[CC511] Homework 8 20204817 Federico Berto

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1 Homework 8 - Federico Berto

```
[95]: # 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 norm
  import math
```

1.1 9.2.8

```
[26]: data=pd.read_excel("DS9.2.8.xls")
  dat_A= data['Standard Antibiotic']
  dat_B= data['New Antibiotic']
  # We create the difference distribution
  print("Difference distribution:\n", diff.describe())
```

Difference distribution:

```
count
         8.000000
mean
         1.375000
         1.784657
std
min
        -1.100000
25%
         0.025000
50%
         1.250000
75%
         2.550000
         4.100000
max
dtype: float64
```

We have the difference $z_i=x_i-y_i$ has a mean $\bar{z}=\mu_S-\mu_N$ where S is the standard antibiotic and N is the new one. So, $\bar{z}=1.375$ and the sample standard deviation is s=1.7847 The hypoteses are: - $H_0: \mu \leq 0$ - $H_A: \mu > 0$

Test statistics:

$$t = \frac{\sqrt{n}\bar{z}}{s} = \frac{\sqrt{8} \times 1.375}{1.7847} = 2.179 \tag{1}$$

```
[27]: print(("P-value using sf (1-cdf) = {:.4f}").format(t.sf(2.179, 7)))
```

P-value using sf (1-cdf) = 0.0329

So, there is some evidence, although it is not strong at all, that the new antibiotic is quicker than the standard one

$1.2 \quad 9.3.2$

We first declare some variables we will use based on the distribution:

```
[81]: # Population A
n = 14
x_bar = 32.45
s_x = 4.30

# Population B
m = 14
y_bar = 41.45
s_y = 5.23
alpha = 0.01 # = 1 - confidence level
```

The pooled variance is given by the following expression:

$$s_p^2 = \frac{(n-1)s_x^2 + (m-1)s_y^2}{n+m-2} \tag{2}$$

a) Confidence interval with 99.00% confidence level: (-14.0282)(-3.9718)

Now, we need to calculate the degrees of freedom by the following formula:

$$v^* = \frac{(s_x^2/n + s_y^2/m)^2}{s_x^4/n^2(n-1) + s_y^4/m^2(m-1)}$$
(3)

b)

Confidence interval with 99.00% confidence level: (-14.0440)(-3.9560)

We consider the statistics is

$$T = \frac{\bar{x} - \bar{y} - (\mu_A - \mu_B)}{\sqrt{\frac{s_x^2}{n} + \frac{s_y^2}{m}}} \tag{4}$$

The hypoteses are: - $H_0: \mu_A = \mu_B$ - $H_A: \mu_A \neq \mu_B$

```
[88]: # Calculate the t-statistics
def T_statistic(x_bar, y_bar, s_x, s_y, n, m, mu_difference = 0):
    return ( (x_bar - y_bar - mu_difference) / math.sqrt(s_x**2/n + s_y**2/m))

t_stat = T_statistic(x_bar, y_bar, s_x, s_y, n, m, mu_difference = 0)
print("|t| value: ", abs(t_stat))
print("Critical point: ", t.ppf(1- alpha/2, n + m -2 ) )
```

|t| value: 4.973595778437414

Critical point: 2.7787145333289134

The null hypotesis is rejected because |t| is greater than the critical point. The *p*-value can be calculated as: $2 \times P(t_{26} \ge 4.97) = 0.000$

```
[91]: print(("P-value = {:.4f}").format(2 * t.cdf(t_stat, n + m -2)))
```

P-value = 0.0000

$1.3 \quad 9.3.10$

From now on, let's directly solve the exercises in Python

```
[92]: # Population A
       n = 38
       x_bar = 5.782
       # Population B
       m = 40
       y_bar = 6.443
       sigma = 2.0
       alpha = 0.01 # = 1 - confidence level
      We calculate the p-value based on H_0: \mu_A - \mu_B > 0
[99]: # Calculate the t-statistics
       def Z_statistic(x_bar, y_bar, s_x, s_y, n, m, mu_difference = 0):
           return ( (x_bar - y_bar - mu_difference) / math.sqrt(s_x**2/n + s_y**2/m))
       Z_stat = Z_statistic(x_bar, y_bar, sigma, sigma, n, m)
       print("a)")
       print("Z-statistics: ", Z_stat)
       print("p-value: ", norm.cdf(Z_stat))
      a)
      Z-statistics: -1.4589686381754354
      p-value: 0.07228686987180125
[107]: def wing_span(alpha, n, m, s_x, s_y):
           return norm.ppf(1-alpha) * math.sqrt( s_x**2/n + s_y**2/m)
       def print_confidence_interval(alpha, mu, wing_span, lower_bound = False, u
        →upper_bound = False):
           if upper_bound:
               print(("Confidence interval with \{:.2f\}% confidence level: (-\omega)(\{:.
        \rightarrow4f})").format(((1-alpha)*100), mu + wing_span))
               return
           if lower bound:
               print(("Confidence interval with {:.2f}% confidence level: ({:.
        \rightarrow4f})(\emptyset)").format(((1-alpha)*100), mu - wing_span))
               return
           # Default case: double bounded
           print(("Confidence interval with {:.2f}% confidence level: ({:.4f})({:.
        \rightarrow 4f)").format(((1-alpha)*1, mu - wing_span, mu + wing_span)))
```

```
diff = x_bar - y_bar
wing_span = wing_span(alpha, n, m, sigma, sigma)
print("b)")
print_confidence_interval(alpha, diff, wing_span, upper_bound = True)
```

b)

Confidence interval with 99.00% confidence level: $(-\omega)(0.3930)$

1.4 9.3.14

```
[110]: print("t value: ", t.ppf(1-alpha/2, n+m-2))
```

t value: 2.7787145333289134

The length follows the following equation:

$$L = 2 \times t_{\alpha/2,\nu} \sqrt{\frac{s_A^2}{n} + \frac{s_B^2}{m}} \le 5 \tag{5}$$

So we get:

$$n = m \ge \frac{4t_{\alpha/2,\nu}(s_A^2 + s_B^2)}{L_0^2} = \frac{4 \times 2.779^2 \times (4.3^2 + 5.23^2)}{5^2} = 56.646$$
 (6)

So, 57 samples will suffice, which means 57 - 14 = 43 more samples should be collected from each population

$1.5 \quad 9.3.22$

```
[115]: # Population A
n = 16
x_bar = 1.053
s_x = 0.058

# Population B
m = 16
y_bar = 1.071
s_y = 0.062
```

```
sign_level = 0.05
```

The hypoteses are: - H_0 : $\mu_A - \mu_B \ge 0$ - H_A : $\mu_A - \mu_B < 0$

Degrees of freedom 29
Test statistics: -0.8480571253767827
P-value using cdf = 0.2017

Therefore, there is not sufficient evidence that the null hypotesis is true

1.6 9.3.26

```
[134]: # Population A
n = 10
x_bar = 7.76
s_x = 1.07

# Population B
m = 9
y_bar = 6.88
s_y = 0.62

alpha = 0.01 # = 1 - confidence level
```

```
print(("Confidence interval with \{:.2f\}\% confidence level: (-\omega)(\{:.
        \rightarrow4f})").format(((1-alpha)*100), mu + wing_span))
                return
           if lower bound:
                print(("Confidence interval with {:.2f}% confidence level: ({:.
        \rightarrow4f})(\emptyset)").format(((1-alpha)*100), mu - wing_span))
           # Default case: double bounded
           print(("Confidence interval with {:.2f}% confidence level: ({:.4f})({:.
        \rightarrow4f})").format(((1-alpha)*1, mu - wing_span, mu + wing_span)))
       def degrees_of_freedom(n, m, s_x, s_y):
           nu = ((s_x**2/n + s_y**2/m)**2) / (s_x**4 / (n**2*(n-1)) + s_y**4 / (n**2*(n-1)))
        \rightarrow (m**2*(m-1)))
           return math.floor(nu)
       nu = degrees_of_freedom( n, m, s_x, s_y)
       print("Degrees of freedom: ", nu)
       t_stat = T_statistic(x_bar, y_bar, s_x, s_y, n, m, mu_difference = 0)
       diff = x_bar - y_bar
       ws = wing_span(alpha, nu, n, m, s_x, s_y)
       print("a)")
       print_confidence interval(alpha, diff, ws, lower_bound = True)
       ws = wing_span(0.05, nu, n, m, s_x, s_y)
       print("b)")
       print_confidence_interval(0.05, diff, ws, lower_bound = True)
      Degrees of freedom: 14
      Confidence interval with 99.00% confidence level: (-0.1606)(\(\omega\))
      Confidence interval with 95.00% confidence level: (0.1817)(0))
      The value of c increases, since the confidence level has decreased: we have a "tighter" constraint
      The hypoteses are: - H_0: \mu_A \leq \mu_B - H_A: \mu_A > \mu_B
[141]: # Calculate the t-statistics
       def T_statistic(x_bar, y_bar, s_x, s_y, n, m, mu_difference = 0):
           return ( (x_bar - y_bar - mu_difference) / math.sqrt(s_x**2/n + s_y**2/m))
       t_stat = T_statistic(x_bar, y_bar, s_x, s_y, n, m)
```

print("T statistic: ", t_stat)

print("Size alpha = 0.01: ", t.ppf(1-0.01, nu))

```
print(("P-value using sf (1-cdf) = {:.4f}").format(t.sf(t_stat, nu)))
```

T statistic: 2.2194985542975125 Size alpha = 0.01: 2.624494067560231 P-value using sf (1-cdf) = 0.0217

We can accept the null hypothesis since $t \le t_{0.01,14}$. The p value is $P(t_{14} \ge 2.22) = 0.0217$

$1.7 \quad 9.7.10$

```
[150]: # Population A
n = 48
x_bar = 432.7
s_x = 20.39

# Population B
m = 10
y_bar = 403.5
s_y = 15.62

alpha = 0.01 # = 1 - confidence level
```

The hypoteses are: - $H_0: \mu_A \leq \mu_B$ - $H_A: \mu_A > \mu_B$

16 T statistics: 5.07845732358246 P-value using sf (1-cdf) = 0.0001

Given the low p-value we can accept the alternative hypothesis and conclude that the new driving route will be quicker on average than the old one

1.8 9.7.14

```
[195]: data=pd.read_excel("DS9.6.7.xls")
  dat_A= data['Procedure 1']
  dat_B= data['Procedure 2']
  # We create the difference distribution
  z = dat_A - dat_B
```

The hypoteses are: - $H_0: \mu_A = \mu_B$ - $H_A: \mu_A \neq \mu_B$

```
[197]: print("Mean =", z.mean())
    print("Standard deviation = ", math.sqrt(z.var()))
    t_stat = 3*z.mean() /math.sqrt(z.var())
    print("t statistics: ", t_stat)
    print(("P-value using sf (1-cdf) = {:.4f}").format(2 * t.sf(t_stat, 8)))
```

```
Mean = -0.02222222222222143

Standard deviation = 0.5911382616989399

t statistics: -0.1127767748869198

P-value using sf (1-cdf) = 1.0870
```

Given the high p-value, there is no evidence to conclude there is any difference between the two procedures

1.9 9.7.20

```
[169]: data=pd.read_excel("DS9.6.11.xls")
  dat_A= data['Joystick Design 1']
  dat_B= data['Joystick Design 2']
  # We create the difference distribution
  z = dat_A - dat_B
```

The hypoteses are: - $H_0: \mu_A = \mu_B$ - $H_A: \mu_A \neq \mu_B$

```
[203]: print("Mean =", z.mean())
    print("Standard deviation = ", math.sqrt(z.var()))
    t_stat = 3*z.mean() /math.sqrt(z.var())
    print("t statistics: ", t_stat)
    print(("P-value using sf (1-cdf) = {:.4f}").format(2 * t.sf(abs(t_stat), 8)))
```

```
Mean = -0.02222222222222143

Standard deviation = 0.5911382616989399

t statistics: -0.1127767748869198

P-value using sf (1-cdf) = 0.9130
```

We can say that there is some evidence, although not overwhelming, that the two joysticks result in different error rate measurements

Confidence interval with 0.99% confidence level: (-0.0151)(0.0565)

1.10 9.7.22

```
[205]: data=pd.read_excel("DS9.6.13.xls")
  dat_A= data['Sphygmomanometer']
  dat_B= data['Finger Monitor']
  # We create the difference distribution
  z = dat_A - dat_B
```

The hypoteses are: - $H_0: \mu_A = \mu_B$ - $H_A: \mu_A \neq \mu_B$

```
[206]: print("Mean =", z.mean())
    print("Standard deviation = ", math.sqrt(z.var()))
    t_stat = math.sqrt(15)*z.mean() /math.sqrt(z.var())
    print("t statistics: ", t_stat)
    print(("P-value using sf (1-cdf) = {:.4f}").format(2 * t.sf(abs(t_stat), 14)))
```

```
Mean = 0.4
Standard deviation = 1.9566735620873066
t statistics: 0.7917484901417817
P-value using sf (1-cdf) = 0.4417
```

We can conclude from the p-value that there is enough evidence to state that the means of measurement of the two instruments are different