Hypothesis Assignment

211091055 Gaytri Thakre

```
'''1) A F&B manager wants to determine whether there is any
In [4]:
        significant difference in the diameter of the cutlet between two
        units. A randomly selected sample of cutlets was collected from
        both units and measured? Analyze the data and draw inferences
        at 5% significance level. Please state the assumptions and tests
        that you carried out to check validity of the assumptions.'''
Out[4]: '1) A F&B manager wants to determine whether there is any\nsignificant difference in the diameter of the cut
        let between two\nunits. A randomly selected sample of cutlets was collected from\nboth units and measured? A
        nalyze the data and draw inferences\nat 5% significance level. Please state the assumptions and tests\nthat
        you carried out to check validity of the assumptions.'
In [5]: import pandas as pd
        import numpy as np
        from scipy.stats import ttest ind
        from scipy.stats import f oneway
        from scipy.stats import stats
In [6]: #Reading data
        data1 = pd.read csv(r'C:\Users\Hypothesis Assignment\Cutlets.csv')
        data1.head()
Out[6]:
            Unit A Unit B
```

Unit A Unit B 6.8090 6.7703 6.4376 7.5093 6.9157 6.7300 7.3012 6.7878 7.4488 7.1522

```
In [7]: unit1 diameters = data1['Unit A']
         unit2 diameters = data1['Unit B']
In [8]: #Defining significance level
         alpha = 0.05
         #Performing Ttest as 'N' is less
         t_stat, p_value = ttest_ind(unit1_diameters, unit2_diameters)
In [9]: p_value
Out[9]: 0.4722394724599501
In [10]: t_stat
Out[10]: 0.7228688704678063
In [11]: # Deciding Correct hypothesis
         if p value < alpha:</pre>
             print("Rejecting null hypothesis as there is a significant difference in the diameter of cutlets between
         else:
             print("Fail to reject null hypothesis there is no significant difference in the diameter of cutlets between
```

Fail to reject null hypothesis there is no significant difference in the diameter of cutlets between the two units.

In [12]: ''' 2) A hospital wants to determine whether there is any
difference in the average Turn Around Time (TAT) of reports
 of the laboratories on their preferred list. They collected a
 random sample and recorded TAT for reports of 4
 laboratories. TAT is defined as sample collected to report
 dispatch.

Analyze the data and determine whether there is any difference in average TAT among the different laboratories at 5% significance level. '''

Out[12]: ' 2) A hospital wants to determine whether there is any\ndifference in the average Turn Around Time (TAT) of reports\nof the laboratories on their preferred list. They collected a\nrandom sample and recorded TAT for reports of 4\nlaboratories. TAT is defined as sample collected to report\ndispatch.\n\nAnalyze the data and determine whether there is any\ndifference in average TAT among the different laboratories\nat 5% significance level. '

In [13]: data2 = pd.read_csv(r'C:\Users\Hypothesis Assignment\LabTAT.csv')
 data2

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	Laboratory 1	Laboratory 2	Laboratory 3	Laboratory 4
0	185.35	165.53	176.70	166.13
1	170.49	185.91	198.45	160.79
2	192.77	194.92	201.23	185.18
3	177.33	183.00	199.61	176.42
4	193.41	169.57	204.63	152.60
115	178.49	170.66	193.80	172.68
116	176.08	183.98	215.25	177.64
117	202.48	174.54	203.99	170.27
118	182.40	197.18	194.52	150.87
119	182.09	215.17	221.49	162.21

120 rows × 4 columns

```
In [14]: | tat laboratory1 = data2['Laboratory 1']
         tat laboratory2 = data2['Laboratory 2']
         tat laboratory3 = data2['Laboratory 3']
         tat_laboratory4 = data2['Laboratory 4']
In [15]: #Defining Significance Level
         alpha = 0.05
         #Performing anova test
         f stat, p value = stats.f oneway(tat laboratory1, tat laboratory2, tat laboratory3, tat laboratory4)
         C:\Users\HOME\AppData\Local\Temp\ipykernel 8148\1099572350.py:4: DeprecationWarning: Please use `f oneway` f
         rom the `scipy.stats` namespace, the `scipy.stats.stats` namespace is deprecated.
           f_stat, p_value = stats.f_oneway(tat_laboratory1, tat_laboratory2, tat laboratory3, tat laboratory4)
In [16]: | f stat
Out[16]: 118.70421654401437
In [17]: p value
Out[17]: 2.1156708949992414e-57
In [18]: # Deciding Correct hypothesis
         if p value < alpha:</pre>
             print("Rejecting null hypothesis as there is a significant difference in the diameter of cutlets between
         else:
             print("Fail to reject null hypothesis there is no significant difference in the diameter of cutlets between
         Rejecting null hypothesis as there is a significant difference in the diameter of cutlets between the two un
         its.
         '''3) Sales of products in four different regions is tabulated for males and females.
In [19]:
         Find if male-female buyer rations are similar across regions.'''
Out[19]: '3) Sales of products in four different regions is tabulated for males and females.\nFind if male-female buy
         er rations are similar across regions.'
```

```
In [20]: from scipy.stats import chi2 contingency
In [21]: # Creating table of given data
         data3 = np.array([[50, 142, 131, 70],
                          [435, 1523, 1356, 750]])
In [22]: #Defining alpha
         alpha = 0.05
         #performing chi-square test
         chi, p value, dof, expected = chi2 contingency(data3)
Out[22]: 1.595945538661058
In [23]: dof
Out[23]: 3
In [24]: p_value
Out[24]: 0.6603094907091882
In [25]: # Deciding Correct hypothesis
         if p value < alpha:</pre>
             print("Rejecting null hypothesis as there is a significant difference in the diameter of cutlets between
         else:
             print("Fail to reject null hypothesis there is no significant difference in the diameter of cutlets between
```

Fail to reject null hypothesis there is no significant difference in the diameter of cutlets between the two units.

```
In [26]:
    '''4) TeleCall uses 4 centers around the globe to process
    customer order forms. They audit a certain % of the
    customer order forms. Any error in order form
    renders it defective and has to be reworked before
    processing. The manager wants to check whether
    the defective % varies by centre. Please analyze the
    data at 5% significance level and help the manager
    draw appropriate inferences'''
```

Out[26]: '4) TeleCall uses 4 centers around the globe to process\ncustomer order forms. They audit a certain % of the \ncustomer order forms. Any error in order form\nrenders it defective and has to be reworked before\nprocess ing. The manager wants to check whether\nthe defective % varies by centre. Please analyze the\ndata at 5% si gnificance level and help the manager\ndraw appropriate inferences'

```
In [27]: #Reading data
data4 = pd.read_csv(r'C:\Users\Hypothesis Assignment\Costomer+OrderForm.csv')
data4.head()
```

```
Out[27]:

Phillippines Indonesia Malta India

0 Error Free Error Free Defective Error Free

1 Error Free Error Free Error Free Defective

2 Error Free Defective Defective Error Free

3 Error Free Error Free Error Free Error Free
```

Error Free Error Free Defective Error Free

```
In [28]: country1 = data4['Phillippines']
country2 = data4['Indonesia']
country3 = data4['Malta']
country4 = data4['India']
```

```
In [29]: data4.replace({'Error Free': 1, 'Defective': 0}, inplace=True)
data4
```

Out[29]:	Phillippines	Indonesia	Malta	India
0	1	1	0	1
1	1	1	1	0
2	1	0	0	1
3	1	1	1	1
4	1	1	0	1
295	1	1	1	1
296	1	1	1	1
297	1	1	0	1
298	1	1	1	1
299	1	0	0	1

300 rows × 4 columns

```
In [30]: #Defining Significance Level
alpha = 0.05
#Performing anova test
f_stat, p_value = stats.f_oneway(country1, country2, country3, country4)
```

C:\Users\HOME\AppData\Local\Temp\ipykernel_8148\3786107032.py:4: DeprecationWarning: Please use `f_oneway` f
rom the `scipy.stats` namespace, the `scipy.stats.stats` namespace is deprecated.
 f stat, p value = stats.f oneway(country1, country2, country3, country4)

```
In [31]: f_stat
```

Out[31]: 1.286168556089167

```
In [32]: p_value
Out[32]: 0.2776780955705948
```

```
In [33]: # Deciding Correct hypothesis

if p_value < alpha:
    print("Rejecting null hypothesis as there is a significant difference in the diameter of cutlets between else:
    print("Fail to reject null hypothesis there is no significant difference in the diameter of cutlets between else:</pre>
```

Fail to reject null hypothesis there is no significant difference in the diameter of cutlets between the two units.