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#### CS20BTECH11063

#### **Data Science Analysis Assignment 4**

```
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

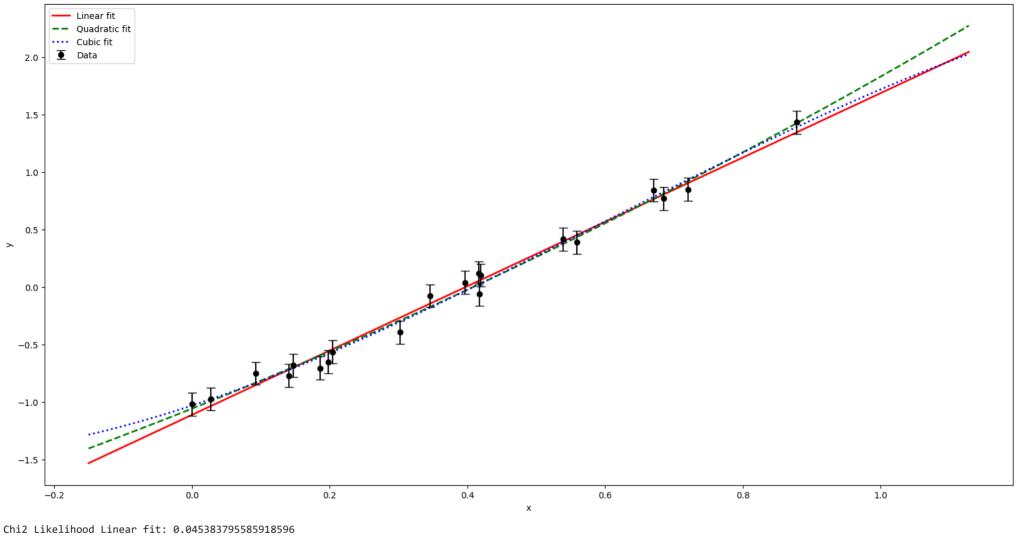
```
In [31]: from scipy.optimize import curve fit
         df = pd.read csv('testdata.csv', sep=' ')
         def linear(x, a, b):
             return a + b*x
         def quadratic(x, a, b, c):
             return a + b*x + c*x**2
         def cubic(x, a, b, c, d):
             return a + b*x + c*x**2 + d*x**3
         linear_fit, linear_cov = curve_fit(linear, df['x'], df['y'], sigma=df['sigma_y'])
         quadratic_fit, quadratic_cov = curve_fit(quadratic, df['x'], df['y'], sigma=df['sigma_y'])
         cubic_fit, cubic_cov = curve_fit(cubic, df['x'], df['y'], sigma=df['sigma_y'])
         # Print expression of all fits
         print('Linear fit: y = {:.2f} + {:.2f}x'.format(linear_fit[0], linear_fit[1]))
         print('Quadratic fit: y = \{:.2f\} + \{:.2f\}x + \{:.2f\}x^2'.format(quadratic fit[0], quadratic fit[1], quadratic fit[2]))
         print('Cubic fit: y = \{:.2f\}x + \{:.2f\}x^2 + \{:.2f\}x^3'.format(cubic_fit[0], cubic_fit[1], cubic_fit[2], cubic_fit[3]))
         # plot the data
         fig, ax = plt.subplots(figsize=(20, 10))
         x = np.linspace(-0.15, max(df['x']) + 0.25, 100)
         plt.plot(x, linear(x, linear_fit[0], linear_fit[1]), color='r', linestyle='-', linewidth=2)
         plt.plot(x, quadratic(x, quadratic_fit[0], quadratic_fit[1], quadratic_fit[2]), color='g', linestyle='--', linewidth=2)
         plt.plot(x, cubic(x, cubic fit[0], cubic fit[1], cubic fit[2], cubic fit[3]), color='b', linestyle=':', linewidth=2)
         ax.errorbar(df['x'], df['y'], yerr=df['sigma_y'], fmt='ok', ecolor='black', lw=1.5, capsize=5)
         plt.xlabel('x')
         plt.ylabel('y')
         plt.title('Data')
         plt.legend(['Linear fit', 'Quadratic fit', 'Cubic fit', 'Data'])
         # plt.xlim(0, 0.3)
```

```
# plt.ylim(-1.2, -0.5)
plt.show()
def AIC func(degree, parameters, x, y, sigma y):
    1 \text{ max} = 0
    if degree == 1:
        fit func = linear(x, parameters[0], parameters[1])
        1 max = np.sum(stats.norm.logpdf(y, loc=fit func, scale=sigma y))
    if degree == 2:
        fit func = quadratic(x, parameters[0], parameters[1], parameters[2])
        l_max = np.sum(stats.norm.logpdf(y, loc=fit_func, scale=sigma_y))
    if degree == 3:
        fit_func = cubic(x, parameters[0], parameters[1], parameters[2], parameters[3])
        1 max = np.sum(stats.norm.logpdf(y, loc=fit func, scale=sigma y))
    return -2*1 max + 2*(degree+1)
def BIC func(degree, parameters, x, y, sigma y, N):
    1 \text{ max} = 0
    if degree == 1:
        fit_func = linear(x, parameters[0], parameters[1])
        1 max = np.sum(stats.norm.logpdf(y, loc=fit func, scale=sigma y))
    if degree == 2:
        fit func = quadratic(x, parameters[0], parameters[1], parameters[2])
        l_max = np.sum(stats.norm.logpdf(y, loc=fit_func, scale=sigma_y))
    if degree == 3:
        fit_func = cubic(x, parameters[0], parameters[1], parameters[2], parameters[3])
        1 max = np.sum(stats.norm.logpdf(y, loc=fit func, scale=sigma y))
    return -2*1_max + np.log(N)*(degree+1)
def AICc func(degree, parameters, x, y, sigma y, N):
    1 \text{ max} = 0
    return AIC func(degree, parameters, x, y, sigma y) + 2*(degree+1)*(degree+2)/(N-degree-2)
def error_fit(degree, parameters, x, y, sigma_y):
    if degree == 1:
        fit func = linear(x, parameters[0], parameters[1])
    if degree == 2:
        fit func = quadratic(x, parameters[0], parameters[1], parameters[2])
   if degree == 3:
        fit_func = cubic(x, parameters[0], parameters[1], parameters[2], parameters[3])
    return (y-fit_func)/sigma_y
def chi2_func(degree, parameters, x, y, sigma_y):
    return np.sum(error_fit(degree, parameters, x, y, sigma_y)**2)
def dof_func(x, degree):
    return len(x) - degree - 1
def chi2 likelihood(degree, parameters, x, y, sigma y):
    return stats.chi2.pdf(chi2_func(degree, parameters, x, y, sigma_y), dof_func(x, degree))
# Print chi2 likelihood for all fits
print('Chi2 Likelihood Linear fit: {}'.format(chi2 likelihood(1, linear_fit, df['x'], df['y'], df['sigma_y'])))
```

```
print('Chi2 Likelihood Quadratic fit: {}'.format(chi2 likelihood(2, quadratic fit, df['x'], df['y'], df['sigma y'])))
print('Chi2 Likelihood Cubic fit: {}'.format(chi2 likelihood(3, cubic fit, df['x'], df['y'], df['sigma y'])))
# Print AIC, BIC, AICc for all fits in table format
print('AIC\tBIC\tAICc\tFit')
print('{:.2f}\t{:.2f}\tLinear'.format(AIC_func(1, linear_fit, df['x'], df['y'], df['sigma_y']),
                                             BIC func(1, linear fit, df['x'], df['v'], df['sigma v'], len(df['x'])),
                                             AICc_func(1, linear_fit, df['x'], df['y'], df['sigma_y'], len(df['x']))))
print('{:.2f}\t{:.2f}\t{:.2f}\tQuadratic'.format(AIC_func(2, quadratic_fit, df['x'], df['y'], df['sigma_y']),
                                                BIC_func(2, quadratic_fit, df['x'], df['y'], df['sigma_y'], len(df['x'])),
                                                AICc_func(2, quadratic_fit, df['x'], df['y'], df['sigma_y'], len(df['x']))))
print('{:.2f}\t{:.2f}\tCubic'.format(AIC_func(3, cubic_fit, df['x'], df['y'], df['sigma y']),
                                            BIC func(3, cubic fit, df['x'], df['y'], df['sigma y'], len(df['x'])),
                                            AICc_func(3, cubic_fit, df['x'], df['y'], df['sigma_y'], len(df['x']))))
# print p-value of quadratic fit and cubic fit wrt linear fit
print('p-value quadratic fit wrt linear fit as null hypothesis: {}'.format(stats.chi2.sf(-chi2_func(2, quadratic_fit, df['x'], df['y'], df['sigma_y'])
                                                                                        + chi2_func(1, linear_fit, df['x'], df['y'], df['sigma_y']),
                                                                     dof_func(df['x'], 1) - dof_func(df['x'], 2))))
print('p-value cubic fit wrt linear fit as null hypothesis: {}'.format(stats.chi2.sf(-chi2 func(3, cubic fit, df['x'], df['y'], df['sigma y'])
                                                                                     + chi2_func(1, linear_fit, df['x'], df['y'], df['sigma_y']),
                                                                 dof func(df['x'], 1) - dof func(df['x'], 3))))
```

Linear fit: y = -1.11 + 2.80xQuadratic fit:  $y = -1.06 + 2.38x + 0.50x^2$ Cubic fit:  $y = -1.03 + 1.97x + 1.74x^2 + -0.97x^3$ 





```
Chi2 Likelihood Quadratic fit: 0.036608447550140304
Chi2 Likelihood Cubic fit: 0.04215280601005979
AIC BIC AICc Fit
-40.04 -38.05 -39.33 Linear
-39.85 -36.86 -38.35 Quadratic
-38.26 -34.28 -35.59 Cubic
p-value quadratic fit wrt linear fit as null hypothesis: 0.1781327569531638
p-value cubic fit wrt linear fit as null hypothesis: 0.3288788441952259
```

It can be seen from the above results that Linear Model would be the best fit as it has the highest  $\chi^2$  Likelyhood value. Linear Model also has the least AIC, BIC, AICc values. Hence, Linear Model is the best fit for the given data.

```
In [23]: data = np.array([[ 0.42,  0.72,  0. ,  0.3 ,  0.15,
                          0.09, 0.19, 0.35, 0.4, 0.54,
                          0.42, 0.69, 0.2, 0.88, 0.03,
                          0.67, 0.42, 0.56, 0.14, 0.2 ],
                         [0.33, 0.41, -0.22, 0.01, -0.05,
                         -0.05, -0.12, 0.26, 0.29, 0.39,
                          0.31, 0.42, -0.01, 0.58, -0.2,
                          0.52, 0.15, 0.32, -0.13, -0.09],
                         [ 0.1 , 0.1 , 0.1 , 0.1 , 0.1 ,
                          0.1, 0.1, 0.1, 0.1, 0.1,
                          0.1, 0.1, 0.1, 0.1, 0.1,
                          0.1, 0.1, 0.1, 0.1, 0.1]])
         x, y, sigma_y = data
        linear fit, linear cov = curve fit(linear, x, y, sigma=sigma y)
         quadratic fit, quadratic cov = curve fit(quadratic, x, y, sigma=sigma y)
         # print AIC and BIC for linear fit and quadratic fit in table format
         print('AIC\tBIC\tFit')
         print('{:.2f}\t{:.2f}\tLinear'.format(AIC_func(1, linear_fit, x, y, sigma_y),
                                            BIC func(1, linear fit, x, y, sigma y, len(x)))
         print('{:.2f}\t{:.2f}\tQuadratic'.format(AIC_func(2, quadratic_fit, x, y, sigma_y),
                                               BIC func(2, quadratic_fit, x, y, sigma_y, len(x))))
         AIC
                BIC
                       Fit
```

```
AIC BIC Fit
-40.02 -38.03 Linear
-39.88 -36.90 Quadratic
```

As we can see that Linear Model has the smaller AIC and BIC value, hence it is the best fit model.

AIC and BIC is used to compare models based on strength of evidence rules. If a model has an AIC value that is 2 or more units smaller than the AIC value of another model, then the first model is considered to be significantly better than the second model. Similarly, lower BIC value indicates a better model.

The above results agree with the one shown on JVDP's blog.

## Q3

We will be using the paper titled, First-Passage Time Distribution of Brownian Motion as a Reliability Model by Y. S. Sherif and M. L. Smith, to understand the usage of Kolmogoorov-Smirnov test.

This paper tries to fit the inversr Gaussian Distribution model to observed failure data of high speed steel tools.

The Inverse Gaussian Distribution is given by:

$$f(x;\mu,\lambda) = \sqrt{rac{\lambda}{2\pi x^3}} \expigg(-rac{\lambda(x-\mu)^2}{2\mu^2 x}igg)$$

where  $\mu$  and  $\lambda$  are the mean and shape parameter respectively.

They substitute  $\lambda$  with  $\mu\gamma^2$ , where  $\gamma$  is the dimensionless shape parameter.

They use the failure data reported by Wager and Barash in the paper titled Study of the distribution of tool life in high speed steel cutting tools.

They hypothesize that the failure data is a sample from the Inverse Gaussian Distribution. They use the Kolmogorov-Smirnov test to test the hypothesis.

The value of K-S test value  $D_{max} = 0.1120$  and is smaller than the expected value at 5% significance level,  $D_{105}^{0.05} = 0.1130$ . Using lognormal distribution, the value of  $D_{max}$  is 0.1331 and they concluded that Inverse Gaussian Distribution is a better fit for the data.

However, the concern here is that KS test was used incorrectly here as KS test should not be used to compare distributions when the hypothesized distribution is calculated from the data itself as mentioned on the Penn State website.

References:

https://asaip.psu.edu/Articles/beware-the-kolmogorov-smirnov-test/

https://ieeexplore.ieee.org/stamp/stamp.jsp?arnumber=5220910

https://medium.com/@pabaldonedo/kolmogorov-smirnov-test-may-not-be-doing-what-you-think-when-parameters-are-estimated-from-the-data-2d5c3303a020

https://www.jstor.org/stable/pdf/2280095.pdf?refreqid=excelsior%3Ae3b35e6171e506d41fd5ad8bdf0340fc&ab\_segments=&origin=&initiator=&acceptTC=1

#### Q4

```
In [38]: # P vals for Higgs Boson
         pval higgs = [10**-1, 10**-2, 10**-3, 10**-5, 10**-7, 10**-9]
         significance_higgs = stats.norm.isf(pval_higgs)
         print('Significance for Higgs Boson from graph')
         for i in range(len(pval_higgs)):
             print('Significance for Higgs Boson for p value: {} : {}'.format(pval higgs[i], significance higgs[i]))
         # print('Significance for Higgs Boson from graph: {}'.format(significance higgs))
         pval higgs = 1.7 * 10 ** (-9)
         significance_higgs = stats.norm.isf(pval_higgs)
         print('\nSignificance for Higgs Boson discovery for p value: {} : {}'.format(pval higgs, significance_higgs))
         # P vals for LIGO
         pval ligo = 2 * 10 ** (-7)
         significance_ligo = stats.norm.isf(pval_ligo)
         print('Significance for LIGO: {}'.format(significance_ligo))
         # pval ligo = 2 * 10 ** (-7)
         # Z score = stats.norm.ppf(1 - pval ligo/2)
         # significance ligo = Z score
         # print('Significance for LIGO: {}'.format(significance_ligo))
```

```
# find chi2 goodness of fit for Super K

chi2_val = 65.2

dof_val = 67
chi2_dof = 1 - stats.chi2.cdf(chi2_val, dof_val)

print('\nGoodness of Fit for Super K: {}'.format(chi2_dof))

Significance for Higgs Boson from graph
Significance for Higgs Boson for p value: 0.1 : 1.2815515655446004
Significance for Higgs Boson for p value: 0.01 : 2.3263478740408408
Significance for Higgs Boson for p value: 0.001 : 3.090232306167813
Significance for Higgs Boson for p value: 1e-05 : 4.264890793922825
Significance for Higgs Boson for p value: 1e-07 : 5.1993375821928165
Significance for Higgs Boson for p value: 1e-09 : 5.9978070150076865

Significance for Higgs Boson discovery for p value: 1.7000000000000001e-09 : 5.911017938341624
Significance for LIGO: 5.068957749717791
```

Goodness of Fit for Super K: 0.5394901931099038