

[CC511] Homework 9 20204817 Federico Berto

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1 Homework 9 - Federico Berto

```
[2]: # Importing useful libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from scipy import stats
import statsmodels.stats.weightstats as sms
from scipy.stats import t
from scipy.stats import z
from scipy.stats import norm
import math
```

1.1 Exercise 10.1.4

The 95% confidence can be calculated via $z_{0.05}$

```
[8]: z = norm.ppf(1-0.05)
print(z)
```

1.6448536269514722

The confidence interval is:

$$\left(\frac{35}{44} - \frac{1.645}{44} \times \sqrt{\frac{35 \times (44 - 35)}{44}}, 1 \right) = (0.695, 1) \quad (1)$$

1.2 Exercise 10.1.8

We know that $L = 2z_{\alpha/2} \sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$, therefore:

$$n \geq \frac{4z_{\alpha/2}^2 \hat{p}(1-\hat{p})}{L^2} \quad (2)$$

```
[9]: z = norm.ppf(1-0.005)
print(z)
```

2.5758293035489004

Given $L = 0.04$, then we have for $\hat{p} = 0.50$:

$$n \geq \frac{4 \times 2.5758 \times 0.50(1 - 0.50)}{0.04^2} = 1609.9 \quad (3)$$

So, n has to be at least 1610.

If $\hat{p} = 0.40$:

$$n \geq \frac{4 \times 2.5758 \times 0.40(1 - 0.40)}{0.04^2} = 1545.5 \quad (4)$$

In this case, n has to be at least 1546.

1.3 Exercise 10.1.18

a) The hypotheses are:

- $H_0 : p_A \leq 0.05$
- $H_A : p_B > 0.05$

We calculate the statistics for the normal approximation as:

$$z = \frac{x - np_0 - 0.5}{\sqrt{np_0(1 - p_0)}} = \frac{13 - 62 \times 0.05 - 0.5}{\sqrt{62 \times 0.05 \times (1 - 0.05)}} = 5.48 \quad (5)$$

```
[23]: print('p-value: ', (1 - norm.cdf(5.48)) )
```

p-value: 2.1266291838628604e-08

We can conclude that with the p value close to 0, there is sufficient evidence to conclude that the probability of breakdown is above 5%.

b) The 95% confidence can be calculated via $z_{0.05}$

```
[ ]: z = norm.ppf(1-0.05)
      print(z)
```

Thus the confidence interval is:

$$\left(\frac{13}{62} - \frac{1.645}{62} \times \sqrt{\frac{13 \times (62 - 13)}{62}}, 1 \right) = (0.125, 1) \quad (6)$$

1.4 Exercise 10.2.2

a) The 95% confidence can be calculated via $z_{0.005}$

```
[26]: z = norm.ppf(1-0.005)
      print(z)
```

2.5758293035489004

The confidence interval can be calculated as such:

$$\hat{p}_a - \hat{p}_b \pm z_{\alpha/2} \sqrt{\frac{\hat{p}_a(1 - \hat{p}_a)}{n} + \frac{\hat{p}_b(1 - \hat{p}_b)}{m}} \quad (7)$$

So, the confidence interval is:

$$\frac{4}{50} - \frac{10}{50} \pm 2.576 \times \sqrt{\frac{4 \times (50 - 4)}{50^3} + \frac{10 \times (50 - 10)}{50^3}} = (-0.296, 0.056) \quad (8)$$

b) We can use the pooled probability estimate, which is $\hat{p} = \frac{x+y}{n+m} = \frac{4+10}{50+50} = 0.14$

The test statistics is:

$$z = \frac{\hat{p}_A - \hat{p}_B}{\sqrt{\hat{p}(1 - \hat{p})(\frac{1}{n} + \frac{1}{m})}} \quad (9)$$

Therefore:

$$z = \frac{\frac{4}{50} - \frac{10}{50}}{\sqrt{0.14 \times (1 - 0.14) \times (\frac{1}{50} + \frac{1}{50})}} = -1.729 \quad (10)$$

```
[33]: print('p value: ', 2*norm.cdf(-1.729))
```

p value: 0.08380909507894739

c) The confidence interval becomes:

$$\frac{40}{500} - \frac{100}{500} \pm 2.576 \times \sqrt{\frac{40 \times (500 - 40)}{500^3} + \frac{100 \times (500 - 100)}{500^3}} = (-0.176, 0.064) \quad (11)$$

We can use the pooled probability estimate, which is $\hat{p} = \frac{x+y}{n+m} = \frac{40+100}{500+500} = 0.14$

$$z = \frac{\frac{40}{500} - \frac{100}{500}}{\sqrt{0.14 \times (1 - 0.14) \times (\frac{1}{500} + \frac{1}{500})}} = -5.468 \quad (12)$$

```
[35]: print('p value: ', 2*norm.cdf(-5.468))
```

p value: 4.551418826709858e-08

In this case, the p-value becomes almost zero.

1.5 Exercise 10.2.12

```
[36]: z = norm.ppf(1-0.05)
print(z)
```

1.6448536269514722

We can construct the upper confidence bound for $p_A - p_B$, where p_A is the probability of following the link in the original design and p_B is the probability after the modification, as following:

$$\left(-1, \frac{22}{542} - \frac{64}{601} + 1.645 \times \sqrt{\frac{22 \times (542 - 22)}{542^3} + \frac{64 \times (601 - 64)}{601^3}} \right) = (-1, -0.041) \quad (13)$$

a) The hypotheses are:

- $H_0 : p_A \geq p_B$
- $H_A : p_A < p_B$

We can use the pooled probability estimate, which is $\hat{p} = \frac{x+y}{n+m} = \frac{22+64}{542+601} = 0.0752$

The test statistics becomes

$$z = \frac{\frac{22}{542} - \frac{64}{601}}{\sqrt{0.0752 \times (1 - 0.0752) \times (\frac{1}{542} + \frac{1}{601})}} = -4.22 \quad (14)$$

```
[38]: print('p value: ', norm.cdf(-4.22))
```

```
p value: 1.2215115925253025e-05
```

Being the p-value almost zero, there is sufficient evidence to conclude that the probability of the link being following has increased.

1.6 Exercise 10.3.6

Soft drink type | Formulation

Formulation I 225

Formulation II 223

Formulation III 152

If the formulations are equally likely, then the expected cell frequencies are: $e_i = 600 \times \frac{1}{3} = 200$
We can use the Pearson chi-square statistics to calculate the *p-value*:

$$X^2 = \sum_{i=1}^k \frac{(x_i - e_i)^2}{e_i} \quad (15)$$

yielding

$$X^2 = \frac{(225 - 200)^2}{200} + \frac{(223 - 200)^2}{200} + \frac{(152 - 200)^2}{200} = 17.29 \quad (16)$$

The p-value is given by $P(\chi_2^2 \geq 17.29)$:

```
[41]: from scipy.stats import chi2
print('p-value: ', chi2.sf(17.29, 2))
```

```
p-value: 0.00017600467513708998
```

We can state that it is not plausible that the formulations of the soft drinks are equally likely.

1.7 Exercise 10.3.14

We can calculate the probabilities given the Weibull distribution with $\lambda = 0.065$ and $a = 0.45$: - $p_1^* = P(X \leq 24) = 1 - e^{-(\lambda x)^\alpha} = 1 - e^{-(0.065 \times 24)^{0.45}} = 0.705$ - $p_2^* = P(X \leq 48) = 1 - e^{-(\lambda x)^\alpha} = 1 - e^{-(0.065 \times 48)^{0.45}} = 0.812$ - $p_3^* = P(X \leq 72) = 1 - e^{-(\lambda x)^\alpha} = 1 - e^{-(0.065 \times 72)^{0.45}} = 0.865$

The observed cell frequencies are: 12, 53, 39, 21. Therefore: - $e_1 = np_1^* = 125 \times (0.705) = 88.125$ - $e_2 = np_2^* = 125 \times (0.812 - 0.705) = 13.375$ - $e_3 = np_3^* = 125 \times (0.865 - 0.812) = 6.625$ - $e_4 = np_4^* = 125 \times (1 - 0.865) = 16.87$

We use the Pearson X^2 statistics which yield

$$X^2 = \sum_{i=1}^k \frac{(x_i - e_i)^2}{e_i} \sim 342.4 \quad (17)$$

```
[43]: print('p-value: ', chi2.sf(342.4, 3))
```

p-value: 6.595603058895007e-74

Which is basically 0. Thus, the null hypothesis of the Weibull distribution approximation is clearly rejected.

1.8 Exercise 10.4.2

We do the experiment via the software package:

```
[56]: from scipy.stats import chi2_contingency
aptocc = np.array([[48,111,186,142],[71,89,174,181],[63,95,181,190]])
aptocc = pd.DataFrame(data=aptocc,
index=['No Fertilizer', 'Fertilizer I', 'Fertilizer II'], columns=['Dead', 'Slow_
→Growth', 'Medium Growth', 'Strong growth'])
print("Observed cell frequencies:\n", aptocc)
```

Observed cell frequencies:

	Dead	Slow Growth	Medium Growth	Strong growth
No Fertilizer	48	111	186	142
Fertilizer I	71	89	174	181
Fertilizer II	63	95	181	190

```
[57]: chi, pvalue, dof, expctd = chi2_contingency(aptocc)
print("Pearson's Chi-squared test \nX-squared = %.4f, p-value = %.4f, df = %d,"
→%(chi,
pvalue, dof))
print("Expected cell frequencies:\n", pd.DataFrame(expctd, index=['No_
→Fertilizer', 'Fertilizer I', 'Fertilizer II'], columns=['Dead', 'Slow_
→Growth', 'Medium Growth', 'Strong growth']))
```

Pearson's Chi-squared test

X-squared = 13.6591, p-value = 0.0337, df = 6,

Expected cell frequencies:

	Dead	Slow Growth	Medium Growth	Strong growth
No Fertilizer	57.892880	93.837361	172.088178	163.181581
Fertilizer I	61.221424	99.232528	181.982364	172.563684
Fertilizer II	62.885696	101.930111	186.929458	177.254735

There is a suggestion that the growth pattern is different for the different growing conditions, but there is no overwhelming evidence.

1.9 Exercise 10.4.6

For the 2×2 contingency table we obtain:

	c1	c2	Sum up
r1	x11	x12	$x1 = x11 + x12$
r2	x21	x22	$x2 = x21 + x22$
Sum up	$x1 = x11 + x21$	$x2 = x12 + x22$	$n = x1 + x2 = x1 + x2$

$$e_{ij} = \frac{x_{i.} \cdot x_{.j}}{n}$$

So, we have:

$$\frac{(x_{11} - e_{11})^2}{e_{11}} = \frac{(x_{11} - \frac{(x_{11}+x_{12})(x_{11}+x_{21})}{x_{11}+x_{21}+x_{12}+x_{22}})^2}{\frac{x_{1.} \cdot x_{.1}}{x_{11}+x_{21}+x_{12}+x_{22}}} = \frac{x_{2.} \cdot x_{.2} (x_{11}x_{22} - x_{12}x_{21})^2}{nx_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}}$$

We also have in a similar way:

$$\frac{(x_{21} - e_{21})^2}{e_{21}} = \frac{x_{1.} \cdot x_{.2} (x_{11}x_{22} - x_{12}x_{21})^2}{nx_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}}$$

and we can easily obtain the other coefficients in the same way. Finally, we can prove the result by substituting:

$$\begin{aligned} \chi^2 &= \sum_{i=1}^2 \sum_{j=1}^2 \frac{(x_{ij} - e_{ij})^2}{e_{ij}} = \frac{x_{2.} \cdot x_{.2} (x_{11}x_{22} - x_{12}x_{21})^2}{nx_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}} + \frac{x_{2.} \cdot x_{.1} (x_{11}x_{22} - x_{12}x_{21})^2}{nx_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}} \\ &\quad + \frac{x_{1.} \cdot x_{.1} (x_{11}x_{22} - x_{12}x_{21})^2}{nx_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}} + \frac{x_{1.} \cdot x_{.2} (x_{11}x_{22} - x_{12}x_{21})^2}{nx_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}} \\ &= \frac{(x_{11}x_{22} - x_{12}x_{21})^2}{nx_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}} (x_{2.} \cdot x_{.2} + x_{2.} \cdot x_{.1} + x_{1.} \cdot x_{.2} + x_{1.} \cdot x_{.1}) \\ &= \frac{(x_{11}x_{22} - x_{12}x_{21})^2}{nx_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}} (x_2 + x_1)^2 = \frac{n(x_{11}x_{22} - x_{12}x_{21})^2}{x_{2.} \cdot x_{.2} x_{1.} \cdot x_{.1}} \end{aligned}$$

1.10 Exercise 10.4.10

We may conduct the experiment via the software package

```
[73]: from scipy.stats import chi2_contingency
aptocc = np.array([[31, 17, 9], [36,9,4], [56, 19, 15]])
aptocc = pd.DataFrame(data=aptocc,
index=['A', 'B', 'C'], columns=['Minor Cracking', 'Medium cracking', 'Severe_
↪cracking'])
print("Observed cell frequencies:\n", aptocc)
```

Observed cell frequencies:

	Minor Cracking	Medium cracking	Severe cracking
A	31	17	9
B	36	9	4
C	56	19	15

```
[74]: chi, pvalue, dof, expctd = chi2_contingency(aptocc)
print("Pearson's Chi-squared test \nX-squared = %.4f, p-value = %.4f, df = %d,"
↪%(chi,
pvalue, dof))
print("Expected cell frequencies:\n", pd.DataFrame(expctd, index=['A', 'B',
↪'C'], columns=['Minor Cracking', 'Medium cracking', 'Severe cracking']))
```

Pearson's Chi-squared test

X-squared = 5.0237, p-value = 0.2849, df = 4,

Expected cell frequencies:

	Minor Cracking	Medium cracking	Severe cracking
A	35.770408	13.086735	8.142857
B	30.750000	11.250000	7.000000
C	56.479592	20.663265	12.857143

Therefore, the null hypothesis of independence is plausible and we have no overwhelming evidence to state the three types of asphalt are different with respect to cracking.