TP1- ROUSSEAU Data Analysis

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1 Data Analysis - TP 1

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This document has been done using python on Jupyter Notebook with the librairies:

- Numpy to manipulate arrays
- matplotlib to plot graphics
- pandas to import csv
- scipy for mathematicals usage
- maths for sqrt, pi, exp

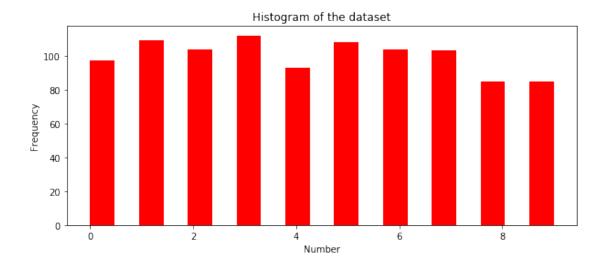
2 Exercice A : Discrete series

```
In [34]: #Import of libraries

    import matplotlib as plt
    import numpy as np
    import matplotlib.pyplot as plt
    import pandas as pd
    from scipy import stats
    from math import sqrt,pi,exp

In [35]: #We Generate a random series of 1000 element and we plot them with matplotlib

A = np.random.randint(0,10,1000)
    figure,axe = plt.subplots(figsize=(10,4))
    plt.hist(A,bins=19,color='red')
    axe.set_title("Histogram of the dataset")
    axe.set_xlabel("Number")
    axe.set_ylabel("Frequency")
Out[35]: Text(0,0.5, 'Frequency')
```



```
In [36]: #Compute the mean/max/mode
         #Compute the mean : the mean is the sum of all the element
         #divided by the len of the array
         mean = sum(A)/len(A)
         #Compute the median : we select the mean between the central element
         #and central element +1 and we convert it into an integer
         A_sorted = np.sort(A)
         index1 = len(A_sorted)/2
         index2=len(A_sorted)/2+1
         index_median = (index1+index2)/2
         median=A_sorted[int(index_median)]
         #Compute the mode : we use numpy.unique who return a 2D array with the number of
         #occurence of each element
         unique, counts = np.unique(A, return_counts=True)
         mode = counts.argmax()
         #Print the result
         print("The mean is {0}".format(mean))
         print("The median is {0}".format(median))
         print("The mode is {0}".format(mode))
The mean is 4.355
```

The median is 4
The mode is 3

The results obtained with the numpy functions are exactly the same as the results of the manually calculated functions. It is normal that the median and the mean are different because the mean is impacted by large and small values while the median only takes into account the number of elements.

```
In [38]: #The we compute the range, the variance and the standard deviation
         #of the same discrete series, firsyt by ourself and then with
         #the numpy functions
         #We compute the range
         range = np.max(A)-np.min(A)
         range_withlib =np.ptp(A,axis=0)
         #We compute the Variance
         num=0
         for x in A:
             num += (x-mean)*(x+mean)
         variance = num/len(A)
         variance_withlib = np.var(A)
         #We compute de standard deviation
         std_withlib = np.std(A)
         std = sqrt(variance)
         #We compare the result
         print("The difference between the range with lib and without is {0}"
               .format(range_withlib-range))
         print("The difference between the variance with lib and without is {0}"
               .format(variance_withlib-variance))
```

```
The difference between the range with lib and without is 0
The difference between the variance with lib and without is -1.163513729807164e-13
The difference between the standard deviation with lib and without is -2.042810365310288e-14
```

It can be seen that the difference between the two methods creates very small deviations for the variance and the standard variation. The standard deviation is less than 10 et close to the means which means that de series is homogeneous.

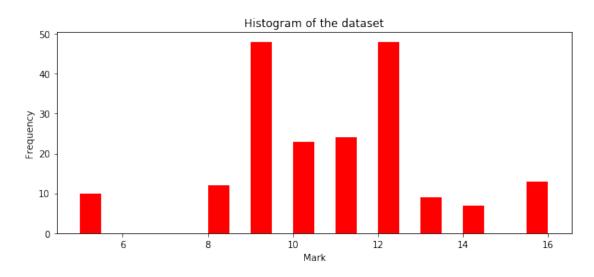
3 Exercice B : Grouped discrete series

```
In [39]: #We input the following array

Mark = [5,8,9,10,11,12,13,14,16]
   Number = [10,12,48,23,24,48,9,7,13]

#Then we plot these arrays, as an histogram
   figure,axe = plt.subplots(figsize=(10,4))
   plt.hist(Mark,bins=22,color='red',weights=Number)
   axe.set_title("Histogram of the dataset")
   axe.set_xlabel("Mark")
   axe.set_ylabel("Frequency")
```

totalmark=[a*b for a,b in zip(Mark,Number)]



```
In [40]: #We compute the mean, the median, the variance,
         #the standard deviation, the min and the max
         mean = np.sum(totalmark)/np.sum(Number)
         dispersion_mark_median = stats.binned_statistic(Number,Mark,statistic="median",bins=1)
         median = dispersion_mark_median[0][0]
         numvar = []
         i=0
         while i < len(Mark):
             numvar.append(Number[i]*(Mark[i]-mean)**2)
             i+=1
         variance = sum(numvar)/sum(Number)
         dispersion_mark_min = stats.binned_statistic(Number, Mark, statistic="min", bins=1)
         min = dispersion_mark_min[0][0]
         dispersion_mark_max = stats.binned_statistic(Number, Mark, statistic="max", bins=1)
         max = dispersion_mark_max[0][0]
         mode = Mark[np.argmax(Number)]
         sumarray = np.dot(Mark, Number)
         #We display everything
         print("Min: {0} ".format(min))
         print("Mean: {0} ".format(mean))
         print("Variance: {0} ".format(variance))
         print("Standard deviation: {0} ".format(std))
         print("Median: {0} ".format(median))
         print("Max: {0} ".format(max))
         print("Mode: {0} ".format(mode))
Min: 5.0
Mean: 10.675257731958762
Variance: 5.848150706770113
Standard deviation: 2.8055257974219585
Median: 11.0
Max: 16.0
```

D:\Programme\Anaconda\envs\ialab\lib\site-packages\scipy\stats_binned_statistic.py:607: Future result = result[core]

We notice that this distribution is bimodal because it has 2 peaks at 9 and 12. This may be related to the fact that there are two different types of populations in this data set. Those who passed the exam and those who did not.

4 Exercice C: Normal distributions

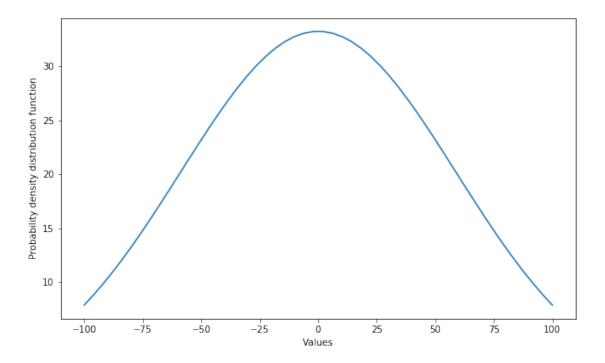
```
In [41]: #We generate a sample of n random variables that follow a normal
    #distribution of mean m and standard deviation sd

x = np.linspace(-100,100)
std = x.std()
m= x.mean()

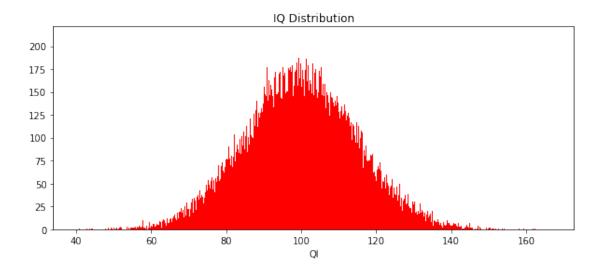
probaility_density_fct =np.exp((-0.5*(x-m)**2)/std**2)*(1/(np.sqrt(np.pi))*std)

#We plot it
fig,axe2 = plt.subplots(figsize=(10,6))
axe2.plot(x,probaility_density_fct)
axe2.set_xlabel('Values')
axe2.set_ylabel('Probability_density_distribution_function')
```

Out[41]: Text(0,0.5,'Probability density distribution function')



Out[42]: Text(0.5,0,'QI')



Mean: 99.97417399964813

Standard deviation: 14.966496430905504

Variance: 223.9960154163072

The standard deviation is very far from the mean, which means that there are many disparities among the different IQs present in the data set. The average is 100, slightly higher than the French average (98) but still low compared to Hong Kong: 108.

```
In [44]: #We create a function who return the probability for a QI to be above or
         #below a certain value
         def proba_isSup(val,isSup):
             count = 0
             for element in QI:
                 if element < val and isSup == True:
                     count+=1
                 if element > val and isSup == False:
                     count+=1
             return (count/len(QI))*100
         range_95 = stats.norm.interval(0.95,loc=mean_norm, scale=std_norm)
         print("Percentage of the sample that has an IQ bellow 60", proba_isSup(60,True))
         print("Percentage of the sample that has an IQ above 130", proba_isSup(130,False))
         print("Range of values that contains 95 percent of the sample", range_95)
Percentage of the sample that has an IQ bellow 60 0.387
Percentage of the sample that has an IQ above 130 2.33699999999997
Range of values that contains 95 percent of the sample (70.64038002032608, 129.30796797897017)
```

5 Exercice D: IQ analysis

```
In [45]: #We generate 3 different samples of size 10, 1000 and 100000
    #(set10, set100, set100000) with a mean value
    #of 100 and a standard deviation of 15

std = 15
    mean = 100

set10= np.random.normal(mean, std, 10)
    set100= np.random.normal(mean, std, 100)
    set100000= np.random.normal(mean, std, 100000)

mean_value10 = np.mean(set10)
    mean_value100 = np.mean(set100)
    mean_value100000 = np.mean(set100000)

std_value10 = np.std(set10)
    std_value100 = np.std(set100)
    std_value100000 = np.std(set100000)
```

```
print("Comparaison mean 10 :", mean_value10-mean)
         print("Comparaison mean 100 :", mean_value100-mean)
         print("Comparaison mean 100000 :", mean_value100000-mean,'\n')
         print("Comparaison std 10 :", std_value10-std)
         print("Comparaison std 100 :", std_value100-std)
         print("Comparaison std 100000 :", std_value100000-std,'\n')
         range_10 = stats.norm.interval(0.95,loc=mean_value10, scale=std_value10)
         range_100 = stats.norm.interval(0.95,loc=mean_value100, scale=std_value100)
         range_100000 = stats.norm.interval(0.95,loc=mean_value100000, scale=std_value100000)
         typerror10 = std/sqrt(10)
         typerror100 = std/sqrt(100)
         typerror100000 = std/sqrt(100000)
         print("Interval 10 :", range_10)
         print("Interval 100 :", range_100)
         print("Interval 100000 :", range_100000,'\n')
        print("typerror10 : " , typerror10)
        print("typerror100 : " , typerror100)
         print("typerror1000000 : " , typerror100000)
         a= mean_value100 + 1.96*(std_value100/100)
         print ("val = ", a)
Comparaison mean 10 : -0.8596166502354379
Comparaison mean 100 : 0.0976825460552817
Comparaison mean 100000 : -0.09479871876243351
Comparaison std 10 : 1.2904007951243806
Comparaison std 100 : 0.2940737884636775
Comparaison std 100000 : -0.01794482301835565
Interval 10: (67.21178449759812, 131.06898220193102)
Interval 100 : (70.12184874376841, 130.07351634834214)
Interval 100000 : (70.54091271996168, 129.26948984251345)
typerror10 : 4.743416490252569
typerror100 : 1.5
typerror100000 : 0.04743416490252569
val = 100.39744639230916
```

The more data there are, the more the confidence interval increases, the closer the mean is to the theoretical mean and the closer the standard deviation is to the theoretical mean. On the

other hand, with a small dataset, the errors are huge (see typerror) and the confidence interval is reduced.

```
In [27]: #We import the csv file with pandas
        import pandas as pd
         dataframe = pd.read_csv("malnutrition.csv")
        mean_malnutrition = np.mean(dataframe.values)
         std_malnutrition = np.std(dataframe.values)
        malnutrition_interval = stats.norm.interval(0.95,loc=mean_malnutrition,
                                                     scale=std_malnutrition)
In [49]: #Comparaison with set1000
        diff_mean = mean_malnutrition-mean_value100
         diff std = std malnutrition-std value100
         diff_intervalMin = diff_mean - 1.96*sqrt((std_malnutrition**2/len(dataframe))
                                                  +((std_value100**2/len(set100))))
         diff_intervalMax = diff_mean + 1.96*sqrt((std_malnutrition**2/len(dataframe))
                                                  +((std_value100**2/len(set100))))
         print("Mean difference : " ,diff_mean)
         print("Standard deviation difference : " ,diff_std)
        print("Confidence interval difference: " ,diff_intervalMin, ";"
               ,diff_intervalMax)
Mean difference: -12.208793657166396
Standard deviation difference: -5.659446750566907
Confidence interval difference: -15.75673132394913; -8.660855990383663
```

Conclusion

According to the results, people suffering from malnutrition have an IQ that is 12.5 points lower on average, which is a lot. The standard deviation is higher, which means that malnutrition has a more or less significant impact on individuals. We also note with the confidence interval that it is those with high IQ that decline more than those with low IQ already. It would be interesting to know to what extent each individual is affected by malnutrition with indicators such as their weight or their cholesterol level for example. However, it would be interesting to ask whether malnutrition leads to a decrease in IQ or whether a lower IQ increases the chances of malnutrition.