import statsmodels.api as sm import statsmodels.stats.multicomp from statsmodels.formula.api import ols from statsmodels.stats.anova import anova_lm import matplotlib.pyplot as plt from scipy import stats import seaborn as sns In [2]: #Read CSV LabTAT=pd.read_csv("C:/Users/rjas/Downloads/ds assignments/3a/LabTAT.csv") In [4]: LabTAT Out[4]: Laboratory 1 Laboratory 2 Laboratory 3 Laboratory 4 0 185.35 165.53 176.70 166.13 170.49 198.45 185.91 160.79 1 2 192.77 194.92 201.23 185.18 3 199.61 177.33 183.00 176.42 4 193.41 169.57 204.63 152.60 115 178.49 170.66 193.80 172.68 215.25 176.08 183.98 177.64 116 117 202.48 174.54 203.99 170.27 118 182.40 197.18 194.52 150.87 119 182.09 215.17 221.49 162.21 120 rows × 4 columns In [5]: LabTAT.dtypes Out[5]: Laboratory 1 float64 float64 Laboratory 2 Laboratory 3 float64 Laboratory 4 float64 dtype: object In [6]: LabTAT.size Out[6]: 480 In [7]: LabTAT. shape Out[7]: (120, 4) In [8]: LabTAT.columns Out[8]: Index(['Laboratory 1', 'Laboratory 2', 'Laboratory 3', 'Laboratory 4'], dtype='object') In [9]: LabTAT.axes Out[9]: [RangeIndex(start=0, stop=120, step=1), Index(['Laboratory 1', 'Laboratory 2', 'Laboratory 3', 'Laboratory 4'], dtype='object')] In [10]: LabTAT.ndim Out[10]: 2 In [11]: LabTAT.values Out[11]: array([[185.35, 165.53, 176.7 , 166.13], [170.49, 185.91, 198.45, 160.79], [192.77, 194.92, 201.23, 185.18], [177.33, 183. , 199.61, 176.42], [193.41, 169.57, 204.63, 152.6], [179.45, 197. , 181.51, 161.12], [191.37, 166.36, 214.21, 154.02], [166.81, 169.6 , 183.43, 163.25], [158.81, 175.36, 191.6 , 152.79], [165.88, 198.68, 208.43, 161.98], [174.75, 189.12, 231. , 171.22], [193.37, 140.55, 198.37, 183.67], [184.75, 160.44, 226.62, 142.95], [178.54, 167.03, 214.44, 152.37], [180.19, 182.67, 159.69, 163.81], [172.17, 155.73, 214.14, 156.06], [172. , 183.07, 212.29, 176.44], [184.92, 177.7, 209.25, 173.68], [187.38, 191.62, 198.54, 161.49], [194.17, 186.85, 229.02, 175.05], [192.03, 161.09, 214.71, 150.22], [193.77, 173.01, 172.27, 148.17], [178.24, 199.43, 205.31, 166.21], [171.41, 185.67, 180.35, 145.4], [178.3 , 182.69, 214.58, 166.42], [176.13, 180.45, 231.76, 132.21], [180.67, 182.37, 215.78, 143.26], [177.38, 189.99, 201.78, 154.06], [190.36, 167.55, 191.64, 179.08], [199.99, 190.09, 217.56, 144.82], [181.36, 168.01, 193.04, 184.64], [187.37, 176.5 , 217.66, 188.49], [172.93, 173.54, 198.68, 198.69], [187.18, 153.75, 206.49, 142.43], [179.03, 212.8 , 198.81, 169.68], [181.69, 179.32, 200.58, 174.79], [185.28, 173.56, 199.91, 133.31], [157.14, 217.86, 190.34, 176.42], [184.42, 187.57, 181.87, 176.67], [183.45, 163.69, 214.39, 158.2], [175.5 , 186.49, 189.61, 157.02], [169.55, 171.31, 203.58, 154.55], [166.86, 202.29, 196.64, 157.58], [166.33, 197.68, 225.42, 161.96], [173.88, 156.89, 175.13, 168.14], [189.84, 168.03, 228.21, 180.14], [184.88, 173.09, 229.33, 170.28], [188.63, 187.52, 170.58, 166.68], [216.39, 193.28, 177.76, 161.42], [180.57, 178.04, 207.34, 153.97], [177.54, 197.45, 178.07, 167.5], [182.34, 172.35, 188.32, 165.43], [187.35, 167.63, 193.26, 150.83], [185.87, 194.24, 201.17, 149.96], [188.01, 179.43, 192.75, 161.94], [182.62, 183.62, 192.81, 146.46], [179.08, 157.92, 183.1 , 174.83], [189.96, 173.55, 181.92, 181.66], [169.15, 180.86, 209.43, 164.82], [169.83, 161.02, 180.11, 181.35], [181.45, 178.42, 172.92, 176.21], [196.13, 165.12, 205.01, 152.4], [172.02, 183.75, 206.46, 159.63], [181.83, 186.77, 192.48, 156.56], [170.41, 177.59, 237.49, 168.11], [161.99, 202.78, 173.08, 145.07], [138.3 , 176.63, 193.08, 128.93], [170.61, 169.11, 203.24, 165.95] [201.27, 148.7 , 215.17, 181.76], [202.12, 154.19, 186.83, 168.17], [172.62, 189.65, 199.7, 162.11], [159.13, 186.93, 225.14, 177.03], [143.52, 168.2 , 213.18, 124.06], [195.85, 159.14, 237.23, 138.48], [186.22, 182.17, 189.98, 167.83], [168.62, 198.69, 193.63, 165.07], [164.88, 167.9 , 184.7 , 153.74], [155.29, 178.21, 175.87, 172.72], [168.39, 170.73, 180.94, 159.97], [190.02, 199.62, 211.19, 167.26], [188.99, 186.03, 205.54, 164.98], [175.49, 156.23, 219.55, 154.06], [190.88, 158.08, 197.33, 168.92], [178.08, 163.74, 205.62, 168.9], [210.38, 161.45, 238.7 , 157.48], [164.27, 163.45, 192.5 , 164.03], [166.97, 178.17, 208.35, 162.76], [176.16, 194.1 , 208.04, 173.37], [199.72, 182.6 , 222.47, 152.13], [176.93, 168.51, 207.6 , 187.08], [185.16, 191.89, 194.59, 162.84], [161.05, 180.53, 184.28, 162.67], [186.32, 165.45, 174.55, 157.21], [149.24, 168.8 , 188.53, 156.79], [168.62, 159.92, 204.28, 172.38], [171.64, 188.49, 202.95, 167.62], [185.42, 205.27, 177.62, 172.72], [162.43, 185.63, 201.99, 167.55], [178.81, 197.38, 195.6 , 132.49], [172.65, 186.71, 178.86, 166.18], [164.65, 194.81, 211.23, 176.79], [178.52, 198.09, 213.29, 197.98], [170.11, 164.07, 199.12, 184.78], [167.15, 170.48, 187.97, 180.51], [190.32, 184.86, 180.93, 167.27], [167.25, 189.11, 213.22, 159.34], [179.2 , 179.68, 175.36, 142.35], [171.65, 142.38, 187.6 , 167.07], [166.35, 167.72, 208.33, 126.94], [175.41, 199.9 , 172.86, 205.18], [172.05, 174.43, 211.64, 152.31], [181.88, 168.2 , 183.19, 191.26], [201.11, 195.07, 203.05, 160.98], [154.21, 193.79, 197.88, 192.23], [145.89, 175.16, 203.39, 139.34], [178.49, 170.66, 193.8 , 172.68], [176.08, 183.98, 215.25, 177.64], [202.48, 174.54, 203.99, 170.27], [182.4 , 197.18, 194.52, 150.87], [182.09, 215.17, 221.49, 162.21]]) In [12]: LabTAT['Laboratory 1'].value_counts() Out[12]: 168.62 176.16 1 181.69 1 185.87 1 188.99 1 192.03 1 180.19 193.41 1 169.83 1 199.72 1 Name: Laboratory 1, Length: 119, dtype: int64 In [13]: LabTAT.describe() Out[13]: Laboratory 1 Laboratory 2 Laboratory 3 Laboratory 4 120.00000 count 120.000000 120.000000 120.000000 178.361583 178.902917 199.913250 163.68275 mean 13.173594 14.957114 16.539033 15.08508 std 138.300000 140.550000 159.690000 124.06000 min **25**% 170.335000 168.025000 188.232500 154.05000 **50**% 178.530000 178.870000 199.805000 164.42500 **75**% 186.535000 189.112500 211.332500 172.88250 216.390000 217.860000 238.700000 205.18000 max In [14]: LabTAT.mean() Out[14]: Laboratory 1 178.361583 Laboratory 2 178.902917 Laboratory 3 199.913250 Laboratory 4 163.682750 dtype: float64 In [16]: LabTAT.std() Laboratory 1 13.173594 Out[16]: Laboratory 2 14.957114 Laboratory 3 16.539033 Laboratory 4 15.085080 dtype: float64 In [17]: LabTAT.median() Out[17]: Laboratory 1 178.530 Laboratory 2 178.870 199.805 Laboratory 3 Laboratory 4 164.425 dtype: float64 In [18]: plt.hist(LabTAT["Laboratory 1"], facecolor="orange", edgecolor="black", bins=10) Out[18]: (array([3., 1., 6., 17., 26., 30., 23., 8., 4., 2.]), array([138.3 , 146.109, 153.918, 161.727, 169.536, 177.345, 185.154, 192.963, 200.772, 208.581, 216.39]), <BarContainer object of 10 artists>) 30 25 20 15 10 180 190 In [29]: LabTAT_CI_1=stats.norm.interval(0.975,loc= 178.361583,scale=13.173594) In [30]: LabTAT_CI_1 Out[30]: (148.83425347603986, 207.88891252396016) In [23]: plt.hist(LabTAT["Laboratory 2"], facecolor="green", edgecolor="red", bins=10) out[23]: (array([2., 4., 13., 23., 18., 26., 14., 15., 2., 3.]), array([140.55 , 148.281, 156.012, 163.743, 171.474, 179.205, 186.936, 194.667, 202.398, 210.129, 217.86]), <BarContainer object of 10 artists>) 25 20 15 10 160 170 180 190 200 210 220 LabTAT_CI_2=stats.norm.interval(0.975,loc=178.902917,scale=14.957114) In [28]: LabTAT_CI_2 Out[28]: (145.3780008833019, 212.42783311669814) In [31]: plt.hist(LabTAT["Laboratory 3"], facecolor="blue", edgecolor="green", bins=10) Out[31]: (array([1., 8., 15., 11., 23., 22., 21., 8., 6., 5.]), array([159.69 , 167.591, 175.492, 183.393, 191.294, 199.195, 207.096, 214.997, 222.898, 230.799, 238.7]), <BarContainer object of 10 artists>) 20 15 10 5 160 170 180 190 200 210 220 230 In [32]: LabTAT_CI_3=stats.norm.interval(0.975,loc=199.913250,scale=16.539033) In [33]: LabTAT_CI_3 Out[33]: (162.8426163218518, 236.98388367814823) In [34]: plt.hist(LabTAT["Laboratory 4"], facecolor="yellow", edgecolor="green", bins=10) Out[34]: (array([3., 5., 9., 17., 26., 30., 16., 9., 2., 3.]), array([124.06 , 132.172, 140.284, 148.396, 156.508, 164.62 , 172.732, 180.844, 188.956, 197.068, 205.18]), <BarContainer object of 10 artists>) 30 25 20 15 10 0 140 150 160 170 180 130 190 200 In [35]: LabTAT CI 4=stats.norm.interval(0.975,loc=163.682750,scale=15.085080) In [36]: LabTAT_CI_4 Out[36]: (129.8710105418612, 197.49448945813884) In [38]: Laboratory_1=LabTAT['Laboratory 1'].mean() Laboratory_2=LabTAT['Laboratory 2'].mean() Laboratory_3=LabTAT['Laboratory 3'].mean() Laboratory_4=LabTAT['Laboratory 4'].mean() print('Laboratory 1 Mean = ', Laboratory_1, '\nLaboratory 2 Mean = ', Laboratory_2, '\nLaboratory 3 Mean = ', Laboratory_3, '\nLaboratory 4 Mean = ', L print('Laboratory 1 Mean < Laboratory 2 Mean < Laboratory 3 Mean > Laboratory 4 Mean = ',Laboratory_1<Laboratory_2<Laboratory_3>Laboratory_4) Laboratory 1 Mean = 178.36158333333333Laboratory 2 Mean = 178.902916666668Laboratory 3 Mean = 199.91325000000003Laboratory 1 Mean < Laboratory 2 Mean < Laboratory 3 Mean > Laboratory 4 Mean = True In [39]: sns.distplot(LabTAT['Laboratory 1']) sns.distplot(LabTAT['Laboratory 2']) sns.distplot(LabTAT['Laboratory 3']) sns.distplot(LabTAT['Laboratory 4']) plt.legend(['Laboratory 1', 'Laboratory 2', 'Laboratory 3', 'Laboratoratory']) C:\Users\rjas\anaconda3\lib\site-packages\seaborn\distributions.py:2557: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level fu nction for histograms). warnings.warn(msg, FutureWarning) C:\Users\rjas\anaconda3\lib\site-packages\seaborn\distributions.py:2557: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level fu nction for histograms). warnings.warn(msg, FutureWarning) C:\Users\rjas\anaconda3\lib\site-packages\seaborn\distributions.py:2557: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level fu nction for histograms). warnings.warn(msg, FutureWarning) C:\Users\rjas\anaconda3\lib\site-packages\seaborn\distributions.py:2557: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level fu nction for histograms). warnings.warn(msg, FutureWarning) Out[39]: <matplotlib.legend.Legend at 0xc0ae310> 0.040 Laboratory 1 Laboratory 2 0.035 Laboratory 3 Laboratoratory 0.030 0.025 0.020 0.015 0.010 0.005 0.000 120 140 160 240 100 180 200 220 260 Laboratory 4 In [40]: sns.boxplot(data=[LabTAT['Laboratory 1'], LabTAT['Laboratory 2'], LabTAT['Laboratory 3'], LabTAT['Laboratory 4']], notch=True) plt.legend(['Laboratory 1','Laboratory 2','Laboratory 3','Laboratory 4']) Out[40]: <matplotlib.legend.Legend at 0xc247b50> 240 Laboratory 1 Laboratory 2 220 Laboratory 3 Laboratory 4 200 180 160 140 120 In [14]: alpha=0.05Laboratory1=pd.DataFrame(LabTAT['Laboratory 1']) Laboratory2=pd.DataFrame(LabTAT['Laboratory 2']) Laboratory3=pd.DataFrame(LabTAT['Laboratory 3']) Laboratory4=pd.DataFrame(LabTAT['Laboratory 4']) print(Laboratory1, Laboratory2, Laboratory3, Laboratory4) Laboratory 1 0 185.35 1 170.49 2 192.77 3 177.33 4 193.41 115 178.49 116 176.08 202.48 117 118 182.40 119 182.09 [120 rows x 1 columns] Laboratory 2 0 165.53 1 185.91 2 194.92 183.00 3 4 169.57 170.66 115 183.98 116 117 174.54 118 197.18 119 215.17 [120 rows x 1 columns] Laboratory 3 176.70 1 198.45 2 201.23 3 199.61 4 204.63 193.80 115 116 215.25 117 203.99 118 194.52 119 221.49 [120 rows x 1 columns] Laboratory 4 166.13 0 160.79 1 2 185.18 3 176.42 4 152.60 115 172.68 116 177.64 117 170.27 150.87 118 119 162.21 [120 rows x 1 columns] In [6]: df = LabTATIn [8]: stats.f_oneway(df.iloc[:,0],df.iloc[:,1],df.iloc[:,2],df.iloc[:,3]) Out[8]: F_onewayResult(statistic=118.70421654401437, pvalue=2.1156708949992414e-57) # Anova Test-One way H0:Average of all laboratory are same Ha:Average of atleast 1 laboratory are different # P-value is 0.00 < 0.05 = Accept Ha, hence Average of atleast 1 laboratory are different As per results we can say that these are not equal i.e Average of atleast 1 laboratory are different

In [1]:

import pandas as pd
import numpy as np

import os
import random

import scipy.stats as stats