# 實驗物理學 (二) 實驗日誌

Fundamental Python Chi-square fitting - 3

Group 2

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# 1 實驗步驟、初步結果

### 1.1 Practice 1

1. By using previous knowledge, choose your own function to generate data points.

2. Generate 1000 sets of data points with the same function with normally distributed random noise.

```
def parabolic(x, a, b, c):
    return a * x**2 + b * x + c

✓ 0.0s

# Generate synthetic data

true_a = 2.0

true_b = 1.0

true_c = 0.5

sigma0 = 3.0

# Generate data

n_data = 10

n_sets = 1000

x = np.linspace(-5, 5, n_data)

✓ 0.0s

Python
```

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3. Fit these data points with the function you used to generate the data points.

4. Calculate the  $\chi^2$  values of each fitting result.

```
all_data = []
fit_results = []
chi2_values = []

> 0.0s

Python

for i in range(n_sets):
    # Generate synthetic data with noise
    y = parabolic(x, true_a, true_b, true_c) + np.random.normal(0, sigma0, size=x.shape)
    all_data.append(y)
    # Fit the data
    popt, pcov = curve_fit(parabolic, x, y)

# Calculate chi-squared
    residuals = y - parabolic(x, *popt)
    chi2_value = np.sum((residuals / sigma0)**2)

fit_results.append(popt)
    chi2_values.append(chi2_value)

> 0.1s
Python
```

5. Create your own **p-value** calculator to calculate the **p-value** of each  $\chi^2$  instead of using functions in scipy.stats.

```
from math import gamma

def chi2_p_value(chi2_value, dof):
    """
    Calculate the p-value for a given chi-square value and degrees of freedom (dof).
    """
    # Compute the incomplete gamma function (upper tail)
    incomplete_gamma = gamma(dof / 2) - sum((chi2_value / 2)**k / gamma(k + 1) for k in range(dof // 2))

# Compute the p-value
    p_value = incomplete_gamma / gamma(dof / 2)
    return p_value

# Example usage
    dof = 7 # degrees of freedom
    p_values = [chi2_p_value(chi2, dof) for chi2 in chi2_values]
    print(p_values[:10]) # Print the first 10 p-values
```

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```
plt.figure(figsize=(10, 6))
x_plot = np.linspace(-8, 8, 500)
plt.plot(x_plot, parabolic(x_plot, true_a, true_b, true_c), color='C4', label='True model',
          linewidth=4, linestyle='-')
sampled_sets = np.random.choice(n_sets, 200, replace=False)
for i in sampled_sets:
    plt.scatter(x, all_data[i, :], color='C1', alpha=0.3)
    plt.plot(x_plot, parabolic(x_plot, *fit_results[i]), color='plum', alpha=0.1,
             linewidth=1.5, linestyle=':')
plt.scatter([], [], color='C1', alpha=0.3, label='Sample points')
plt.plot([], [], color='plum', alpha=0.5, linewidth=1.5, label='Fitted lines', linestyle=':')
plt.title('Sample scatter plot', fontsize=16)
plt.xlim(-5.5, 5.5)
plt.ylim(-10, 70)
plt.xlabel('x', fontsize=14)
plt.xticks(fontsize=12)
plt.ylabel('y', fontsize=14)
plt.yticks(fontsize=12)
plt.legend(fontsize=14, loc='upper left', framealpha=1)
plt.savefig("output1_1.pdf", transparent=True)
plt.show()
                                                                                                                                      Python
```

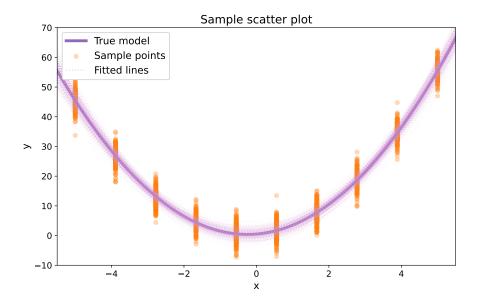


Figure 1: 使用已有的知識選擇的數學函數,並加上隨機雜訊產生之數據,及每組數據的擬合曲線

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#### 6. Plot the histogram of $\chi^2$ .

```
plt.figure(figsize=(10, 6))
plt.hist(chi2_values, bins=30, density=True, alpha=0.7, label='\chi^2', color='C6',
       histtype='stepfilled')
x_{chi} = np.linspace(0, 50, 500)
linewidth=3.5)
plt.plot([np.mean(chi2_values), np.mean(chi2_values)], [0, 0.2], linestyle='--',
       color="purple", lw=2, label='__no_legend__')
plt.text(np.mean(chi2_values)+0.5, 0.12, f'$\mu$={np.mean(chi2_values):.3f}',
        fontsize=14, color='purple')
plt.title('Chi-square Distribution (dof=8)', fontsize=16)
plt.xlabel('Chi-square value', fontsize=14)
plt.ylabel('Probability density', fontsize=14)
plt.xlim(left=0, right=30)
plt.ylim(bottom=0, top=0.13)
plt.legend(fontsize=14, framealpha=1)
plt.savefig("output1_2.pdf", transparent=True)
plt.show()
```

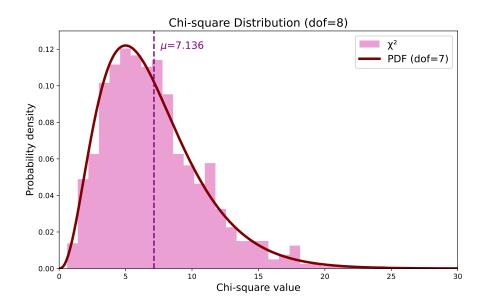


Figure 2:  $\chi^2$ 直方圖與dof = 7時的PDF

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7. Mark a certain  $\chi^2$  value with its corresponding **p-value** on the histogram.

```
specific_chi2_value = chi2_values[i] # Use the χ² value at index i
specific_p_value = chi2.sf(specific_chi2_value, df=dof) # Calculate p-value using survival function
plt.figure(figsize=(10, 6))
plt.hist(chi2\_values, \ bins=30, \ density=True, \ alpha=0.7, \ label='\chi^2', \ color='C6', \ histtype='stepfilled')
# Plot the χ² PDF
x_chi = np.linspace(0, 50, 500)
plt.plot(x_chi, chi2.pdf(x_chi, df=dof), linestyle='-', color='maroon', label='PDF (dof=7)', linewidth=3.5)
plt.axvline(specific\_chi2\_value, color='blue', linestyle='--', linewidth=2, label=f'\chi^2=\{specific\_chi2\_value:.2f\}')\\ plt.text(specific\_chi2\_value + 0.5, 0.1, f'p=\{specific\_p\_value:.3f\}', fontsize=14, color='blue')
# Add labels, title, and legend plt.title('Chi-square Distribution (dof=8)', fontsize=16)
plt.xlabel('Chi-square value', fontsize=14)
plt.ylabel('Probability density', fontsize=14)
plt.xlim(left=0, right=30)
plt.ylim(bottom=0, top=0.13)
plt.legend(fontsize=14, framealpha=1)
# Save and show the plot
plt.savefig("output1_3.pdf", transparent=True)
plt.show()
0.1s
```

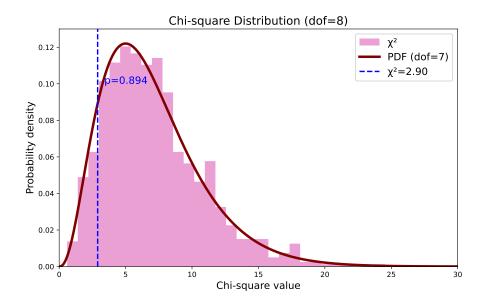


Figure 3: 標上對應之P-value

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```
# Sort the chi-squared values
sorted_chi2_values = np.sort(chi2_values)

# Compute the cumulative probabilities
cumulative_probabilities = np.arange(1, len(sorted_chi2_values) + 1) / len(sorted_chi2_values)

# Plot the CDF
plt.figure(figsize=(10, 6))
plt.plot(sorted_chi2_values, cumulative_probabilities, label='Empirical CDF', color='C0', linewidth=3.5)

# Overlay the theoretical CDF for comparison
theoretical_cdf = chi2_cdf(sorted_chi2_values, df=dof)
plt.plot(sorted_chi2_values, theoretical_cdf, label='Theoretical CDF (dof=7)', color='C3', linestyle='--', linewidth=3.5)

# Add labels, title, and legend
plt.title('Cumulative pistribution Function (CDF)', fontsize=16)
plt.xlabel('Cimulative probability', fontsize=14)
plt.ylabel('Cumulative probability', fontsize=14)
plt.legend(fontsize=14, framealpha=1)

# Save and show the plot
plt.savefig("output1_4.pdf", transparent=True)
plt.show()
```

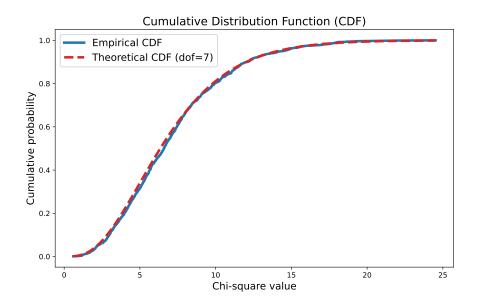


Figure 4: Cumulative Distribution Function

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#### 1.2 Practice 2

1. Using prior knowledge, select an appropriate function with noise to generate your dataset.

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```
plt.figure(figsize=(10, 6))
x_plot = np.linspace(-8, 8, 500)
plt.plot(x_plot, parabolic(x_plot, true_a, true_b, true_c), color='C4', label='True model',
         linewidth=4, linestyle='-')
sampled_sets = np.random.choice(n_sets, 200, replace=False)
for i in sampled_sets:
    plt.scatter(x, all_data[i, :], color='C1', alpha=0.3)
    plt.scatter([], [], color='C1', alpha=0.3, label='Sample points')
plt.plot([], [], color='plum', alpha=0.5, linewidth=1.5, label='Fitted lines', linestyle=':')
plt.title('Sample scatter plot', fontsize=16)
plt.xlim(-5.5, 5.5)
plt.ylim(-10, 70)
plt.xlabel('x', fontsize=14)
plt.xticks(fontsize=12)
plt.ylabel('y', fontsize=14)
plt.yticks(fontsize=12)
plt.legend(fontsize=14, loc='upper left', framealpha=1)
plt.savefig("output2_1.pdf", transparent=True)
plt.show()
                                                                                                                       Python
```

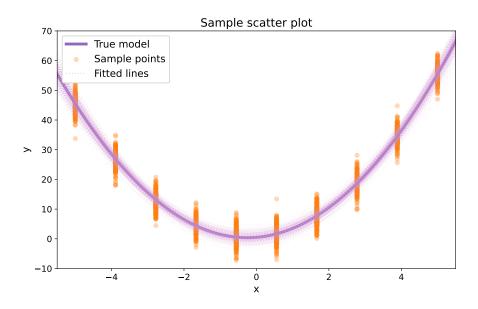


Figure 5: 使用已有的知識選擇的數學函數,並加上隨機雜訊產生之數據,及每組數據的擬合曲線

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2. Plot a histogram of the  $\chi^2$  values obtained from fitting each dataset.

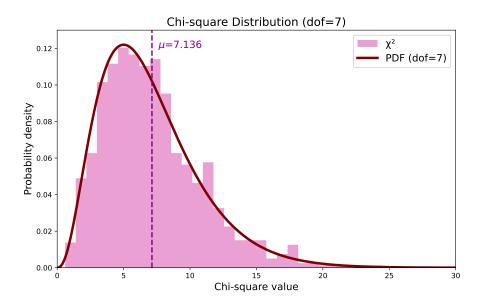


Figure 6: 每組數據對應的 $\chi^2$ 值之直方圖,與dof = 7時對應的PDF

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- 3. Mark the **p-value** get from the fitting result of mean (y).
- 4. You might get a  $\chi^2$  of the mean (y) fitting result close to zero and a **p-value** close to 1.

```
mean_y = np.mean(all_data, axis=0)
parms_mean, _ = curve_fit(parabolic, x, mean_y)
chisq_mean = np.sum((mean_y - parabolic(x, *parms_mean))**2 / sigma0**2)
p_mean = 1 - chi2.cdf(chisq_mean, df=7)

print('chi-square of mean(y)= ', chisq_mean)
print('p-value of mean(y)= ', p_mean)
```

```
chi-square of mean(y)= 0.004356074591544417 p-value of mean(y)= 0.99999999586144
```

Figure 7: 平均數據的 $\chi^2$ 值和其對應的P-value

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5. Also mark 5% significance level, and 95% significance level on the histogram.

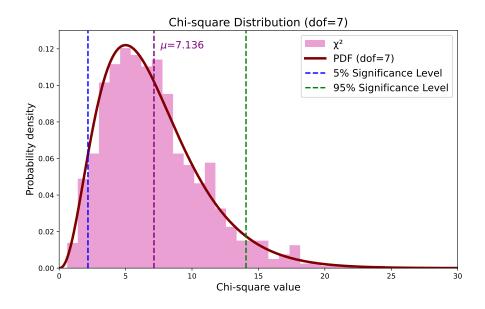


Figure 8: 由Fig.6的基礎下加入5%和95%的Significance level

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## 2 初步分析

#### 2.1 Practice 1

觀察Fig.1,紫色粗線爲true model,設定爲: $y=2x^2+x+0.5$ ;擬合線爲紫色虛線。可以看出部分擬合線非常貼近眞實模型,表示這部分擬合線的資料誤差小、擬合結果準確,有些則有明顯偏離,表示該組資料雜訊大或者擬合出來的參數有誤差。擬合線與資料點大致上皆吻合,但不一定會完全與眞實曲線重合。雖然使用相同的擬合方式,但因爲受到noise的隨機數值影響,擬合出的結果會不相同,即是擬合結果的variability (變異性)。

而Fig.2是擬合結果的 $\chi^2$ 分布直方圖,並與理論上的卡方分布PDF進行比較,自由度爲 dof=10`3=7。從圖中可見實際 $\chi^2$ 值的直方圖與理論的卡方 PDF 曲線高度相符,且圖中 $\chi^2$ 平均值(虛線)與理論自由度相符,表示擬合過程是合理的,且高斯雜訊與誤差模型( $\sigma=3$ )和擬合模型之間的假設是一致的,也沒有明顯系統性偏誤。實際 $\chi^2$ 分布與理論曲線在形狀上相符,代表殘差的統計性質符合高斯假設。整體 $\chi^2$ 值沒有出現明顯右偏或重尾現象,代表大多數擬合皆落在合理範圍以內,沒有出現明顯異常。

Fig.3在原有 $\chi^2$ 分布直方圖基礎上,再多標示出一組擬合結果的 $\chi^2=2.90$ 及其對應之p-value=0.894。此處所選 $\chi^2$ 值爲X(以藍線標示),對應的 p-value 爲 Y,表示該擬合結果在所有結果中的相對位置。若 p-value 明顯偏小(例如 <0.05),可視爲模型擬合不良;若 p-value 適中,則該擬合與理論模型一致,未有顯著偏差。

最後,Fig.4為Chi – square 分布的CDF。藍色實線爲實際模擬資料的經驗累積分布( $Empirical\ CDF$ );紅色虛線爲理論上的卡方分布累積函數( $Theoretical\ CDF$ )(自由度 = 7)。這張圖畫出了模擬出的 $\chi^2$ 值之累積分布(藍線)與理論卡方分布(自由度7)的CDF(紅線)對比結果。模擬結果與理論分布相符,顯示生成資料的統計性質與理論預期相差不大。這種比較可以當駔是擬合殘差是否爲常態分布的間接驗證方式之一,如果有明顯的差異,可能代表模型設計、誤差估計或noise生成中出現問題。

#### 2.2 Practice 2

從Fig.5可以看出我們成功在 $y=ax^2+bx+c$ 這個模型上加入不同程度的雜訊;並可看出越接近true model時,每組雜訊所對應的擬合曲線就越密集。從Fig.6來看,可見我們所建立的不同資料組的擬合所對應是 $\chi^2$ 值有符合當自由爲7時,其PDF的趨勢,而 $\chi^2$  value平均值爲7.136,透過先前所學,將其除以自由度:

$$\chi_{\nu}^2 = \frac{\chi^2}{\nu} = \frac{7.136}{7} \approx 1.019 \approx 1$$

可確認其擬合程度很好。而Fig.8可以看見我們 $\chi^2$ 的平均值是落在 $5\%\sim95\%$  significance level之間,表示我們的擬合結果是在正常的範圍內,是合理的模型;最後,Fig.7得到利用平均的數據做擬合時,得到的 $\chi^2$  value趨近於 $0 \times P$ -value接近1,符合理論結果。

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## 3 具體說明實驗遇到的問題,或分析可能的問題

在practice 2中,我們原先計算平均數據的 $\chi^2$  value並沒有接近0,經過檢查才發現將其寫成parms\_mean, \_ = curve\_fit (parabolic, x, y),這會導致我們使用的y值是最後一組數據的y值,並非其平方,導致結果不符預期;將其更正爲parms\_mean, \_ = curve\_fit (parabolic, x, mean\_y),就獲得到理想結果了(詳情見步驟4)。

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