.038781	0.310249	0.024931	0.085873
sread	0.129778	0.021333	0.272	0.010667	0.058667
scall	0.305725	0.009151	0.307975	0.007651	0.033153
lwrite	0.0	0.0	0.12	0.0	0.04
Iread	0.021277	0.0	0.319149	0.0	0.106383
	0	.l	2	ż	4

In this method, we convert variables with different scales of measurements into a single scale. First we check what are the values that needs to be changed. Standard Scaler normalizes the data using the formula (x-mean)/standard deviation.

#### e. Working with Categorical variables

i. Identify the categorical data given as part of the data set?



On performing the given code we can see that , runqsz has the categorical values that needs to be changed. Hence perform get dummies function to replace these values with one and 0.(Given in the HTML file)

ii.) Perform encoding and provide detailed comments and reasoning for the encoding approach.

Most of the machine learning models are designed to work on numeric data. Hence, we need to convert categorical text data into numerical data for model building

One-Hot-Encoding is used to create dummy variables to replace the categories in a categorical variable into features of each category and represent it using 1 or 0 based on the presence or absence of the categorical value in the record

- 3. Perform univariate, bivariate & Multivariate analysis
- a. Perform Univariate analysis for each variable and write comments

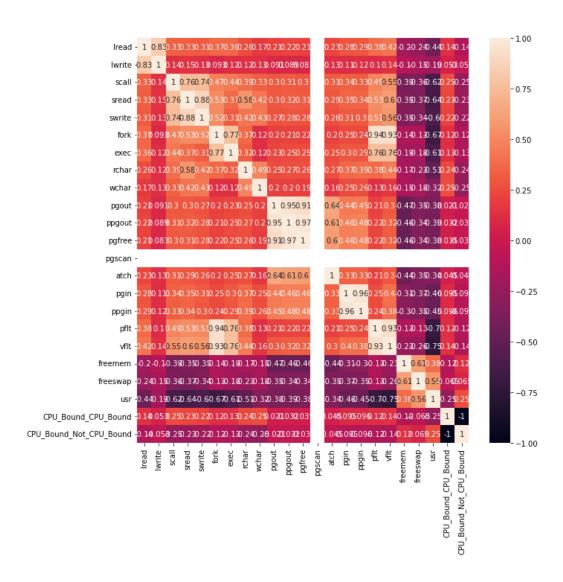
We have performed univariate analysis for each variable(in the HTML file.) For Iread its is positively skewed and greater frequency. For lwrite its similar to Iread but its less frequent. For all of the graphs itself we can see an unique peak in the beginning and then a drop which is constant and then a sudden rise as well. For pg scan we can see there is no data to be distributed.

b. Perform bivariate analysis and make necessary inference about the relation between the variables. Refer html file for the plot.

```
sns.pairplot(df_scaled, diag_kind='kde',size = 3)
plt.show()
```

- c. Perform Multivariate analysis and make necessary inferences about the relation between variables.
- d. Check Covariance and Correlation and identify positively and negatively correlated variables.

```
plt.figure(figsize=(20,20))
sns.heatmap(df.corr(), annot=True)
```



-1.50
-1.25
-1.00
-0.75
-0.50

- 0.00

-175

e. Identify the variables which has multicollinearity. Check for multi collinearity and drop the variables.

```
]: vif_data = pd.DataFrame()
   vif_data["feature"] = X.columns
   # calculating VIF for each feature
   vif_data["VIF"] = [variance_inflation_factor(X.values, i)
                             for i in range(len(X.columns))]
   cols_to_remove=list(vif_data.loc[vif_data['VIF']>=5]['feature'])
   print(cols_to_remove)
   vif_data.sort_values('VIF', ascending=False)
   print(vif_data)
   ['lread', 'sread', 'swrite', 'fork', 'pgout', 'ppgout', 'pgfree', 'pgin', 'ppgin', 'pflt', 'vflt', 'CPU_Bound_CPU_Bound', 'CPU_
   Bound_Not_CPU_Bound']
                       feature
   0
                                 5.232535
                         lread
                                 4.260737
   1
                        lwrite
   2
                         scall
                                 2.989088
   3
                         sread
                                 6.517014
   4
                        swrite
                                 5.639798
                          fork 12.947438
                                3.137525
                          exec
                                 2.087446
                         rchar
   8
                         wchar
                                 1.605667
   g
                         pgout 11.474544
   10
                        ppgout
                                30.685098
   11
                        pgfree 17.175122
                        pgscan
                                      NaN
   12
                                 1.859658
   13
                          atch
   14
                          pgin 13.733882
   15
                         ppgin 13.962094
                          pflt 11.536918
                          vflt 15.204685
                       freemem
                                 1.969949
   18
   19
                      freeswap
                                 1.825789
           CPU_Bound_CPU_Bound 13.213383
   20
   21 CPU_Bound_Not_CPU_Bound 13.225624
 new_df=X.drop(['lread', 'sread', 'swrite', 'fork', 'pgout', 'ppgout', 'pgfree', 'pgin', 'ppgin', 'pflt', 'vflt', 'CPU_Bound_CPU_f
        Iwrite
              scall exec
                           rchar
                                 wchar pgscan atch freemem freeswap
         0.0 2147.0 0.20 40671.0 53995.0 0.0 0.0 4659.125 1730946.0
         0.0
              170.0 0.20
                           448.0 8385.0
                                          0.0 0.0 4659.125 1869002.0
         3.0 2162.0 2.40 125473.5 31950.0 0.0 1.2 702.000 1021237.0
         0.0
             160.0 0.20 125473.5 8670.0 0.0 0.0 4659.125 1863704.0
         1.0 330.0 0.40 125473.5 12185.0 0.0 0.0 633.000 1760253.0
        12.0 3009.0 5.81 405250.0 85282.0 0.0 0.6 387.000 986647.0
   8188
         0.0 1596.0 1.80 89489.0 41764.0 0.0 0.8 263.000 1055742.0
         5.0 3116.0 0.60 325948.0 52640.0 0.0 0.4 400.000 969106.0
       25.0 5180.0 1.20 62571.0 29505.0 0.0 0.4 141.000 1022458.0
   8191 0.0 985.0 4.80 111111.0 22256.0 0.0 0.2 659.000 1756514.0
  8192 rows × 9 columns
```

First we check for multicollinearity and remove all the features that are multicollinear using variance inflation factor. Mathematically, the VIF for a regression model variable is equal to the ratio of the overall model variance to the variance of a model that includes only that single independent variable.

- 4. Building a Linear Regression Model
- a. Prepare data for model building

Preparing the data for model building, we drop the values of usr and copy it in another dataset so that we can use this data frame to train and test our model.

```
X = df.drop('usr', axis=1)
y = df[['usr']]
```

We get the table that we need to test and train.(refer to HTML file.)

b. Build linear regression model.

```
: X = new_df
  y = df[['usr']]
: from sklearn.model_selection import train_test_split
   X_{train}, X_{test}, y_{train}, y_{test} = train_{test_split}(X, y, test_{size}=0.25, random_{state}=1)
|: model.predict(X_train)
: 5745
            68.397776
   1930
           53.316204
   5622
            60.481046
          102.002734
           98.826406
   2255
   7935
            74.811305
   5192
            96.452824
   3980
            69.561082
   235
            70.027195
   5157
            94.588283
   Length: 6144, dtype: float64
|: regression_model = LinearRegression()
   regression_model.fit(X_train, y_train)
```

Following all the required codes to achieve the model to further find the required measures to check accuracy of our model.

c. Find the features that add value to the model. Identify the list of Variables which highly impact the prediction based on the correlation Metrix given for regression (target variable)

```
: vif = [variance_inflation_factor(X.values, ix) for ix in range(X.shape[1])]
  i=0
  for column in X.columns:
      if i < 15:
          print (column ,"--->", vif[i])
  lread ---> 5.232534882470838
  lwrite ---> 4.260737491684919
  scall ---> 2.9890877339622937
  sread ---> 6.517014292159949
  swrite ---> 5.639797517443181
  fork ---> 12.947438478657318
  exec ---> 3.1375249704246078
  rchar ---> 2.087445737239059
  wchar ---> 1.6056667145387191
  pgout ---> 11.474543882811679
  ppgout ---> 30.685097541787272
  pgfree ---> 17.17512180642015
  pgscan ---> nan
  atch ---> 1.8596581052295806
  pgin ---> 13.733882306297474
```

Again we can find the best correlated values that suit for our model. First we check for multicollinearity and remove all the features that are multicollinear using variance inflation factor. Mathematically, the VIF for a regression model variable is equal to the ratio of the overall model variance to the variance of a model that includes only that single independent variable.

d. Print test and train results with all variables and best fit line.

Linear Regression Model

```
sns.set(rc = {'figure.figsize':(7,7)})
ax=sns.regplot(x=y_test,y=y_pred,ci=None,color ='blue');
ax.set_xlabel('Actual')
ax.set_ylabel('Predicted')

Text(0, 0.5, 'Predicted')

100
90
80
60
50
40
30
```

Heavily clustered but in a consistent manner, but the growth seems to be very consistent.

100

#### 5. Model Performance:

a. Check for the performance measures for linear regression (Hint: RMSE, R square, etc.)

```
: # Let us check the intercept for the model
  intercept = regression\_model.intercept\_[\emptyset]
  print("The intercept for our model is {}".format(intercept))
  The intercept for our model is 52.703408616461346
: # R square on training data
  regression_model.score(X_train, y_train)
: 0.5513661898690928
: #R square on testing data
  regression_model.score(X_test, y_test)
: 0.5447312405803018
: #RMSE on training data
  \label{eq:predicted_train} \begin{split} & \texttt{predicted\_train-regression\_model.fit}(X\_\texttt{train}, \ y\_\texttt{train}). \\ & \texttt{predicted\_train}). \end{split} \\ & \texttt{predicted\_train}) \end{split}
: 12.16989876561295
: ##RMSE on testing data
  predicted\_test = regression\_model.fit(X\_train, y\_train).predict(X\_test)
  np.sqrt(mean_squared_error(y_test,predicted_test))
: 12.870554763335232
: # concatenate X and y into a single dataframe
data_train = pd.concat([X_train, y_train], axis=1)
  data_test=pd.concat([X_test,y_test],axis=1)
  data_train.head()
         lwrite scall exec rchar wchar pgscan atch freemem freeswap usr
   5745 10.0 2007.0 0.60 32665.0 49643.0 0.0 0.6 329.0 989029.0 94
   1930 4.0 837.0 0.80 6255.0 23670.0 0.0 0.0 3052.0 1013758.0 96
   \textbf{5622} \qquad 2.0 \quad 2227.0 \quad 6.70 \quad 108370.0 \quad 28568.0 \qquad 0.0 \quad 0.2 \qquad 314.0 \quad 1108418.0 \quad 77
   5340 0.0 2132.0 0.20 90813.0 28590.0 0.0 0.0 488.0 1742493.0 93
   2255 0.0 3517.0 2.99 310439.0 214462.0 0.0 0.2 420.0 1547657.0 86
```

From above relations we can see that my RMSE value is 12.16 for test and train almost. For R square the value is 0.544.

b. Experiment with data transformation and suggest if we can improve the model performance.

```
100 - (rmse/y_train.mean())*100

usr 85.523778

dtype: float64
```

To improve the performance of this model we can decrease the RMSE values and increase the R\*2 mean value.