

**1) Aim: Write a python program to find the best fit straight line and draw the scatter plot.**

**Source Code:**

```
#Import libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from statistics import mean

#Reading data
data=pd.read_csv('csdata.csv')

print(data)

x=data['x']
y=data['y']

def linear_regression(x,y):
    sumx=0
    for i in range(len(x)):
        sumx=sumx+x[i]
    sumy=0
    for i in range(len(y)):
        sumy=sumy+y[i]
    meanx=sumx/len(x)
    meany=sumy/len(y)
    print("mean of x :",meanx)
    print("mean of y :",meany)
    n=sum((x-meanx)*(y-meany))
    d=sum((x-meanx)**2)
    b1=n/d
    b0=meany-b1*meanx
    return b0,b1

b0,b1=linear_regression(x,y)
print("B0 :",round(b0,5)," , B1 :",round(b1,5))

print("Equation :")
print("y =",round(b0,5)," + x *",round(b1,5))

y_hat=b0+x*b1
print("y_hat\n",y_hat)

def plot1(x,y,y_hat):
```

```
plt.scatter(x,y)
plt.xlabel("x values")
plt.ylabel("y values")
plt.title("Data")
plt.show()
plt.plot(x,y,'ro-')
plt.plot(x,y_hat,'bo-')
plt.show()
```

#plot the graph

plot1(x,y,y\_hat)

```
def cost_function(y,y_hat):
    ybar=mean(y)
    sst=sum((y-ybar)**2)
    ssr=sum((y_hat-ybar)**2)
    r2=ssr/sst
    return r2
```

```
r2=cost_function(y,y_hat)
print("Cost function is",round(r2,5))
if r2<0.9:
    print("Not best fit")
else:
    print("Best fit")
```

### Output 1

```
x  y
0 1 2.4
1 2 3.0
2 3 3.6
3 4 4.0
4 6 5.0
5 8 6.0
```

```
mean of x : 4.0
mean of y : 4.0
B0 : 1.97647 , B1 : 0.50588
```

Equation :  
 $y = 1.97647 + x * 0.50588$

Regd. No :

y\_hat :

0 2.482353

1 2.988235

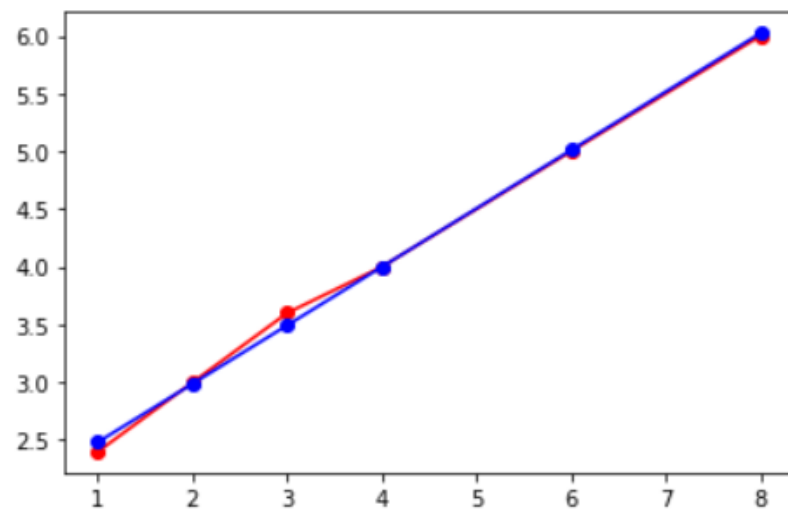
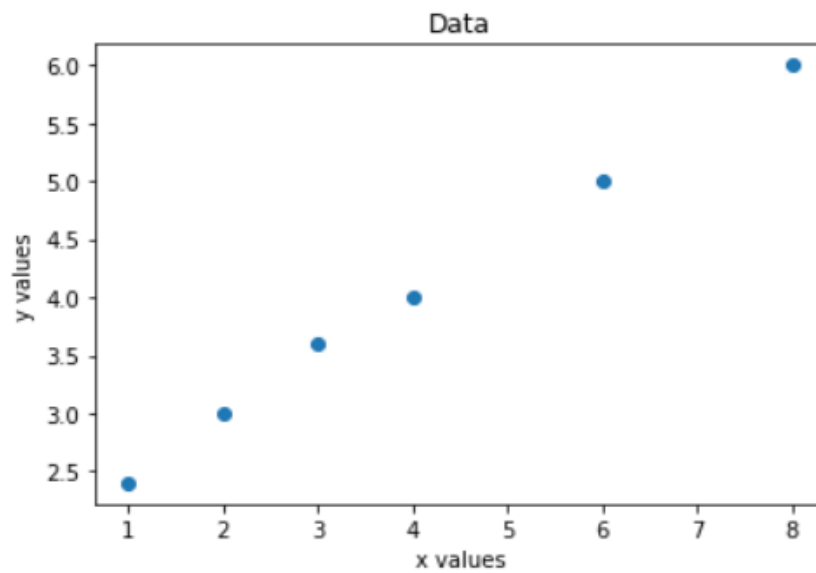
2 3.494118

3 4.000000

4 5.011765

5 6.023529

Name: x, dtype: float64



Cost function is 0.99784

Best fit

## Output 2

```
x y
0 1 7.2
1 2 5.0
2 3 1.0
3 4 8.0
4 5 5.5
5 6 6.0
6 7 2.5
7 8 8.0
```

mean of x : 4.5

mean of y : 5.4

B0 : 5.1 , B1 : 0.06667

Equation :

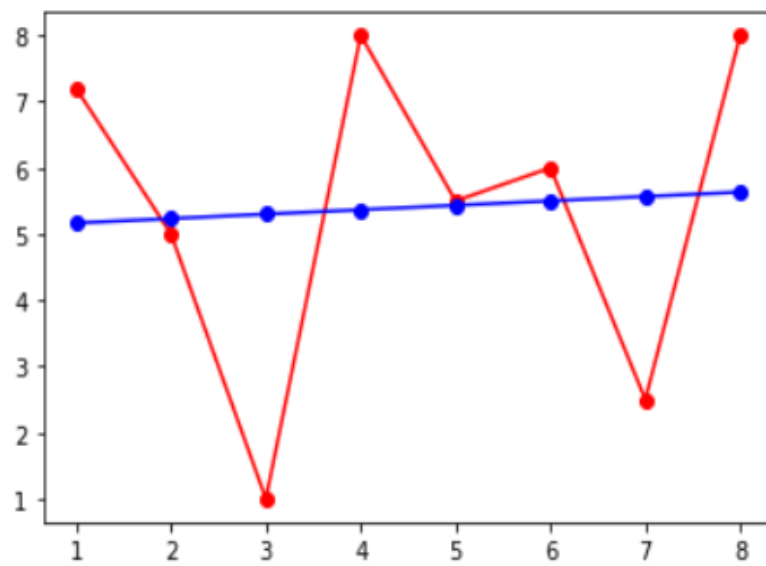
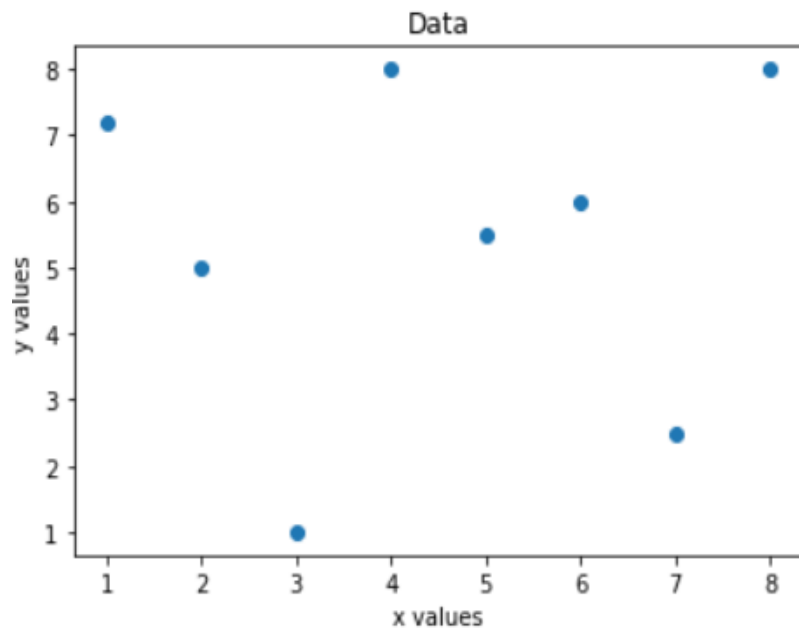
$y = 5.1 + x * 0.06667$

y\_hat :

```
0 5.166667
1 5.233333
2 5.300000
3 5.366667
4 5.433333
5 5.500000
6 5.566667
7 5.633333
```

Name: x, dtype: float64

Regd. No :



Cost function is 0.00414  
Not best fit

**2) Aim: Write a python program to fit a second degree parabola of the form  $y=a+bx+cx^2$  and draw the scatter plot.**

**Source Code:**

```
#importing libraries
import numpy as np
x=np.array([float(x) for x in input().split(" ")])
y=np.array([float(x) for x in input().split(" ")])
n=len(x)

sumx=np.sum(x)
sumy=np.sum(y)
sumxy=np.sum(x*y)
sumx2=np.sum(x*x)
sumx3=np.sum(x*x*x)
sumx4=np.sum(x*x*x*x)
sumx2y=np.sum(x*x*y)

#calculating determinant
def getMinor(m,i,j):
    return [row[:j] + row[j+1:] for row in (m[:i]+m[i+1:])]
def getDetermninant(m):
    if len(m) == 2:
        return m[0][0]*m[1][1]-m[0][1]*m[1][0]
    determinant = 0
    for c in range(len(m)):
        determinant += ((-1)**c)*m[0][c]*getDetermninant(getMinor(m,0,c))
    return determinant

#by using cramer's rule (without built-in)
p=getDetermninant([[n,sumx,sumx2],[sumx,sumx2,sumx3],[sumx2,sumx3,sumx4]])
q=getDetermninant([[sumy,sumxy,sumx2y],[sumx,sumx2,sumx3],[sumx2,sumx3,sumx4]])
r=getDetermninant([[n,sumx,sumx2],[sumy,sumxy,sumx2y],[sumx2,sumx3,sumx4]])
s=getDetermninant([[n,sumx,sumx2],[sumx,sumx2,sumx3],[sumy,sumxy,sumx2y]])
a=round(q/p,3)
b=round(r/p,3)
c=round(s/p,3)

print("The equation of parabola is  $y={}+{}x+{}x^2$ ".format(a,b,c))
import matplotlib.pyplot as plt
plt.scatter(x,y)
plt.xlabel("x values")
plt.ylabel("y values")
```

```
plt.title("Data")  
plt.show()
```

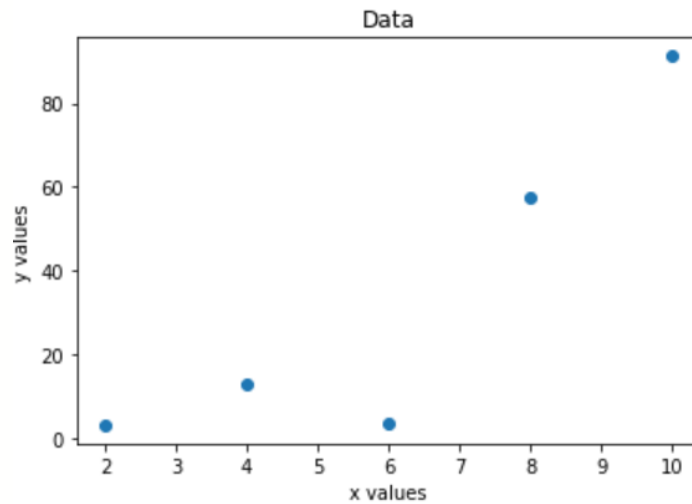
**Input 1:**

2 4 6 8 10

3.07 12.85 3.47 57.38 91.29

**Output 1:**

The equation of parabola is  $y=23.096+-12.855x+1.992x^2$



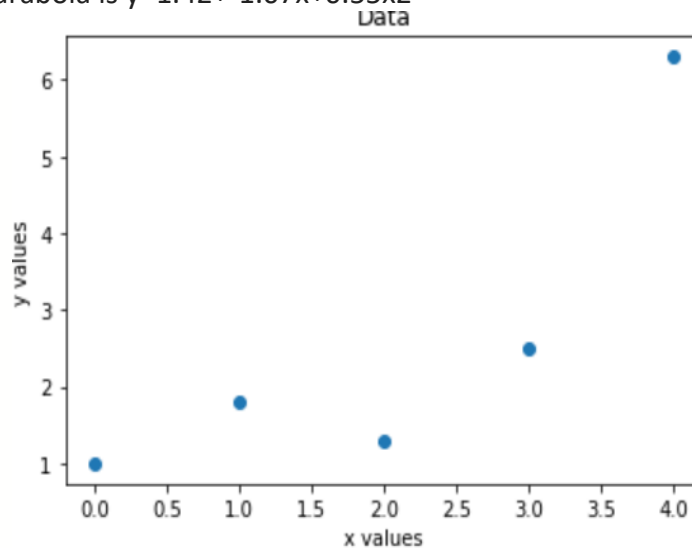
**Input 2:**

0 1 2 3 4

1 1.8 1.3 2.5 6.3

**Output 2:**

The equation of parabola is  $y=1.42+-1.07x+0.55x^2$



**3) Aim: Write a python program to find Karl Pearson's correlation coefficient between x and y variables.**

**Source Code:**

```
from math import sqrt
import numpy as np
print("Enter x :")
x=[int(x) for x in input().split()]
print("Enter y :")
y=[int(x) for x in input().split()]
x=np.array(x)
y=np.array(y)
xy=x*y
x2=x**2
y2=y**2
n=len(x)
num=n*sum(xy)-(sum(x)*sum(y))
den=sqrt((n*sum(x2)-sum(x)**2)*(n*sum(y2)-sum(y)**2))
r=num/den
print("Correlation coefficient is :",round(r,4))
```

**Output 1**

```
Enter x :
3 7 4 2 0 4 1 2
Enter y :
11 18 9 4 7 6 3 8

Correlation coefficient is : 0.7867
```

**Output 2**

```
Enter x :
65 66 67 67 68 69 70 72
Enter y :
67 68 65 68 72 72 69 71

Correlation coefficient is : 0.603
```



**4) Aim: Write a python program to find the Spearman's correlation coefficient between x and y variables.**

**Source Code:**

```
import pandas as pd
s=input()
a=[float(i) for i in s.split(" ")]
s1=input()
b=[float(i) for i in s1.split(" ")]
n=len(a)
data=pd.DataFrame({'A':a,'B':b})
```

```
def rank(a):
    s=sorted(a)
    n=len(a)
    s=s[::-1]
    i=0
    d=[]
    count=[]
    while i<n:
        k=s.count(s[i])
        if k==1:
            d.append(i+1)
            i=i+1
        else:
            m=0
            for j in range(i+1,i+k+1):
                m=m+j
            m=m/k
            for j in range(k):
                d.append(m)
            i=i+k
            count.append(k)
    r=[]
    for i in range(n):
        j=s.index(a[i])
        r.append(d[j])
    return r,count
```

```
r_x,c_x=rank(a)
r_y,c_y=rank(b)
data['Rank of x']=r_x
data['Rank of y']=r_y
```

```

di=[]
di2=[]
for i in range(len(a)):
    k=r_x[i]-r_y[i]
    di.append(k)
    di2.append(k**2)
data['di']=di
data['di2']=di2
print(data)

def correction_factor(c):
    if len(c)!=0:
        m=c[0]
        cf=(m*(m**2-1))/12
        return cf
    else:
        return 0
cf_x=correction_factor(c_x)
cf_y=correction_factor(c_y)
sum_di2=sum(di2)+cf_x+cf_y
print("Correction factor of a",cf_x)
print("Correction factor of b",cf_y)
print("di2 after correction factor is added",sum_di2)
r=1-((6*sum_di2)/(n*(n**2-1)))
print("Rank Correlation coefficient : ",round(r,4))

```

**Input 1**

```

68 64 75 50 64 80 75 40 55 64
62 58 68 45 81 60 68 48 50 70

```

**Output 1:**

	A	B	Rank of x	Rank of y	di	di2
0	68.0	62.0	4.0	5.0	-1.0	1.0
1	64.0	58.0	6.0	7.0	-1.0	1.0
2	75.0	68.0	2.5	3.5	-1.0	1.0
3	50.0	45.0	9.0	10.0	-1.0	1.0
4	64.0	81.0	6.0	1.0	5.0	25.0
5	80.0	60.0	1.0	6.0	-5.0	25.0
6	75.0	68.0	2.5	3.5	-1.0	1.0
7	40.0	48.0	10.0	9.0	1.0	1.0
8	55.0	50.0	8.0	8.0	0.0	0.0
9	64.0	70.0	6.0	2.0	4.0	16.0

Correction factor of a 0.5  
 Correction factor of b 0.5  
 di2 after correction factor is added 73.0

Rank Correlation coefficient : 0.5576

**Input 2:**

115 109 112 87 98 120 98 100 98 118  
 75 73 85 70 76 82 65 73 68 80

**Output 2:**

	A	B	Rank of x	Rank of y	di	di2
0	115.0	75.0	3.0	5.0	-2.0	4.00
1	109.0	73.0	5.0	6.5	-1.5	2.25
2	112.0	85.0	4.0	1.0	3.0	9.00
3	87.0	70.0	10.0	8.0	2.0	4.00
4	98.0	76.0	8.0	4.0	4.0	16.00
5	120.0	82.0	1.0	2.0	-1.0	1.00
6	98.0	65.0	8.0	10.0	-2.0	4.00
7	100.0	73.0	6.0	6.5	-0.5	0.25
8	98.0	68.0	8.0	9.0	-1.0	1.00
9	118.0	80.0	2.0	3.0	-1.0	1.00

Correction factor of a 2.0  
 Correction factor of b 0.5  
 di2 after correction factor is added 45.0

Rank Correlation coefficient : 0.7273

**5) Aim: Write a python program to classify the data based on one way Anova.****Source Code:**

```

import numpy as np
import scipy.stats as stats
print("Enter treatment 1:")
a=[int(x) for x in input().split()]
print("Enter treatment 2:")
b=[int(x) for x in input().split()]
print("Enter treatment 3:")
c=[int(x) for x in input().split()]

#read level of significance
print("Level of significance :")
alpha=float(input())

fa=np.array(a)
fb=np.array(b)
fc=np.array(c)

print("Treatment 1 :",fa)
print("Treatment 2 :",fb)
print("Treatment 3 :",fc)

#calculate rss,cf,sst,sstr,sse
N=np.size(fa)+np.size(fb)+np.size(fc)
rss=np.sum(fa**2)+np.sum(fb**2)+np.sum(fc**2)
cf=(np.sum(fa)+np.sum(fb)+np.sum(fc))**2/(N)
sst=rss-cf
sstr=(np.sum(fa)**2/np.size(fa)+np.sum(fb)**2/np.size(fb)+np.sum(fc)**2/np.size(fc))-cf
sse=sst-sstr
print("rss=",rss)
print("cf=",cf)
print("sst=",sst)
print("sstr=",sstr)
print("sse=",sse)

#degree of freedom
k=3
d1=k-1
d2=N-k
print("Degree of freedom of treatments =",d1)
print("Degree of freedom of error =",d2)

```

```
#calculate f
msstr=sstr/d1
msse=sse/d2
F=msstr/msse
if F<1:
    F=msse/msstr
print("Calculated value :",F)

#table value
tablevalue=stats.f.ppf(1-alpha,d1,d2)
print("Table value :",round(tablevalue,4))

#testing
if tablevalue>F:
    print("H0 is accepted.")
else:
    print("H0 is rejected.")
```

**Input 1:**

Enter treatment 1:

13 10 8 11 8

Enter treatment 2:

13 11 14 14

Enter treatment 3:

4 1 3 4 2 4

Level of significance :

0.05

**Output 1:**

Treatment 1 : [13 10 8 11 8]

Treatment 2 : [13 11 14 14]

Treatment 3 : [4 1 3 4 2 4]

rss= 1262

cf= 960.0

sst= 302.0

sstr= 270.0

sse= 32.0

Degree of freedom of treatments = 2

Degree of freedom of error = 12

Calculated value : 50.625

Table value : 3.8853

H<sub>0</sub> is rejected.

**Input 2:**

Enter treatment 1:

90 82 79 98 83 91

Enter treatment 2:

105 89 93 104 89 95 86

Enter treatment 3:

83 89 80 94

Level of significance :

0.05

**Output 2:**

Treatment 1 : [90 82 79 98 83 91]

Treatment 2 : [105 89 93 104 89 95 86]

Treatment 3 : [83 89 80 94]

rss= 138638

cf= 137700.0

sst= 938.0

sstr= 234.4523809523671

sse= 703.5476190476329

Degree of freedom of treatments = 2

Degree of freedom of error = 14

Calculated value : 2.3327016142676427

Table value : 3.7389

H<sub>0</sub> is accepted.

**6) Aim: Write a python program to classify the data based on two way Anova****Source Code:**

```

import scipy.stats as stats
import pandas as pd

def input_data(k):
    l=[]

    for i in range(k):
        print("Enter treatment ",i+1)
        s=[int(x) for x in input().split()]
        l.append(s)
    return l
print("Enter number of treatments :")
k=int(input())
print("Enter number of blocks :")
h=int(input( ))
l=input_data(k)

def dataframe(l):
    df=pd.DataFrame(l)
    col=[]
    for i in range(h):
        col.append("B"+str(i+1))
    df.columns=col
    index=[]
    for i in range(k):
        index.append("T"+str(i+1))
    df.index=index
    print("Given data :")
    print(df)
dataframe(l)

def calculations(l):
    G=0
    flag_ftr=0
    flag_fb=0
    Ti2=0
    for i in range(k):
        G=G+sum(l[i])
        Ti2=Ti2+sum(l[i])**2
    bj2=0

```

```

rss=0
for j in range(h):
    bj=0
    for i in range(k):
        bj=bj+l[i][j]
        rss=rss+l[i][j]**2
    bj2=bj2+bj**2
cf=(G**2)/(k*h)
st2=rss-cf
str2=Ti2*(1/h)-cf
sb2=bj2*(1/k)-cf
se2=st2-str2-sb2
print("Row sum of squares =",rss)
print("Correction factor =",cf)
print("Sum of squares due to total =",st2)
print("Sum of squares due to treatments =",str2)
print("Sum of squares due to blocks =",sb2)
print("Sum of squares due to error =",se2)
mstr=str2/(k-1)
msb=sb2/(h-1)
mse=se2/((k-1)*(h-1))
ftr=mstr/mse
fb=msb/mse
if ftr<1:
    ftr=mse/mstr
    flag_ftr=1
if fb<1:
    fb=mse/msb
    flag_fb=1
return ftr,fb,flag_ftr,flag_fb
ftr,fb,flag_ftr,flag_fb=calculations(l)
print("Calculated values")
print("Treatments :",round(ftr,4))
print("Blocks :",round(fb,4))
if flag_ftr==1:
    ft_tr=stats.f.ppf(0.95,(k-1)*(h-1),(k-1))
else:
    ft_tr=stats.f.ppf(0.95,(k-1),(k-1)*(h-1))
if flag_fb==1:
    ft_b=stats.f.ppf(0.95,(k-1)*(h-1),(h-1))
else:
    ft_b=stats.f.ppf(0.95,(h-1),(k-1)*(h-1))
print("Table values")

```



```
print("Treatments :",round(ft_tr,4))
print("Blocks :",round(ft_b,4))
```

```
if ftr>ft_tr:
    print("H0(tr) is rejected.")
else:
    print("H0(tr) is accepted.")
if fb>ft_b:
    print("H0(b) is rejected.")
else:
    print("H0(b) is accepted.")
```

**Input 1:**

Enter number of treatments :

3

Enter number of blocks :

4

13 7 9 3

6 6 3 1

11 5 15 5

**Output 1:**

Given data :

B1 B2 B3 B4

T1 13 7 9 3

T2 6 6 3 1

T3 11 5 15 5

Row sum of squares = 786

Correction factor = 588.0

Sum of squares due to total = 198.0

Sum of squares due to treatments = 56.0

Sum of squares due to blocks = 90.0

Sum of squares due to error = 52.0

Calculated values

Treatments : 3.2308

Blocks : 3.4615

Table values

Treatments : 5.1433

Blocks : 4.7571

H0(tr) is accepted.

H0(b) is accepted.

**Input 2:**

Enter number of treatments :

4

Enter number of blocks :

5

Enter treatment 1

75 73 59 69 84

Enter treatment 2

83 72 56 70 92

Enter treatment 3

86 61 53 72 88

Enter treatment 4

73 67 62 79 95

**Output 2:**

Given data :

	B1	B2	B3	B4	B5
T1	75	73	59	69	84
T2	83	72	56	70	92
T3	86	61	53	72	88
T4	73	67	62	79	95

Row sum of squares = 110607

Correction factor = 107898.05

Sum of squares due to total = 2708.949999999997

Sum of squares due to treatments = 42.94999999999709

Sum of squares due to blocks = 2326.699999999997

Sum of squares due to error = 339.30000000000029

Calculated values

Treatments : 1.975

Blocks : 20.5721

Table values

Treatments : 8.7446

Blocks : 3.2592

H0(tr) is accepted.

H0(b) is rejected.

**7. Aim: Write a python program to fit a multiple regression model for any given data.**

**Source Code:**

```
#transpose
def transpose(arr):
    t=[]
    for i in range(len(arr[0])):
        T=[]
        for j in range(len(arr)):
            T.append(arr[j][i])
        t.append(T)
    return t

#multiplication
def mul(a,b):
    c=[]
    n=len(a)
    m=len(a[0])
    q=len(b[0])
    for i in range(n):
        C=[]
        for j in range(q):
            multi=0
            for k in range(m):
                multi=multi+a[i][k]*b[k][j]
            C.append(round(multi,4))
        c.append(C)
    return c
```

```
#inverse
def inverse(a):
    n=len(a)
    m=len(a[0])
    cofactor=[]
    det=0
    if n==2 and m==2:
        cofactor.append([a[1][1],-1*a[0][1]])
        cofactor.append(-1*[a[1][0],a[0][0]])
        det=cofactor[0][0]*cofactor[1][1]-cofactor[0][1]*cofactor[1][0]
    else:
        for i in range(n):
            co=[]
            for j in range(m):
                c=[]
                for k in range(n):
                    for o in range(m):
                        if i!=k and j!=o:
                            c.append(a[k][o])
                if (i+j)%2!=0:
                    q=-1*(c[0]*c[3]-c[1]*c[2])
                else:
                    q=(c[0]*c[3]-c[1]*c[2])
                co.append(q)
            if i==0:
```

```

        det=det+a[i][j]*q
    cofactor.append(co)
    inv=transpose(cofactor)
    ine=inv/det
    return ine

import numpy as np
import pandas as pd
import scipy.stats as s

df=pd.read_csv('mutliple.csv')
print(df)

#beta_hat=(x'x)-1(x'y)
df['x0']=[1]*len(df)
x=df[['x0','x1','x2']].to_numpy()
y=df[['y']].to_numpy()
#calculate x'x and x'y and (x'x)(x'y)
x1=transpose(x)
x1x=mul(x1,x)
inv=inverse(x1x)
x1y=mul(x1,y)
beta_hat=mul(inv,x1y)
print("y =",beta_hat[0][0],"+ x1",beta_hat[1][0],"+ x2",beta_hat[2][0])
#test of goodness of fit using coefficient of determination
y_hat=[]
error=[]
for i in range(len(x)):
    s=beta_hat[0][0]+x[i][1]*beta_hat[1][0]+x[i][2]*beta_hat[2][0]

```

```
y_hat.append(round(s,4))
error.append(round(y[i][0]-s,4))
d=pd.DataFrame({'y_hat':y_hat,'error':error})
print(d)
#calculations
sse=sum(np.array(error)**2)
y_bar=sum(y)/len(y)
sst=sum((y-y_bar)**2)
ssr=sst-sse
R2=ssr/sst
print(R2)
if R2<0.9:
    print("The Regression model is not good fit")
else:
    print("The Regression model is good fit")
#degree of freedom
n1=len(x[0])-1
n2=len(x)-len(x[0])
#to test goodness of fit using anova
msr=ssr/n1
mse=sse/n2
f=msr/mse
f=f[0]
if f<1:
    f=mse/msr
print("Calculated value",round(f,4))
```

```

#table value
f_tab=s.f.ppf(0.95,n1,n2)
print("Table value",round(f_tab,4))
if f<f_tab:
    print("H0 is Accepted")
    print("Hence we conclude that there is no regression parameter that influence in the
model.")
else:
    print("H0 is Rejected")
    print("Hence we conclude that there is atleast one regression parameter that influence in the
model.")
#test of individual variables
cij=[inv[0][0],inv[1][1],inv[2][2]]
t_cal=[]
print("Calculated value :")
for i in range(len(cij)):
    se=sqrt(mse*cij[i])
    t=beta_hat[i][0]
    t_cal.append(round(t/se,4))
    print("beta",i,"=",t_cal[i])
#table value
t_tab=stats.t.ppf(1-0.05/2,n2)
print("Table value",round(t_tab,4))
weak_variable=-1
for i in range(len(t_cal)):
    if t_tab>t_cal[i]:
        print("H0 is accepted.\nHence the parameter beta",i,"is not influencing the model")

```

```

weak_variable=i
else:
    print("H0 is rejected.\nHence the parameter beta",i,"is influencing the model")
print("Therefore,The weak variable is beta",weak_variable)

```

### Output 1:

```

x1 x2 y
0 -5 5 11
1 -4 4 11
2 -1 1 8
3 2 -3 2
4 2 -2 5
5 3 -2 5
6 3 -3 4

```

$$y = 6.5714 + 1.0 x_1 + 2.0 x_2$$

```

y_hat error
0 11.5714 -0.5714
1 10.5714 0.4286
2 7.5714 0.4286
3 2.5714 -0.5714
4 4.5714 0.4286
5 5.5714 -0.5714
6 3.5714 0.4286

```

#R2

R2= 0.97674419

The Regression model is good fit

#anova

Calculated value 84.0

Table value 6.9443

H0 is Rejected

Hence we conclude that there is atleast one regression parameter that influence in the model.

#testing weak variables

Calculated value :

beta 0 = 26.558



beta 1 = 2.1523

beta 2 = 4.3046

Table value 2.7764

H<sub>0</sub> is rejected.

Hence the parameter beta 0 is influencing the model

H<sub>0</sub> is accepted.

Hence the parameter beta 1 is not influencing the model

H<sub>0</sub> is rejected.

Hence the parameter beta 2 is influencing the model

Therefore, The weak variable is beta 1

### Output 2:

	month	y	x1	x2
0	1	100	9	62
1	2	110	8	58
2	3	105	7	64
3	4	94	14	60
4	5	95	12	63
5	6	99	10	57
6	7	104	7	55
7	8	108	4	56
8	9	105	6	59
9	10	98	5	61
10	11	105	7	57
11	12	110	6	60

$$y = 133.4605 + -1.2485 x_1 + -0.351 x_2$$

	y_hat	error
0	100.4620	-0.4620
1	103.1145	6.8855
2	102.2570	2.7430
3	94.9215	-0.9215
4	96.3655	-1.3655
5	100.9685	-1.9685
6	105.4160	-1.4160
7	108.8105	-0.8105
8	105.2605	-0.2605
9	105.8070	-7.8070

10 104.7140 0.2860  
11 104.9095 5.0905

[0.5415279]

The Regression model is not good fit

Calculated value 5.3152

Table value 4.2565

H0 is Rejected

Hence we conclude that there is atleast one regression parameter that influence in the model.

Calculated value :

beta 0 = 5.0882

beta 1 = -2.8079

beta 2 = -0.7711

Table value 2.2622

H0 is rejected.

Hence the parameter beta 0 is influencing the model

H0 is accepted.

Hence the parameter beta 1 is not influencing the model

H0 is accepted.

Hence the parameter beta 2 is not influencing the model

Therefore, The weak variable is beta 2

**8) Aim: Write a python program to fit a multivariate regression model for the given data.**

**Source Code:**

```
import numpy as np
import pandas as pd
import scipy.stats as stats
df=pd.read_csv('multivariate.csv')
print(df)

#beta ground=(x'x)-1*(x'y)
df['x0']=[1]*len(df)
x=df[['x0','x1','x2','x3']].to_numpy()
y=df[['y1','y2']].to_numpy()
x1x=mul(transpose(x),x)
x1y=mul(transpose(x),y)
inv=np.linalg.inv(x1x)
beta=np.array(mul(inv,x1y))
print("y1 =",beta[0][0],"+",beta[1][0],"x1 +",beta[2][0],"x2 +",beta[3][0],"x3")
print("y2 =",beta[0][1],"+",beta[1][1],"x1 +",beta[2][1],"x2 +",beta[3][1],"x3")
#test of goodness of fit using coefficient of determination
y_hat=x@beta
error=y-y_hat
sse=np.sum(error**2,axis=0)
mean=np.sum(y,axis=0)/len(y)
sst=np.sum((y-mean)**2,axis=0)
ssr=sst-sse
R2=ssr/sst
def test(r2):
```

```
if r2<0.9:
    print("The Regression model is not good fit")
else:
    print("The Regression model is good fit")
print("For y1 R2 is:",round(R2[0],4))
test(R2[0])
print("For y2 R2 is:",round(R2[1],4))
test(R2[1])
#to test goodness of fit using anova
def cal_value(f_cal):
    if f_cal<1:
        f_cal=mse/msr
        print("Calculated value",round(f_cal,4))
#degree of freedom
n1=len(x[0])-1
n2=len(x)-len(x[0])
msr=ssr/n1
mse=sse/n2
f=msr/mse
cal_value(f[0])
cal_value(f[1])
#table value
f_tab=stats.f.ppf(0.95,n1,n2)
print("Table value",round(f_tab,4))
def test(f_cal):
    if f_cal<f_tab:
```

```

    print("H0 is Accepted")

    print("Hence we conclude that there is no regression parameter that influence in the
model.")

    else:

        print("H0 is Rejected")

        print("Hence we conclude that there is atleast one regression parameter that influence in
the model.")

print("For y1 :")
test(f[0])
print("For y2 :")
test(f[1])

#testing individual parameters
t=[]

for i in range(len(inv)):

    se1=sqrt(mse[0]*inv[i][i])
    se2=sqrt(mse[1]*inv[i][i])
    t1=round(beta[i][0]/se1,4)
    t2=round(beta[i][1]/se2,4)
    t.append([t1,t2])

for i in range(len(t[0])):

    print("t calculated value for y"+str(i+1))

    for j in range(len(t)):

        print("beta"+str(j)+":",t[j][i])

t_tab=round(stats.t.ppf(1-0.05/2,n2),4)

print("Table Value :",t_tab)

#testing the weak parameters

for i in range(len(t[0])):

```

```

print("For y"+str(i+1)+" :")
weak_variable=[]
for j in range(len(t)):
    if t_tab>t[j][i]:
        print("H0 is accepted.\nHence the parameter beta",i,"is not influencing the model")
        weak_variable.append(j)
    else:
        print("H0 is rejected.\nHence the parameter beta",i,"is influencing the model")
print("\nTherefore,The weak variable is beta",weak_variable)

```

**Output 1:**

#output

	month	y1	y2	x1	x2	x3
0	1	10	100	9	62	1.0
1	2	12	110	8	58	1.3
2	3	11	105	7	64	1.2
3	4	9	94	14	60	0.8
4	5	9	95	12	63	0.8
5	6	10	99	10	57	0.9
6	7	11	104	7	55	1.0
7	8	12	108	4	56	1.2
8	9	11	105	6	59	1.1
9	10	10	98	5	61	1.0
10	11	11	103	7	57	1.2
11	12	12	110	6	60	1.2

$$y_1 = 10.897 + -0.0449 x_1 + -0.0877 x_2 + 5.0355 x_3$$

$$y_2 = 91.0972 + -0.064 x_1 + -0.2944 x_2 + 27.8353 x_3$$

For  $y_1$   $R^2$  is: 0.9238

The Regression model is good fit

For  $y_2$   $R^2$  is: 0.8655

The Regression model is not good fit

Calculated value 32.3272

Calculated value 17.1613

Table value 4.0662

For  $y_1$  :

$H_0$  is Rejected

Hence we conclude that there is atleast one regression parameter that influence in the model.

For  $y_2$  :

$H_0$  is Rejected

Hence we conclude that there is atleast one regression parameter that influence in the model.

t calculated value for  $y_1$

beta0: 4.2373

beta1: -0.8276

beta2: -2.2751

beta3: 5.4618

t calculated value for  $y_2$

beta0: 5.2648

beta1: -0.1753

beta2: -1.1351

beta3: 4.4872

Table Value : 2.306

For  $y_1$  :

$H_0$  is rejected.

Hence the parameter beta 0 is influencing the model

$H_0$  is accepted.

Hence the parameter beta 0 is not influencing the model

$H_0$  is accepted.

Hence the parameter beta 0 is not influencing the model

$H_0$  is rejected.

Hence the parameter beta 0 is influencing the model

Therefore, The weak variable is beta [1, 2]

For  $y_2$  :

$H_0$  is rejected.

Hence the parameter beta 1 is influencing the model

$H_0$  is accepted.

Hence the parameter beta 1 is not influencing the model

$H_0$  is accepted.

Hence the parameter beta 1 is not influencing the model

$H_0$  is rejected.

Hence the parameter beta 1 is influencing the model

Therefore, The weak variable is beta [1, 2]



**Output 2:**

	y1	y2	x1	x2	x3
0	9	150	3	62	1.5
1	2	98	8	58	0.7
2	5	75	3	64	3.6
3	2	24	14	60	1.8
4	4	95	15	63	1.8
5	13	39	18	57	6.9
6	10	14	23	55	1.1
7	16	88	4	56	1.0
8	6	15	6	59	1.0
9	2	48	12	61	1.6
10	1	73	9	57	1.3
11	12	91	10	60	1.2

$$y1 = 59.8734 + -0.2111 x1 + -0.8916 x2 + 1.0523 x3$$

$$y2 = -84.0353 + -3.0545 x1 + 3.1483 x2 + -1.7594 x3$$

For y1 R2 is: 0.2755

The Regression model is not good fit

For y2 R2 is: 0.3696

The Regression model is not good fit

Calculated value 1.014

Calculated value 1.5632

Table value 4.0662

For y1 :

H0 is Accepted

Hence we conclude that there is no regression parameter that influence in the model.

For y2 :

H0 is Accepted

Hence we conclude that there is no regression parameter that influence in the model.

t calculated value for y1

beta0: 1.6625

beta1: -0.7635

beta2: -1.5111

beta3: 1.1341

t calculated value for y2

beta0: -0.3096

beta1: -1.4659

beta2: 0.708

beta3: -0.2516

Table Value : 2.306

For y1 :

H0 is accepted.

Hence the parameter beta 0 is not influencing the model

H0 is accepted.

Hence the parameter beta 0 is not influencing the model

H0 is accepted.

Hence the parameter beta 0 is not influencing the model

H0 is accepted.

Hence the parameter beta 0 is not influencing the model

Therefore, The weak variable is beta [0, 1, 2, 3]

For y2 :

H0 is accepted.

Hence the parameter beta 1 is not influencing the model

H0 is accepted.

Hence the parameter beta 1 is not influencing the model

H0 is accepted.

Hence the parameter beta 1 is not influencing the model

H0 is accepted.

Hence the parameter beta 1 is not influencing the model

Therefore, The weak variable is beta [0, 1, 2, 3]

**9) Aim: Write a python program to classify the treatments based on MANOVA Test.****Source Code:**

```
import numpy as np
import scipy.stats as stats
from math import sqrt
m=int(input('Enter number of treatments : '))
t=[]
for i in range(m):
    print("Enter Treatment",(i+1))
    y1=[int(x) for x in input().split()]
    y2=[int(x) for x in input().split()]
    t.append([y1,y2])
for i in range(m):
    print("Treatment",i+1)
    print(t[i][0])
    print(t[i][1])
t_mean=[]
total=[0,0]
t_size=0
for i in range(m):
    y1=np.array(t[i][0])
    y2=np.array(t[i][1])
    t_mean.append([sum(y1)/len(y1),sum(y2)/len(y2)])
    total[0]+=sum(y1)
    total[1]+=sum(y2)
    t_size+=len(y1)
```

```

total[0]=total[0]/t_size
total[1]=total[1]/t_size
print("Yi mean")
for i in range(m):
    print(t_mean[i])
print("Y mean")
print(total)
def calculations(t,t_mean,total,y):
    sse=0
    sst=0
    for i in range(m):
        for j in range(len(t[i][0])):
            if y==-1:
                sse+=t[i][0][j]*t[i][1][j]-t_mean[i][0]*t_mean[i][1]
                sst+=t[i][0][j]*t[i][1][j]-total[0]*total[1]
            else:
                sse+=(t[i][y][j]-t_mean[i][y])**2
                sst+=(t[i][y][j]-total[y])**2
    return sse,sst
sse_y1,sst_y1=calculations(t,t_mean,total,0)
ssr_y1=sst_y1-sse_y1
sse_y2,sst_y2=calculations(t,t_mean,total,1)
ssr_y2=sst_y2-sse_y2
sse_y,sst_y=calculations(t,t_mean,total,-1)
ssr_y=sst_y-sse_y
print("For y1 :")

```

```

print("sse =",sse_y1,end=" , ")
print("sst =",sst_y1,end=" , ")
print("ssr =",ssr_y1)
print("For y2 :")
print("sse =",sse_y2,end=" , ")
print("sst =",sst_y2,end=" , ")
print("ssr =",ssr_y2)
print("Cross product values of y1 and y2 :")
print("sse =",sse_y,end=" , ")
print("sst =",sst_y,end=" , ")
print("ssr =",ssr_y)
#sum of squares
B=np.array([ssr_y1,ssr_y,ssr_y,ssr_y2]).reshape(2,2)
W=np.array([sse_y1,sse_y,sse_y,sse_y2]).reshape(2,2)
T=np.array([sst_y1,sst_y,sst_y,sst_y2]).reshape(2,2)
print("Regression :\n",B)
print("Error :\n",W)
print("Total :\n",T)
#Degree of Freedom
d1=m-1
n=0
for i in range(m):
    n=n+len(t[i][0])
d2=n-m
print("Degree of Freedom :",d1,"",d2)
def det(A):

```

```

return A[0][0]*A[1][1]-A[0][1]*A[1][0]
wilks=det(W)/det(T)
print("Wilk's Value :",round(wilks,4))
f=((n-m-1)/(m-1))*(1-sqrt(wilks))/sqrt(wilks)
print("Calculate value :",round(f,4))
tab=stats.f.ppf(0.95,2*(m-1),2*(n-m-1))
print("Table Value :",round(tab,4))
if f>tab:
    print("H0 is Rejected.Hence we conclude that there is no homogeneity among regression
model")
else:
    print("H0 is Accepted.Hence we conclude that there is homogeneity among regression model")

```

**Input 1:**

Enter number of treatments : 3

Enter Treatment 1

2 3 5 2

3 4 4 5

Enter Treatment 2

4 5 6

8 6 7

Enter Treatment 3

7 8 10 9 7

6 7 8 5 6

**Output 1:**

Treatment 1

[2, 3, 5, 2]

[3, 4, 4, 5]

Treatment 2

[4, 5, 6]

[8, 6, 7]

Treatment 3

[7, 8, 10, 9, 7]

[6, 7, 8, 5, 6]

Yi mean

[3.0, 4.0]

[5.0, 7.0]

[8.2, 6.4]

Y mean

[5.666666666666667, 5.75]

For y1 :

sse = 14.799999999999997 , sst = 76.66666666666667 , ssr = 61.866666666666674

For y2 :

sse = 9.2 , sst = 28.25 , ssr = 19.05

Cross product values of y1 and y2 :

sse = 1.600000000000000156 , sst = 25.999999999999943 , ssr = 24.399999999999928

Regression :

[[61.86666667 24.4 ]

[24.4 19.05 ]]

Error :

[[14.8 1.6]

[ 1.6 9.2]]

Total :

[[76.66666667 26. ]

[26. 28.25 ]]

Degree of Freedom : 2 , 9

Wilk's Value : 0.0897

Calculate value : 9.3575

Table Value : 3.0069

H0 is Rejected.Hence we conclude that there is no homogeneity among regression model

**Input 2:**

Enter number of treatments : 3

Enter Treatment 1

9 6 9

3 2 7

Enter Treatment 2

0 2

4 0

Enter Treatment 3

3 1 2

8 9 7

**Output 2:**

Treatment 1

[9, 6, 9]

[3, 2, 7]

Treatment 2

[0, 2]

[4, 0]



Treatment 3

[3, 1, 2]

[8, 9, 7]

Yi mean

[8.0, 4.0]

[1.0, 2.0]

[2.0, 8.0]

Y mean

[4.0, 5.0]

For y1 :

sse = 10.0 , sst = 88.0 , ssr = 78.0

For y2 :

sse = 24.0 , sst = 72.0 , ssr = 48.0

Cross product values of y1 and y2 :

sse = 1.0 , sst = -11.0 , ssr = -12.0

Regression :

[[ 78. -12.]

[-12. 48.]]

Error :

[[10. 1.]

[ 1. 24.]]

Total :

[[ 88. -11.]

[-11. 72.]]

Degree of Freedom : 2 , 5

Wilk's Value : 0.0385

Calculate value : 8.1989

Table Value : 3.8379

H0 is Rejected. Hence we conclude that there is no homogeneity among regression model

**10) Aim: Write a python program to classify the given observation using Linear Discriminant Analysis.**

**Source Code:**

```
import pandas as pd
import numpy as np

sat=int(input('Enter incoming student SAT : '))
gpa=float(input('Enter incoming student GPA : '))

df=pd.read_csv('discriminant.csv')
print(df)

df['x0']=[1]*len(df)
x=df[['x0','x1','x2']]
x=np.array(x,dtype=float)
y=df[['y']].replace('yes',1)
y=df[['y']].replace('no',0)
y=np.array(y)

#beta hat
x1x=mul(transpose(x),x)
inv=np.linalg.inv(x1x)
x1y=mul(transpose(x),y)
beta_hat=mul(inv,x1y)

print("y = ",beta_hat[0][0],"+",beta_hat[1][0],"x1 +",beta_hat[2][0],"x2")
new_y=beta_hat[0][0]+beta_hat[1][0]*sat+beta_hat[2][0]*gpa
print("new_y =",new_y)

if round(new_y)==0:
    print("The value is nearer to zero.Therefore, the candidate will not graduated.")
else:
```

```
print("The value is nearer to one. Therefore, the candidate will graduated.")
```

**Input 1:**

Enter incoming student SAT : 1000

Enter incoming student GPA : 2.9

**Output 1:**

	x1	x2	y
0	1300	2.7	yes
1	1260	3.7	yes
2	1220	2.9	yes
3	1180	2.5	yes
4	1060	3.9	yes
5	1140	2.1	no
6	1100	3.5	no
7	1020	3.3	no
8	980	2.3	no
9	940	3.1	no

$y = -3.8392 + 0.0032 x_1 + 0.2395 x_2$

$new\_y = 0.055350000000000023$

The value is nearer to zero. Therefore, the candidate will not graduated.

**Output 2:**

Enter incoming student SAT : 1500

Enter incoming student GPA : 1.3

	x1	x2	y
0	1150	2.7	yes
1	1060	3.7	yes
2	1220	2.9	yes

3 1980 2.5 yes

4 1980 3.9 no

5 1840 2.1 no

$y = 2.264495 + -0.00080765 x_1 + -0.11979125 x_2$   
 $new\_y = 0.89727898$

The value is nearer to one. Therefore, the candidate will graduated.

### #Fishers' Linear Discriminant

```
import pandas as pd
import numpy as np
from math import log
print("Enter new matrix")
l=[float(x) for x in input().split()]
df=pd.read_csv('fishers.csv')
print(df)
x_1=[]
x_2=[]
for i in range(len(df)):
    if df['y'][i]==1:
        x_1.extend([df['x1'][i],df['x2'][i]])
    else:
        x_2.extend([df['x1'][i],df['x2'][i]])
x=df[['x1','x2']].to_numpy()
x_1=np.array(x_1).reshape(len(x_1)//2,2)
x_2=np.array(x_2).reshape(len(x_2)//2,2)
print("x :\n",x)
```

```

print("\nx1 :\n",x_1)
print("\nx2 :\n",x_2)
mean=np.mean(x,axis=0)
mean_1=np.mean(x_1,axis=0)
mean_2=np.mean(x_2,axis=0)
print("mean for x :",mean)
print("mean for x_1 :",mean_1)
print("mean for x_2 :",mean_2)
xm=x-mean
c=mul(transpose(xm),xm)
c=np.array(c)/len(x)
c_inv=np.linalg.inv(c)
print("Pooled covariance matrix :\n",c)
def fisherEquation(mean,c_inv,x,p):
    m_c=mul([mean],c_inv)
    f=mul(m_c,x)[0][0]-0.5*mul(m_c,transpose([mean]))[0][0]+log(p)
    return f
f1=fisherEquation(mean_1,c_inv,transpose([1]),len(x_1)/len(x))
f2=fisherEquation(mean_2,c_inv,transpose([1]),len(x_2)/len(x))
print("f1 =",f1)
print("f2 =",f2)
if f1>f2:
    print("The new observation",l,"is classified into group 1")
else:
    print("The new observation",l,"is classified into group 2")

```

**Input 1:**

Enter new matrix

5.1 3.2

**Output 1:**

x1 x2 y

0 1 2 1

1 2 3 1

2 3 3 1

3 4 5 1

4 5 5 1

5 4 2 0

6 5 0 0

7 5 2 0

8 3 2 0

9 5 3 0

10 6 3 0

x :

[[1 2]

[2 3]

[3 3]

[4 5]

[5 5]

[4 2]

[5 0]

[5 2]

[3 2]

[5 3]

[6 3]]

x1 :

[[1 2]

[2 3]

[3 3]

[4 5]

[5 5]]

x2 :

[[4 2]

[5 0]

[5 2]

[3 2]

[5 3]

[6 3]]

mean for x : [3.90909091 2.72727273]

mean for x\_1 : [3. 3.6]

mean for x\_2 : [4.66666667 2. ]

Pooled covariance matrix :

[[2.08264545 0.15702727]

[0.15702727 1.83470909]]

f1 = 6.48594263963573

f2 = 7.393764196429685

The new observation [5.1, 3.2] is classified into group 2

**Input 2:**

Enter new matrix

5 6

**Output 2:**

x1 x2 y

0 4 2 1

1 2 4 1

2 2 3 1

3 3 6 1

4 4 4 1

5 9 10 0

6 6 8 0

7 9 5 0

8 8 7 0

9 10 8 0

x :

[[4 2]

[2 4]

[2 3]

[3 6]

[4 4]

[9 10]

[6 8]

[9 5]

[8 7]

[10 8]]



x1 :

[[4 2]

[2 4]

[2 3]

[3 6]

[4 4]]

x2 :

[[9 10]

[6 8]

[9 5]

[8 7]

[10 8]]

mean for x : [5.7 5.7]

mean for x\_1 : [3. 3.8]

mean for x\_2 : [8.4 7.6 ]

Pooled covariance matrix :

[[86.1 50.1]

[50.1 58.1]]

f1 = 1.9839

f2 = 1.7051

The new observation [5, 6] is classified into group 1

**11) Aim: Write a python program to find Principal Components for the given variables.**

**Source Code:**

```
import numpy as np
import pandas as pd
n=int(input('Enter number of components :'))
l=[]
for i in range(n):
    k=[float(x) for x in input().split()]
    l.extend(k)
    m=len(k)
x=np.array(l).reshape(n,m)
x=transpose(x)
print("x\n",x)
mean=np.sum(x,axis=0)/m
print("Mean\n",mean)
x_mean=x-mean
c=mul(transpose(x_mean),x_mean)/m
print("c\n",c)
e_values,e_vectors=np.linalg.eig(c)
e1=np.argsort(e_values)[::-1]
e_values=e_values[e1]
e_vectors=e_vectors[:,e1]
print('Eigen values:\n',e_values)
print('Eigen vectors:\n',e_vectors)
z=[]
z_name=[]
```

```

sum=0
t_sum=np.sum(e_values)
for i in range(len(e_values)):
    sum=sum+e_values[i]
    z_name.append('z'+str(i+1))
    z.append(round(sum*100/t_sum,2))
print("Principal Components :",z)
threshold=int(input('Enter Threshold value :'))
c=0
for x in z:
    if x<=threshold+2:
        c=c+1
d={'principle components':z_name,'variance explained':e_values,'cumulative proportion of
total variance':z}
df=pd.DataFrame(d)
print(df)
print("Principal Components are:")
for i in range(c):
    print("z"+str(i+1)+" =",end=")
    for y in range(n):
        if y==n-1:
            print(round(e_vectors[y][i],4),"x"+str(y+1))
        else:
            print(round(e_vectors[y][i],4),"x"+str(y+1)+" + ",end=")
p=mul(x,transpose(e_vectors[:c]))
p=transpose(p)
d={}

```

```
for i in range(c):  
d.update({'z'+str(i+1):p[i]})  
df=pd.DataFrame(d)  
df
```

**Input 1:**

Enter number of components :2

2 1 0 -1

4 3 1 0.5

**Output 1:**

x

[[ 2. 4.]

[ 1. 3.]

[ 0. 1.]

[-1. 0.5]]

Mean

[0.5 2.125]

c

[[1.25 1.5625 ]

[1.5625 2.046875]]

Eigen values:

[3.26093826 0.03593674]

Eigen vectors:

[[-0.6135581 -0.78964958]

[-0.78964958 0.6135581 ]]

Principal Components : [98.91, 100.0]

Enter Threshold value :99

Regd. No :

	principle components	variance explained	cummulative proportion of total variance
0	z1	3.260938	98.91
1	z2	0.035937	100.00

Principal Components are:

$$z1 = -0.6136 x1 + -0.7896 x2$$

$$z2 = -0.7896 x1 + 0.6136 x2$$

	z1	z2
0	-4.385715	0.874933
1	-2.982507	1.051025
2	-0.789650	0.613558
3	0.218733	1.096429

**Input 2:**

Enter number of components :3

7 4 6 8 8 7 5 9 7 8

4 1 3 6 5 2 3 5 4 2

3 8 5 1 7 9 3 8 5 2

**Output 2:**

x

[[7. 4. 3.]

[4. 1. 8.]

[6. 3. 5.]

[8. 6. 1.]

[8. 5. 7.]

[7. 2. 9.]

[5. 3. 3.]

[9. 5. 8.]

[7. 4. 5.]

[8. 2. 2.]]

Mean

[6.9 3.5 5.1]

c

[[ 2.09 1.45 -0.39]

[ 1.45 2.25 -1.15]

[-0.39 -1.15 7.09]]

Eigen values:

[7.44654832 3.30851634 0.67493534]

Eigen vectors:

[[ -0.1375708 0.69903712 -0.70172743]

[ -0.25045969 0.66088917 0.70745703]

[ 0.95830278 0.27307986 0.08416157]]

Principal Components : [65.15, 94.1, 100.0]

Enter Threshold value :93

Principle components variance explained \

0 z1 7.446548

1 z2 3.308516

2 z3 0.674935

cumulative proportion of total variance

0 65.15

1 94.10

2 100.00

Principal Components are:

$$z_1 = -0.1376 x_1 + -0.2505 x_2 + 0.9583 x_3$$

$$z_2 = 0.699 x_1 + 0.6609 x_2 + 0.2731 x_3$$

	<b>z1</b>	<b>z2</b>
<b>0</b>	-0.272029	3.012710
<b>1</b>	-5.465065	5.318707
<b>2</b>	-2.236951	4.017195
<b>3</b>	2.391929	2.669115
<b>4</b>	-2.517473	6.252968
<b>5</b>	-5.880468	5.935674
<b>6</b>	-0.695925	2.852740
<b>7</b>	-3.356771	6.709965
<b>8</b>	-1.675484	4.427624
<b>9</b>	-1.105947	0.733015

**12) Aim: Write a python program to group the given variables using Factor Analysis.****Source Code:**

```
import numpy as np
import pandas as pd
from math import floor
from math import sqrt
n=int(input('Enter number of components :'))
l=[]
```

```
for i in range(n):
    k=[float(x) for x in input().split()]
    l.extend(k)
    m=len(k)
x=np.array(l).reshape(n,m)
x=transpose(x)
print("x\n",x)
mean=np.sum(x,axis=0)/m
x_mean=x-mean
s_d=[]
s_d=np.sum((x_mean)**2,axis=0)/(m-1)
for i in range(n):
    s_d[i]=sqrt(s_d[i])
x=x_mean/s_d
c=mul(transpose(x),x)/m
```

```
e_values,e_vectors=np.linalg.eig(c)
e1=np.argsort(e_values)[::-1]
e_values=e_values[e1]
e_vectors=e_vectors[:,e1]
print('Eigen values:\n',e_values)
print('Eigen vectors:\n',e_vectors)
```

```
z=[]
z_name=[]
sum=0
t_sum=np.sum(e_values)
for i in range(len(e_values)):
```



```

sum=sum+e_values[i]
z_name.append('z'+str(i+1))
z.append(round(sum*100/t_sum,2))

print("Principal Components :",z)
threshold=int(input('Enter Threshold value :'))
c=0
for x in z:
    if floor(x)<=threshold+2:
        c=c+1
print("Principal Components are:")
for i in range(c):
    print("z"+str(i+1)+" =",end="")
    for y in range(n):
        if y==n-1:
            print(round(e_vectors[y][i],4),"x"+str(y+1))
        else:
            print(round(e_vectors[y][i],4),"x"+str(y+1)+" + ",end="")

f=[]
for i in range(c):
    f1=[]
    for j in range(n):
        f1.append(sqrt(e_values[i])*e_vectors[j][i])
    f.append(f1)

f1=f
f=transpose(np.array(f))
h2=np.sum(f**2,axis=1)
s=np.sum(f,axis=0)
t_s=np.sum(h2)
f1[0].extend([s[0],s[0]*100/t_s])
f1[1].extend([s[1],s[1]*100/t_s])
h2=list(h2)
h2.extend([t_s,s[0]+s[1]])
df=pd.DataFrame({'variables':['Finance','Marketing','Business Policy','variance explained','%of variance explained'],'F1':f1[0],'F2':f1[1],'h^2':h2})
df

```

**Input 1:**

Enter number of components :3

3 7 10 3 10

6 3 9 9 6

5 3 8 7 5

**Output 1:**

x

[[ 3. 6. 5.]

[ 7. 3. 3.]

[10. 9. 8.]

[ 3. 9. 7.]

[10. 6. 5.]]

Eigen values:

[1.5851705 0.80664506 0.00818443]

Eigen vectors:

[[ 0.0212208 -0.99538347 -0.09360247]

[ 0.70623585 0.08119286 -0.70330551]

[ 0.70765853 -0.05118071 0.70469847]]

Principal Components : [66.05, 99.66, 100.0]

Enter Threshold value :97

Principal Components are:

 $z_1 = 0.0212 x_1 + 0.7062 x_2 + 0.7077 x_3$  $z_2 = -0.9954 x_1 + 0.0812 x_2 + -0.0512 x_3$ 

	variables	F1	F2	h^2
0	Finance	0.026718	-0.893988	0.799928
1	Marketing	0.889176	0.072922	0.795952
2	Business Policy	0.890967	-0.045967	0.795936
3	variance explained	1.806861	-0.867033	2.391816
4	%of variance explained	75.543493	-36.249995	0.939828

**Input 2:**

Enter number of components :3

2 4 1 5

2 8 4 0

8 3 1 2

**Output 2:**

x

[[2. 2. 8.]

[4. 8. 3.]

[1. 4. 1.]

[5. 0. 2.]]

Eigen values:

[0.89208647 0.82928548 0.52862805]

Eigen vectors:

[[-0.5267742 0.63311916 0.56715876]

[-0.34594212 -0.76916788 0.53731259]

[ 0.7764232 0.08683831 0.62420039]]

Principal Components : [39.65, 76.51, 100.0]

Enter Threshold value :75

Principal Components are:

 $z1 = -0.5268 x1 + -0.3459 x2 + 0.7764 x3$  $z2 = 0.6331 x1 + -0.7692 x2 + 0.0868 x3$ 

	variables	F1	F2	h^2
0	Finance	-0.497540	0.576551	0.579957
1	Marketing	-0.326743	-0.700444	0.597383
2	Business Policy	0.733334	0.079079	0.544033
3	variance explained	-0.090949	-0.044814	1.721372
4	%of variance explained	-5.283528	-2.603361	-0.135763