Tanmay Garg

CS20BTECH11063

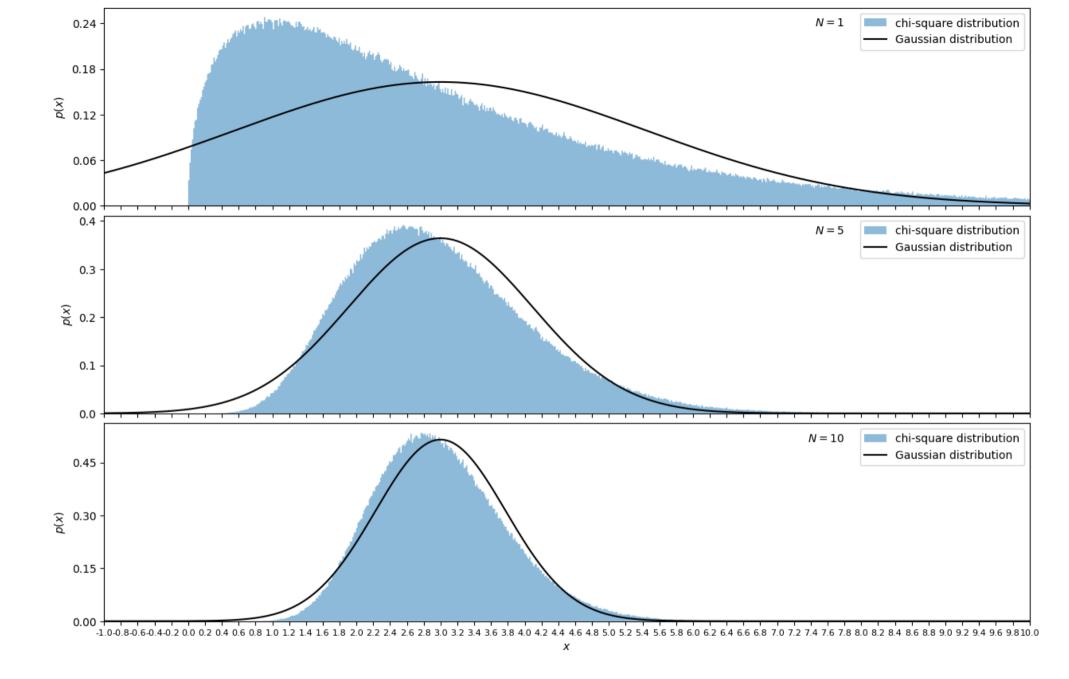
Data Science Analysis Assignment 2

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

Q1

```
In [2]: # Generate the uniform samples
        from scipy.stats import norm
        N = [1, 5, 10]
        x = np.random.chisquare(3, size=(max(N), int(1E6)))
        fig = plt.figure(figsize=(15, 10))
        fig.subplots_adjust(hspace=0.05)
        for i in range(len(N)):
            ax = fig.add_subplot(3, 1, i + 1)
            # take the mean of the first N[i] samples
            x_i = x[:N[i], :].mean(0)
            # histogram the data
            ax.hist(x i, bins=np.linspace(0, 15, 1000),
                    histtype='stepfilled', alpha=0.5, density=True, label='chi-square distribution')
            # plot the expected gaussian pdf
            sigma = (1. / np.sqrt(N[i])) * np.sqrt(6)
            # sigma = sigma * np.sqrt(6)
            dist = norm(mu, sigma)
```

```
x pdf = np.linspace(-1, 10, 1000)
   ax.plot(x pdf, dist.pdf(x pdf), label='Gaussian distribution', color='black')
   ax.set xlim(-1, 10.0)
   # ax.set ylim(0.001, None)
   ax.xaxis.set major locator(plt.MultipleLocator(0.2))
   ax.yaxis.set major locator(plt.MaxNLocator(5))
   ax.text(0.80, 0.95, r"$N = %i$" % N[i],
           ha='right', va='top', transform=ax.transAxes)
   if i == len(N) - 1:
        ax.xaxis.set major formatter(plt.FormatStrFormatter('%.1f'))
        ax.xaxis.set_tick_params(which='major', labelsize=8)
        ax.set xlabel(r'$x$')
    else:
        ax.xaxis.set_major_formatter(plt.NullFormatter())
   ax.set_ylabel('$p(x)$')
   # ax.grid()
   ax.legend()
plt.show()
```



Q2

```
In [3]: df = pd.read_csv('luminosity_redshift.csv', sep=' ')
plt.loglog(df['z'], df['Lx'], '.')
```

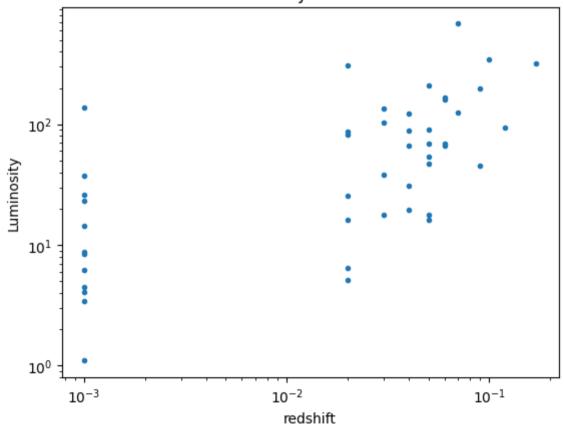
```
plt.xlabel('redshift')
plt.ylabel('Luminosity')
plt.title('Luminosity vs Redshift')
plt.show()

from scipy.stats import spearmanr, pearsonr, kendalltau

spearman_coefficient, spearman_pvalue = spearmanr(df['z'], df['Lx'])
pearson_coefficient, pearson_pvalue = pearsonr(df['z'], df['Lx'])
kendall_coefficient, kendall_pvalue = kendalltau(df['z'], df['Lx'])

print(f'Spearman Correlation Coefficient: {spearman_coefficient}, p-value: {spearman_pvalue}')
print(f'Pearson Correlation Coefficient: {pearson_coefficient}, p-value: {pearson_pvalue}')
print(f'Kendall Correlation Coefficient: {kendall_coefficient}, p-value: {kendall_pvalue}')
```

Luminosity vs Redshift

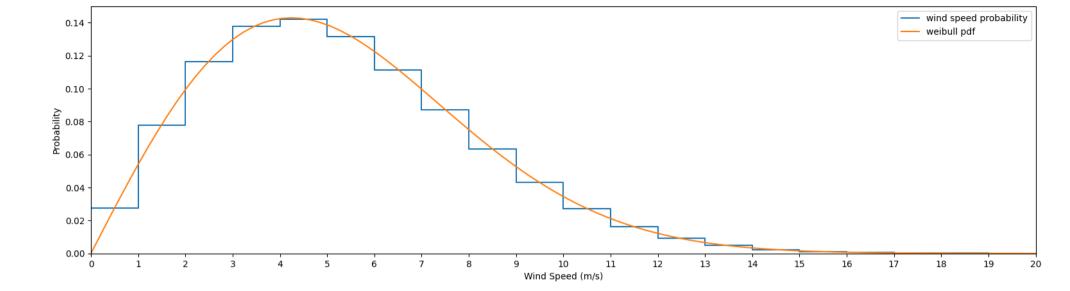


Spearman Correlation Coefficient: 0.6596325957535454, p-value: 6.166489759081011e-07 Pearson Correlation Coefficient: 0.5144497852670242, p-value: 0.0002546471657612425 Kendall Correlation Coefficient: 0.5029584682704178, p-value: 2.9696862274734036e-06

According to the above graph the data seems to have some positive correlation, which can also be verified by the correlation coefficients and p-values.

Q3

```
In [4]: from scipy.stats import weibull min, weibull max
        wind speed = pd.read csv('wind.csv', sep='\t')
        wind speed['probability'] = wind speed['probability'] / 100
        wind speed['lower bound'] = wind speed['speed'].str.extract('(\d+)').astype(int)
        wind speed['upper bound'] = wind speed['speed'].str.extract('-(\d+)').astype(int)
        # weibull distribution
        x = np.linspace(0, 20, 100)
        y = weibull min.pdf(x, 2, loc=0, scale=6)
        plt.figure(figsize=(19, 5))
        plt.step(wind_speed['lower_bound'], wind_speed['probability'], where='post', label='wind speed probability')
        plt.xlabel('Wind Speed (m/s)')
        plt.ylabel('Probability')
        plt.xticks(np.arange(0, 21, 1))
        plt.xlim(0, 20)
        plt.ylim(0, 0.15)
        plt.plot(x, y, label='weibull pdf')
        plt.legend()
        plt.show()
```



Q4

```
In [5]: from scipy.stats import pearsonr, t

arr1 = np.random.normal(0, 1, 1000)
arr2 = np.random.normal(0, 1, 1000)
pearson_coefficient, pearson_pvalue = pearsonr(arr1, arr2)
print(f'Pearson Correlation Coefficient: {pearson_coefficient}, p-value: {pearson_pvalue}')

t_value = pearson_coefficient * np.sqrt((len(arr1) - 2) / (1 - pearson_coefficient**2))

t_distribution = 2 * t.cdf(-np.abs(t_value), len(arr1) - 2)
print(f'p-value from student-t distribution: {t_distribution}')
```

Pearson Correlation Coefficient: 0.007248419402012619, p-value: 0.8189221797940635 p-value from student-t distribution: 0.8189221797941082

We can see from the above results that p-value calculated from pearson correlation matches with the one calculated from student-t distribution