

```

import numpy as np
import matplotlib.pyplot as plt
from scipy.optimize import curve_fit
import scipy.stats
import pandas as pd

from scipy.special import factorial
# (function needed to compute factorials for an array)  ## Not anymore

def P(N,Nbar):
    lnprob=N*np.log(Nbar) - Nbar - scipy.special.gammaln(N+1)
    return np.exp(lnprob)

#Read raw trails from excel sheet (trail number vs observed counts)
filelow=pd.read_csv(r'./geiger.csv')
data=filelow["counts"]
#Check if data is loaded correctly
#print(data)

min_count = min(data)
max_count = max(data)
#print(min_count, max_count)

#Create appropriate boundaries
bin_boundaries = []
for i in range (0,(max_count-min_count+1)+1):
    bin_boundaries.append(i+(min(data)-0.5))

## Plot histogram of the distribution
hist_data = plt.hist(data,bin_boundaries,label='histogram',rwidth=0.90)

```

```

# Extract counts and occurrence from the histogram
counts1=[]
for i in range(min_count,max_count+1):
    counts1.append(i)
counts = np.array(counts1)
occurrence = hist_data[0]

# Mean calculation

m=0
for i in range(0, len(counts)):
    m= m+ counts[i]*occurrence[i]

#Mean of the distribution
mean= m/sum(occurrence)
Mean_counts= mean
#print("Mean of distribution is : " , Mean_counts)

#Scatter plot of the distribution, i.e, heights of bars on histogram (**should be available in
plt.scatter(counts, occurrence,label='observed value',color='red')

# Creating a x_values for plotting a smooth fit
l=list(range(max(min(counts)-5,0),max(counts)+5)) #(N can not be less than zero for P(N))
l1=np.array(l,dtype='double') # Changing data type to accommodate high exponentials
#print("list of x values for plotting are= ",l1)

#Genrate data poitns to make smooth curve
y_for_fit = P(l1,float(mean))
plt.plot(l1,y_for_fit*np.sum(occurrence),'k',label='fit')
#-----

#Scatter plot of expected points on the curve

```

```

expexted_y_values = P(counts,float(mean))*np.sum(occurrence)
plt.scatter(counts,expexted_y_values,color='green',label='expected values')
#-----

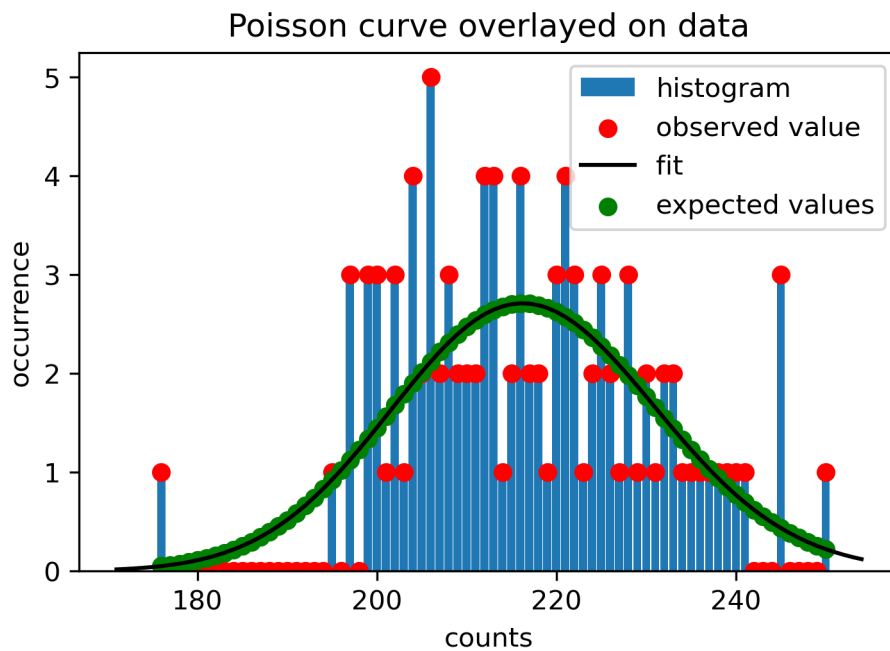
## Chi_square calculation
chi_square=0
for i in range(0,len(counts)):
    chi_square = chi_square + ((occurrence[i]-expexted_y_values[i])**2)/(expexted_y_values[i])
#-----

# Plot graph
plt.xlabel("counts")
plt.ylabel("occurrence")
plt.title("Poisson curve overlayed on data")
plt.legend()
plt.show()

# Print_results
print("Mean of distribution is : " , Mean_counts)
print("Chi square of the fit is :",chi_square)
Dof=len(counts)-1 #Degrees of freedom
print("No of degrees of freedom is :",Dof)

if (chi_square/Dof>1):
    print("Your (chi_square)/DoF is =",chi_square/Dof," , which is greater than 1 => Poisson :")
if (chi_square/Dof<=1):
    print("Your (chi_square)/DoF is  =",chi_square/Dof," , which is less than 1 => Poisson fit")

```



Mean of distribution is : 216.74

Chi square of the fit is : 71.60102781757169

No of degrees of freedom is : 74

Your (chi_square)/DoF is = 0.967581456994212 , which is less than 1 => Poisson fit overlaps