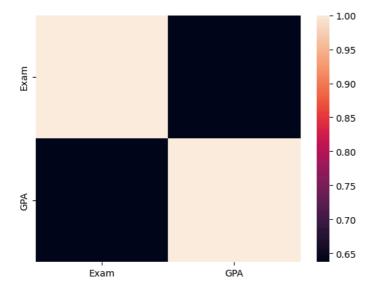
```
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import numpy as np
from scipy import stats
import statsmodels.api as sm
df=pd.read_csv('/content/slr (1).csv')
df.head()
                      \blacksquare
         Exam GPA
      0 1714 2.40
      1 1664 2.52
      2 1760 2.54
      3 1685 2.74
      4 1693 2.83
 Next steps:
              Generate code with df
                                      View recommended plots
DataFrame Observation
#continous_column=["Exam","GPA"]
#dependent=GPA
#independent=Exam
#Supervised regression
Data cleaning using pandas
Handling Null Values
Handling Duplicates
Data Type Conversion
Standardizing or Normalizing Data
Handling Text Data
Handling Date and Time Data
Handling Outliers
df.info()
     <class 'pandas.core.frame.DataFrame'>
     RangeIndex: 84 entries, 0 to 83
     Data columns (total 2 columns):
      # Column Non-Null Count Dtype
     0 Exam
                  84 non-null
                                   int64
      1 GPA
                  84 non-null
                                  float64
     dtypes: float64(1), int64(1) memory usage: 1.4 KB
df.isnull().sum()
             0
     Exam
     GPA
             0
     dtype: int64
df.corr()
                                 \blacksquare
                           GPA
                Exam
      Exam 1.000000 0.637184
      GPA 0.637184 1.000000
sns.heatmap(df.corr())
plt.show()
```



Hypothesis testing

0

Exam

1845.273810

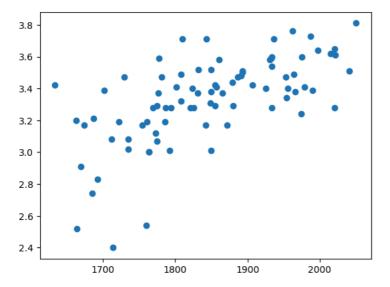
1854.16400

```
continous_column=["Exam","GPA"]
def central_limit_theorem(continous_column,sample_size,rage):
  result_centrallimit={}
  pop=df[continous_column].values
  population_mean=pop.mean()
  allsample=[]
  for i in range(rage):
     sample=np.random.choice(pop,sample_size)
     allsample.append(sample.mean())
  all sample mean=(np.mean(allsample))
  result_centrallimit.update({'Column Name':continous_column,
                        'Population mean':population_mean,
                        'Allsample mean':all_sample_mean
                       })
  if continous_column:
   H0_accepted=0
    H0_rejected=0
    for i in range(rage):
      sample1=df[continous_column].sample(frac=0.04)
      t_test,p_value=stats.ttest_1samp(sample1,df[continous_column].mean())
      if p_value<0.05:
        H0_rejected+=1
      else:
        H0_accepted+=1
      if H0_accepted>H0_rejected:
        result_centrallimit.update({'Onesamplettest':' H0-There is no significant difference','Ttest':t_test,
                                     'pvalue':p_value})
      else:
        result\_central limit.update(\{'Onesamplettest':' \ Ha-There \ is \ significant \ difference', 'Ttest':t\_test, 'pvalue':p\_value'\})
    return(result_centrallimit)
res=[]
for i in continous_column:
  s=central_limit_theorem(i,50,10)
  res.append(s)
pd.DataFrame(res)
                     Population
          Column
                                   Allsample
                                                                                          \blacksquare
                                                    Onesamplettest
                                                                       Ttest
                                                                                pvalue
             Name
                           mean
                                        mean
                                                                                          ılı
                                                      H0-There is no
```

0.514925 0.657867

significant difference

```
def twosamplettest(continous_column1,continous_column2,sample_size,rage):
  H0_accepted=0
  H0 rejected=0
  result={}
  allsample1=[]
   allsample2=[]
   for i in range(rage):
    sample1=df[continous column1].sample(frac=0.2)
    sample2=df[continous_column2].sample(frac=0.2)
     t_test,p_value=stats.ttest_ind(sample1,sample2)
    if p value<0.05:
       H0_rejected+=1
    else:
       H0_accepted+=1
     if H0_accepted>H0_rejected:
         result.update({
             'column':continous_column1+"&"+continous_column2,
             'twosamplettest': 'HO-There is no significant difference',
             't_test_value':t_test,
              'P_value':p_value})
    else:
         result.update({
              'column':continous_column1+"&"+continous_column2,
             'twosamplettest':' Ha-There is significant difference',
              't_test_value':t_test,
              'P_value':p_value})
   if continous_column1:
    H0_accepted=0
    H0 rejected=0
     for i in range(rage):
         column1=df[continous_column1]
         column2=df[continous_column2]
         sample1=np.random.choice(column1,sample_size)
         sample2=np.random.choice(column2,sample_size)
         allsample1.append(sample1.mean())
         allsample2.append(sample2.mean())
     t_test,p_value=stats.ttest_ind(allsample1,allsample2)
    if p_value<0.05:
             H0_rejected+=1
     else:
            H0_accepted+=1
    if H0_accepted>H0_rejected:
              result.update({
             'column':continous_column1+"&"+continous_column2,
             'twosamplettest central':'H0-There is no significant difference',
             't_test_valuone':t_test,
             'P_valueone':p_value})
    else:
              result.update({
              'column':continous_column1+"&"+continous_column2,
             'twosamplettest central':' Ha-There is significant difference',
             't_test_valuone':t_test,
             'P_valueone':p_value})
   return result
columns=continous_column
res1=[]
for i in range (len(columns)-1):
 column1=columns[i]
 for j in range(i+1,len(columns)):
      column2=columns[j]
      j=twosamplettest(column1,column2,50,10)
     res1.append(j)
pd.DataFrame(res1)
                                                    P_value twosamplettest
            column twosamplettest t_test_value
                                                                             t_test_valuon
                                                                     central
# consider independent as x and dependent as y
x1=df["Exam"]
y=df["GPA"]
plt.scatter(x1,y)
plt.show()
```



 $x=sm.add_constant(x1)$

 $\label{local_model_sm.ols} $$ model = sm.OLS(y,x)$$ $$ reating a model using dependent and dependent variable through ols method result=model.fit() $$ Training the model $$$

result.summary()

OLS Regression Results

Dep. Variable: **GPA** R-squared: 0.406 Model: OLS Adj. R-squared: 0.399 Method: Least Squares F-statistic: 56.05 Date: Sat, 30 Mar 2024 Prob (F-statistic): 7.20e-11 Log-Likelihood: 12.672 Time: 15:40:52 No. Observations: 84 AIC: -21.34 Df Residuals: 82 BIC: -16.48 Df Model: 1

Covariance Type: nonrobust

 coef
 std err
 t
 P>|t|
 [0.025 0.975]

 const 0.2750 0.409
 0.673 0.503 -0.538 1.088

 Exam 0.0017 0.000
 7.487 0.000 0.001 0.001
 0.002

 Omnibus:
 12.839
 Durbin-Watson:
 0.950

 Prob(Omnibus):
 0.002
 Jarque-Bera (JB):
 16.155

 Skew:
 -0.722
 Prob(JB):
 0.000310

 Kurtosis:
 4.590
 Cond. No.
 3.29e+04

Notes:

- [1] Standard Errors assume that the covariance matrix of the errors is correctly specified.
- [2] The condition number is large, 3.29e+04. This might indicate that there are

strong multicollinearity or other numerical problems

$\label{eq:continuous_params} $$y$$ hat=result.params[0]+result.params[1]*x1 $$y$$ hat$

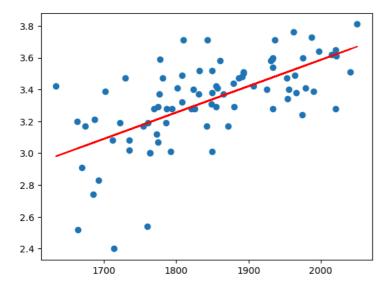
- 0 3.112890 1 3.030105
- 2 3.189051
- 3 3.064875
- 4 3.078120
- 79 3.480452
- 80 3.271836 81 3.564892
- 82 3.523500
- 83 3.669201

Name: Exam, Length: 84, dtype: float64

plt.scatter(x1,y)

plt.plot(x1,yhat,color='red')

plt.show()



Start coding or generate with AI.