Statistics module assignment

PART A:

Task 1: Conditional Probability

$$P(B \mid A) = \frac{P(B \cap A)}{P(A)} = \frac{P(A \cap B)}{P(A)}, \quad P(A) > 0$$

P(S= true | G=female,C=1)

91 / 887 were females in 1st class who survived

94 / 887 were females in 1st class

$$\frac{91}{887}$$
 = 0.102 $\frac{94}{887}$ = 0.105

$$P(S = true \mid G = female, C = 1) = \frac{0.102}{0.105} = 0.971 = 97.1\%$$

P(S= true | G=female,C=2)

76 out of 887 are females who were in 2nd class

70 out of 887 are females who were in 2nd class and survived

$$\frac{70}{887}$$
 = 0.78 $\frac{76}{887}$ = 0.86

$$P(S = true \mid G = female, C = 2) = \frac{0.078}{0.086} = 0.906 = 90.6\%$$

P(S= true | G=female,C=3)

72 / 887 female in 3nd class and survived

144 / 887 female in 3rd class

$$\frac{72}{887}$$
 = 0.081 $\frac{144}{887}$ = 0.162

$$P(S = true \mid G = female, C = 3) = \frac{0.081}{0.162} = 0.5 = 50\%$$

P(S= true | G=male,C=1)

45 / 887 are males who are in 1st class and survived

122 / 887 are males who were in 1st class

$$\frac{45}{887}$$
 = 0.050 $\frac{122}{887}$ = 0.137

$$P(S = true \mid G = male, C = 1) = \frac{0.050}{0.137} = 0.364 = 36.4\%$$

P(S= true | G=male,C=2)

108 / 887 are males who were in 2nd class

17 / 887 were males in 2nd class who survived

$$\frac{17}{887}$$
 = 0.019 $\frac{108}{887}$ = 0.121

$$P(S = true \mid G = male, C = 2) = \frac{0.019}{0.121} = 0.157 = 16\%$$

P(S= true | G=male,C=3)

47 / 887 were males in 3rd class who survived 343 / 887 were males in 3rd class

$$\frac{47}{887}$$
 = 0.052 $\frac{343}{887}$ = 0.386

$$P(S = true \mid G = male, C = 3) = \frac{0.052}{0.386} = 0.134 = 13.4\%$$

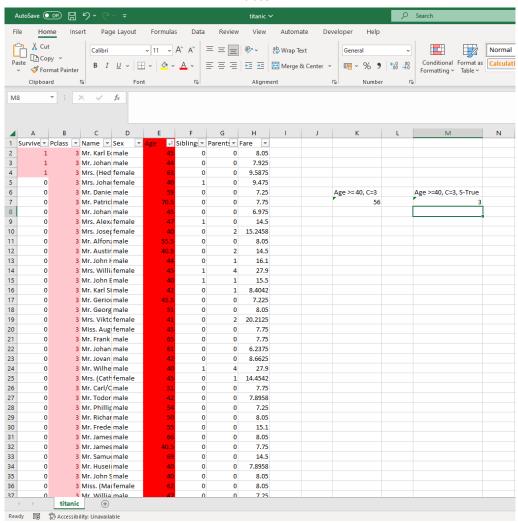
Task b:

$$P(S = True | Age >= 40, C = 3)$$

56 / 887 are 3rd class and have an age of 40 or over 3 / 887 are 3rd class, have an age of 40 or over and survived

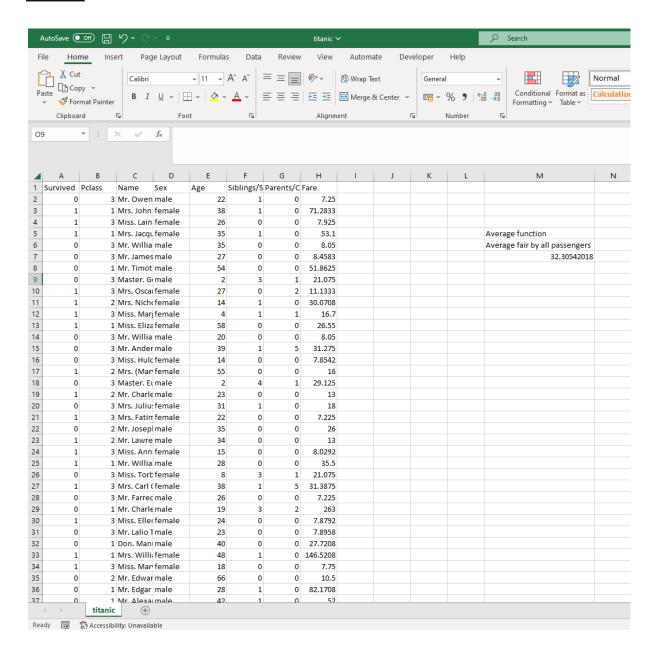
$$\frac{56}{887} = 0.063 \frac{3}{887} = 0.003$$

$$P(S = true \mid Age >= 40, C = 3) = \frac{0.003}{0.063} = 0.047 = 5\%$$



I used Conditional formatting and highlights in Excel so that i could filter through the data and this would allow me to use =COUNT functions to count the cells that included the data i needed which i could then use later on in my calculations, this was done as counting by hand will present the opportunity for human error which will result in incorrect calculations.

Task c:



I simply opened the dataset in Excel and used the =AVG function on all the cells in the "Fare" column as this would efficiently calculate the average fare for me and because this included all passengers i could just highlight all the cells in the column and include them in the calculation. - Average fair paid by all passengers =

32.30542018 which simplifies to **32.3** (There was no information on currency given so i have not used units)

Average fare = 32.3

Task 2: Hypotheses Testing

Survived passengers & class of travel

Chi-Square Calculator

Success! The contingency table below provides the following information: the observed cell totals, (the expected cell totals) and [the chi-square statistic for each cell].

The chi-square statistic, *p*-value and statement of significance appear beneath the table. Blue means you're dealing with dependent variables; red, independent.

			Results		
	1st Class	2nd Class	3rd Class		Row Totals
Male	45 (43.35) [0.06]	17 (27.73) [4.15]	47 (37.93) [2.17]		109
Female	91 (92.65) [0.03]	70 (59.27) [1.94]	72 (81.07) [1.02]		233
Column Totals	136	87	119		342 (Grand Total)

The chi-square statistic is 9.3711. The p-value is .009227. The result is significant at p < .05.

```
from scipy.stats import chi2_contingency
3
       data = [[45, 17, 47], [91, 70, 72]]
5
      stat, p, dof, expected = chi2_contingency(data)
      print("The expected values are : ", expected)
print("The chi-squared value is ", stat)
print("The degree of freedom is ", dof)
 8
9
10
      print("The p value is ", p)
       significance_level = 0.05
14
      if p <= significance_level :</pre>
15
           print('Reject HO (There IS an association)')
16
           print('Accept HO (There IS NOT an association)')
18
      The expected values are : [[43.34502924 27.72807018 37.92690058]
       [92.65497076 59.27192982 81.07309942]]
       The chi-squred value is 9.371139335460681
      The degree of freedom is 2
      The p value is 0.00922747651908542
       Reject HO (There IS an association)
```

```
Degree of freedom = (Rows - 1) (Columns - 1)
Degree of freedom = 1 X 2 = 2
```

The Image above tells us that the Chi-square statistic is 9.3711 which is larger than 5.99, this is the number we find on the Chi squared distribution table when looking at a degree of freedom equivalent to 2 and a p value of 0.05. This means that we reject the null hypothesis which is that there is no relationship between the two variables and therefore we accept the alternative hypothesis which is that there is a relationship between the surviving passengers and the class of travel. This outcome suggests that passengers who boarded the Titanic at a higher class(e.g 1st class) were at a much lower risk of death and were subsequently more safe than the passengers in the lower classes, this makes sense as the 1st and 2nd class cabins were located closer to the boat deck(where the lifeboats were located) therefore making access to lifeboats much easier to 1st and 2nd class passengers than 3rd class passengers.

Survived passengers & gender

Chi-Square Calculator

Success! The contingency table below provides the following information: the observed cell totals, (the expected cell totals) and [the chi-square statistic for each cell].

The chi-square statistic, p-value and statement of significance appear beneath the table. Blue means you're dealing with dependent variables; red, independent.

			Results		
	Male	Female			Row Totals
Survived	109 (220.93) [56.71]	233 (121.07) [103.48]			342
Dead	464 (352.07) [35.59]	81 (192.93) [64.94]			545
Column Totals	573	314			887 (Grand Total)

The chi-square statistic is 260.7153. The p-value is < .00001. The result is significant at p < .05.

```
[6]
      from scipy.stats import chi2_contingency
      data = [[109, 233], [464, 81]]
      stat, p, dof, expected = chi2_contingency(data)
      print("The expected values are : ", expected)
     print("The chi-squared value is ", stat)
print("The degree of freedom is ", dof)
8
10
     print("The p value is ", p)
      significance_level = 0.05
14
     if p <= significance_level :</pre>
15
         print('Reject HO (There IS an association)')
16
          print('Accept HO (There IS NOT an association)')
18
      The expected values are : [[220.93122886 121.06877114]
      [352.06877114 192.93122886]]
      The chi-squared value is 258.39126076789773
      The degree of freedom is 1
      The p value is 3.847574039733855e-58
      Reject HO (There IS an association)
```

Degree of freedom = (Rows - 1) (Columns - 1) Degree of freedom = 1 X 1 = 1

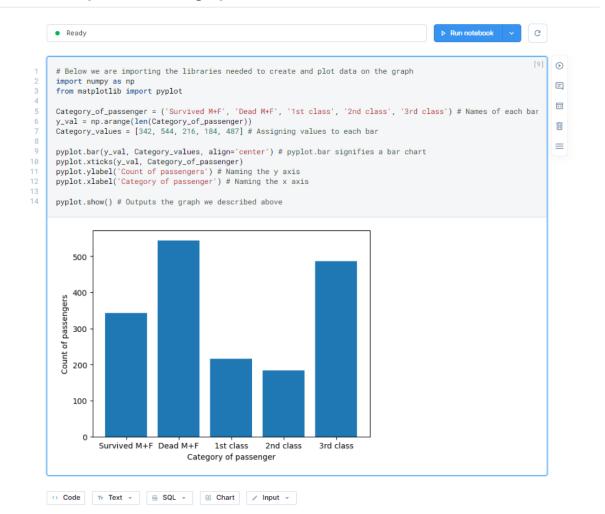
D.f = 1 / P value = 0.05 = 3.84 3.84 < 260.7153

The Image above tells us that the Chi-square statistic is 260.7153 which is larger than 3.84, this is the number we find on the Chi squared distribution table when looking at a degree of freedom equivalent to 1 and a p value of 0.05. This means that we reject the null hypothesis which is that there is no relationship between the passengers that survived and their gender, instead we accept the alternative hypothesis which is that there is a relationship between the passengers that survived and their gender. This outcome suggests that you were more or less likely to survive based on your gender and this could make sense as it is believed that the women

and children were given first priority when boarding the lifeboats, when we compile the data, more women survived than men in every class which further backs up this claim.

Task 2: Visualisation

Create multiple individual graphs



This graph shows all the categories and their corresponding values in one bar graph



This graph shows the number of males and females that survived

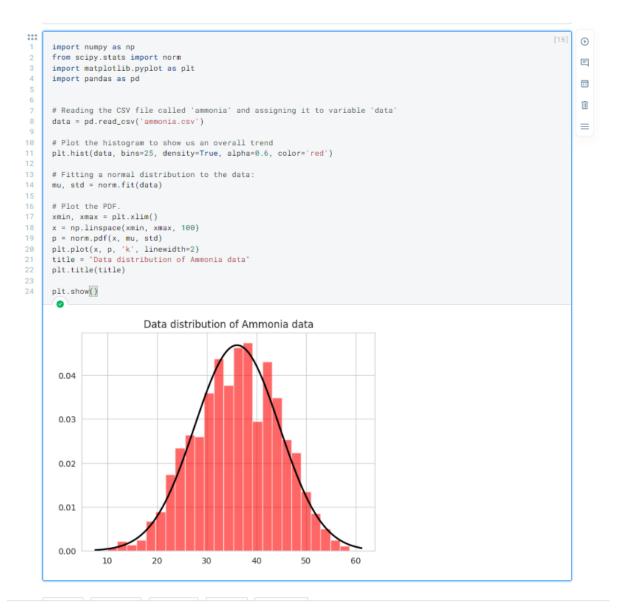


This graph shows the number of passengers in each class



This graph shows the number of males and females that died

PART B: Data distribution



The image above is the coding i have created on Deep note, i have also included the output which shows a histogram of the ammonia data from 'ammonia.csv', from the shape of the bars in the histogram i can infer that there is a normal distribution, i then added some code to overlay a bell curve to further make my observation more apparent.

To answer the questions:

- I believe a normal distribution is the most likely distribution of the dataset.
- I have estimated the mean to be 36.1 and the standard deviation to be 8.51 assuming that the data is from a normal distribution.

The probability of the ammonia concentration being greater than 30 and less than 40 mg/L is 44%.

$$25corc = (X - \mu)$$
> 30 and < 40
$$\frac{40 - 36.094993}{8.515969} = 0.45855111$$

$$\frac{30 - 36.094993}{8.515969} = -0.7157134$$

$$8.515969$$

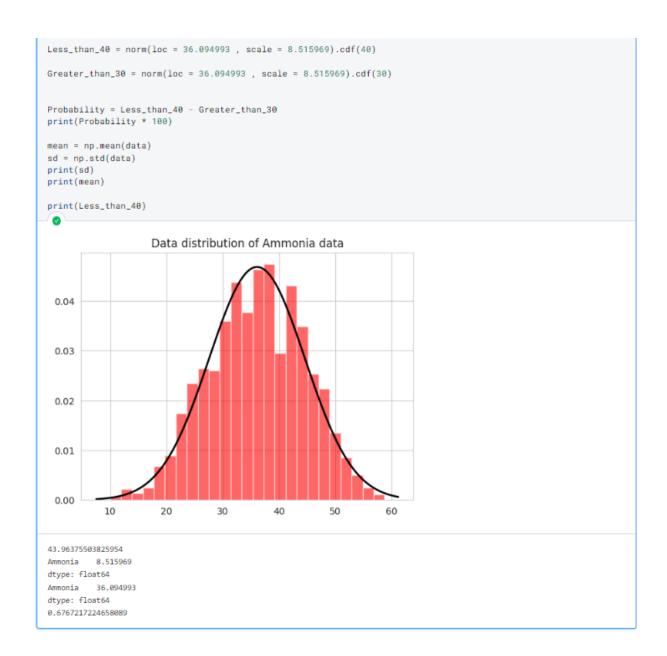
$$0.45855111 = 0.6736$$

$$-0.7157134 = 0.2358$$

$$0.6736 - 0.2358 = 0.4378$$

$$= 44\%$$

Above is the manual calculation for the probability of the ammonia concentration being greater than 30 and less than 40 mg/L. Below is the python calculation of the probability and this also shows 43.96 % which rounded up is 44%



PART C: Combination probability

<u>APPENDIX</u>

Graph showing all categories

```
# Below we are importing the libraries needed to create and plot data on
the graph
import numpy as np
from matplotlib import pyplot

Category_of_passenger = ('Survived M+F', 'Dead M+F', '1st class', '2nd
class', '3rd class') # Names of each bar
y_val = np.arange(len(Category_of_passenger))
Category_values = [342, 544, 216, 184, 487] # Assigning values to each bar
```

```
pyplot.bar(y_val, Category_values, align='center') # pyplot.bar signifies a
bar chart
pyplot.xticks(y_val, Category_of_passenger)
pyplot.ylabel('Count of passengers') # Naming the y axis
pyplot.xlabel('Category of passenger') # Naming the x axis
pyplot.show() # Outputs the graph we described above
Male and females that survived
# Below we are importing the libraries needed to create and plot data on
the graph
import numpy as np
from matplotlib import pyplot
Category_of_passenger = ('Male', 'Female') # Names of each bar
y_val = np.arange(len(Category_of_passenger))
Category_values = [109, 233] # Assigning values to each bar
pyplot.bar(y_val, Category_values, align='center') # pyplot.bar signifies a
bar chart
pyplot.xticks(y_val, Category_of_passenger)
pyplot.ylabel('Count of passengers that survived') # Naming the y axis
pyplot.xlabel('Gender') # Naming the x axis
pyplot.show() # Outputs the graph we described above
Male and females that died
# Below we are importing the libraries needed to create and plot data on
the graph
import numpy as np
from matplotlib import pyplot
Category_of_passenger = ('Male', 'Female') # Names of each bar
y_val = np.arange(len(Category_of_passenger))
```

Category_values = [463, 81] # Assigning values to each bar

```
pyplot.bar(y_val, Category_values, align='center') # pyplot.bar signifies a
bar chart
pyplot.xticks(y_val, Category_of_passenger)
pyplot.ylabel('Count of passengers that died') # Naming the y axis
pyplot.xlabel('Gender') # Naming the x axis

pyplot.show() # Outputs the graph we described above
```

```
Count of passengers and what class they were in
```

```
# Below we are importing the libraries needed to create and plot data on
the graph
import numpy as np
from matplotlib import pyplot

Category_of_passenger = ('1st Class', '2nd class', '3rd Class') # Names of
each bar
y_val = np.arange(len(Category_of_passenger))
Category_values = [216, 190, 486] # Assigning values to each bar

pyplot.bar(y_val, Category_values, align='center') # pyplot.bar signifies a
bar chart
pyplot.xticks(y_val, Category_of_passenger)
pyplot.ylabel('Count of passengers') # Naming the y axis
pyplot.xlabel('Class') # Naming the x axis

pyplot.show() # Outputs the graph we described above
```

Data distribution graph

```
import numpy as np
from scipy.stats import norm
import matplotlib.pyplot as plt
import pandas as pd
```

```
# Reading the CSV file called 'ammonia' and assigning it to variable 'data'
data = pd.read_csv('ammonia.csv')
# Plot the histogram to show us an overall trend
plt.hist(data, bins=25, density=True, alpha=0.6, color='red')
# Fitting a normal distribution to the data:
mu, std = norm.fit(data)
# Plot the PDF.
xmin, xmax = plt.xlim()
x = np.linspace(xmin, xmax, 100)
p = norm.pdf(x, mu, std)
plt.plot(x, p, 'k', linewidth=2)
title = "Data distribution of Ammonia data"
plt.title(title)
plt.show()
\#Less\_than\_40 = norm(loc = 36.094993 , scale = 8.515969).cdf(40)
\#Greater\_than\_30 = norm(loc = 36.094993 , scale = 8.515969).cdf(30)
#Probability = Less_than_40 - Greater_than_30
#print(Probability * 100)
#mean = np.mean(data)
#sd = np.std(data)
#print(sd)
#print(mean)
#print(Less_than_40)
Chi squared test #1
from scipy.stats import chi2_contingency
data = [[45, 17, 47], [91, 70, 72]]
```

```
from scipy.stats import chi2_contingency
data = [[45, 17, 47], [91, 70, 72]]
stat, p, dof, expected = chi2_contingency(data)
print("The expected values are : ", expected)
print("The chi-squared value is ", stat)
print("The degree of freedom is ", dof)
print("The p value is ", p)
significance_level = 0.05
if p <= significance_level :</pre>
    print('Reject HO (There IS an association)')
else:
    print('Accept HO (There IS NOT an association)')
Chi squared test #2
from scipy.stats import chi2_contingency
data = [[109, 233], [464, 81]]
stat, p, dof, expected = chi2_contingency(data)
print("The expected values are : ", expected)
print("The chi-squared value is ", stat)
print("The degree of freedom is ", dof)
print("The p value is ", p)
significance_level = 0.05
if p <= significance_level :</pre>
    print('Reject HO (There IS an association)')
else:
    print('Accept HO (There IS NOT an association)')
```