

Statistics and data analysis II Week13

[Hypothesis test for mean]

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# INIAD

#### Lecture plan

Week1: Introduction of the course and some mathematical preliminaries

Week2: Overview of statistics, One dimensional data(1): frequency and histogram

Week3: One dimensional data(2): basic statistical measures

Week4: Two dimensional data(1): scatter plot and contingency table

Week5: Two dimensional data(2): correlation coefficients, simple linear regression and concepts of Probability /

Probability(1):randomness and probability, sample space and probabilistic events

Week6:Probability(2): definition of probability, additive theorem, conditional probability and independency

Week7:Review and exam(i)

Week8: Random variable(1): random variable and expectation

Week9: Random variable(2): Chebyshev's inequality, Probability distribution(1):binomial and

Poisson distributions

Week10: Probability distribution(2): normal and exponential distributions

Week11: From descriptive statistics to inferential statistics -z-table and confidwncw interval-

Week12: Hypothesis test(1) -Introduction, and distributions of test statistic (t-distribution)-

Week13: Hypothesis test(2) -Test for mean-

Week14: Hypothesis test(3) -Test for difference of mean-





# 2. Hypothesis test for mean

# Agenda

- Hypothesis test for mean
- In case population S.D. is known
- In case population S.D. is unknown
- Exercises



#### 2-1. Hypothesis test for mean



#### Hypothess test for mean

 In the inferential statistics, we estimate the characteristics of the population from observed samples.

 Especially, if we validate the value of the population mean, it is called as the hypothesis test for mean.

#### Usage

 By using samples, validate whether the product specification is correctly applied.

#### Take care...

- 1) Population variance is known? Unknown?
  - → If known, apply z-dist., otherwise, t-dist.
- 2Two-sided? One -sided?
  - → The p-value python returns depends on cases.
    Is it two-sided p-val.? Or one-sided?
- (3) The row data is given? Just some statistical indicators (sample mean, unbiased SD) are given?



2-2. Hypothesis test for mean (In case the population S.D. is known)



# Hypothesis test for mean 1 (In case the population S.D. is known)

- Let the population mean and S.D. of a normal distribution be  $\mu$  and  $\sigma$ , respectively. Then, the n samples extracted from them also follows the normal distribution.
  - $\Box$  The sample mean remains as  $\mu$ , but the S.D. of the samples reduces to  $\frac{\sigma}{\sqrt{n}}$ 
    - 95% C.I. of the namal distribution:

$$-1.96 \le \frac{x - \mu}{\frac{\sigma}{\sqrt{n}}} \le 1.96$$





- Hypothesis test for mean 1
- (In case the population S.D. is known)
  - •In case  $\sigma^2$  is known, we use the following quantity as the test statistic:

$$t = \frac{x - \mu}{\frac{\sigma}{\sqrt{n}}}$$

•Then, what is the p.d.f. of the test statistic?

N(0,1)!





- Hypothesis test for mean 1
- (In case the population S.D. is known)
  - vi) For the significance level a, set the rejection region R that satisfies

$$P(T(X_1,X_2,...,X_N) \in R) = a.$$

- •In case of the two-sided test, it's out of the upper / lower 2.5-percentiles (=±1.96) of z-distribution.
- In case of the two-sided test, it's out of the upper / lower 5-percentiles (=1.64).





# Example 1

In a certain maker of a part (named as "M") of computer, its diameter is described as 1.54[cm] in its product specification. In a certain sample survey, they extracted 8 samples randomly, and observed the following data of measured diameter [cm]:

1.53 1.57 1.54 1.57 1.53 1.55 1.56 1.53

It's known that the population variance is  $\sigma^2$ =0.0001. Then, can you say that this part follow its product specification? Test with the significance level of 5%.



#### Flow of hypothesis test

- i) Set the population ( Similar to confidence interval)
- ii) Set the null hypothesis H<sub>0</sub>.
- iii) Extract samples  $x_1, x_2, ..., x_N$  from the population.
- iv) Find a statistics  $T(x_1,x_2,...,x_N)$  from the sample above.
- v) Calculate the probability density of the statistics  $T(X_1, X_2, ..., X_N)$  for r.v.s  $X_1, X_2, ..., X_N$ .





#### Flow of hypothesis test

vi) For a certain significance level a, find a region R.
 where
 DITIV V

$$P(T(X_1,X_2,...,X_N) \in R) = a$$

holds(This region R is called as the critical region)

• vii) If  $T(x_1, x_2, ..., x_N) \in \mathbb{R}$ , reject the null hypothesis  $H_0$  / otherwise,  $H_0$  cannot be rejected.





Flow of hypothesis test.

• Null hypothesis H<sub>0</sub> "The diameter of part "M" is 1.54cm"

Alternative hypothesis  $H_1$ :" The diameter of part "M" is not 1.54cm"

(No good with too large nor too small results.)

- Find the test statistic  $T(x_1, x_2, ..., x_N)$ .
- •Since the population variance is known, we use

$$t = \frac{x - \mu}{\frac{\sigma}{\sqrt{n}}}$$



• Find the p.d.f. of the test statistic t.

• If X follows  $N(\mu, \sigma^2)$ , then

$$t = \frac{\bar{x} - \mu}{\frac{\sigma}{\sqrt{n}}}$$

follows N(0,1).

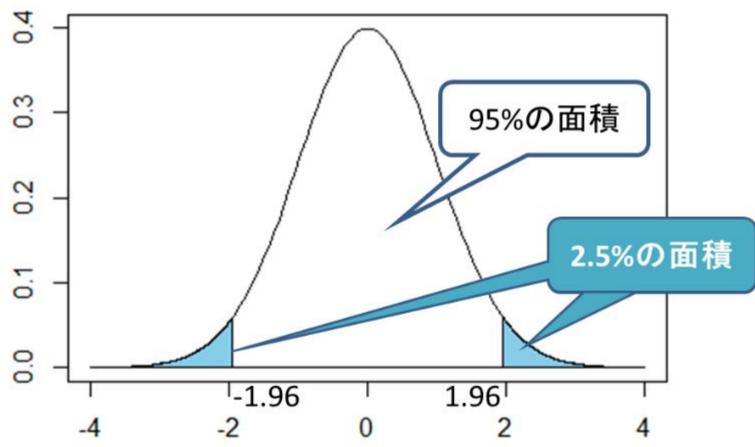
- Find the p.d.f. of the test statistic t.
  - $\Rightarrow$  z-distribution.



- Determine the rejection region.
- ⇒ Two-sided test from the form of the alternative hypothesis.

The rejection region is as follows (out of upper /lower2.5-percentiles of z-distribution).

• |t|>1.96 is the rejection region.





• Find the value of  $T(x_1, x_2, ..., x_N)$ :

$$\bar{x} = \frac{1.53 + 1.57 + \dots + 1.53}{8} = 1.5475$$

$$\mu = 1.54$$
  $\sigma = \sqrt{0.0001} = 0.01$ 

Thus, we have

$$t = \frac{1.5475 - 1.54}{\frac{0.01}{\sqrt{8}}} = 2.121$$

Belong to the rejection region |z|>1.96!



Reject H0 under the significance level of 5%. Employ the alternative hypothesis.

That is, "the diameter of the part M is not 1.54cm."



# Example (1) [answer]: Using Python

```
import numpy as np
from scipy import stats
from scipy.stats import norm
X = np.array([1.53, 1.57, 1.54, 1.57, 1.53, 1.55, 1.56, 1.53])
mu 0=1.54
avg = X.mean()
std = 0.01#X. std()
N=X.size
#print sample mean.
print(avg)
z = (avg - mu 0) / (std / np.sqrt(N))
#print z-value.
print(z)
p = norm.cdf(-np.abs(z), 0, 1) * 2
print(p)
```

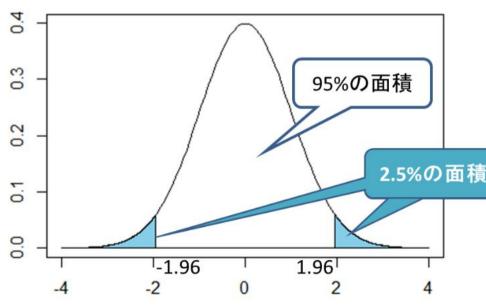


# Example (1) [answer]: Using Python

```
import numpy as np
from scipy import stats
from scipy.stats import norm
X = np.array([1.53, 1.57, 1.54,1.57, 1.53, 1.55, 1.56, 1.53])
mu 0=1.54
                                                 Null
avg = X.mean()
                                                 hypothesis
std = 0.01#X. std()
N=X.size
#print sample mean.
print(avg)
z = (a \lor g - mu_0) / (std / np.sqrt(N))
                                                                     Z-statistic
#print z-value.
print(z)
p = norm.cdf(-np.abs(z), 0, 1) * 2
print(p)

    To find the p-value, you should make it

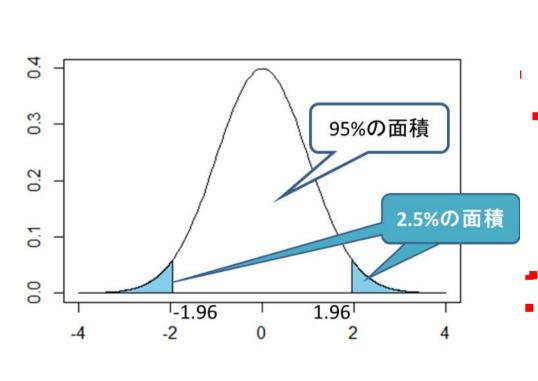
                               twice
                               in case of the two-sided test.
```





### Example (1) [answer]: Using Python

[Output]



```
import numpy as np
from scipy import stats
from scipy.stats import norm
X = np.array([1.53, 1.57, 1.54,1.57, 1.53, 1.55, 1.56, 1.53]) Null
                                                                   hypothesis
avg = X.mean()
std = 0.01#X. std()
N=X.size
#print sample mean.
print(avg)
z = (avg - mu_0) / (std / np.sqrt(N))
#print z-value.
print(z)
                                                                         Z-statistic
p = norm.cdf(- np.abs(z), 0, 1) * 2
print(p)
1.5475
2.1213203435596606
0.033894853524687726
                                                              P-val.<5%
                                                              In the rejection
                                                              region.
```

#### Answer.

"The diameter is not 1.54cm"

or

"Reject HO."

\*Assuming you can answer H0 correctly!



2-3. Hypothesis test for mean (In case population S.D. is unknown)



# Test for mean 2 (population SD is unknown)

- In method ①, the population SD  $\sigma$  was known What about  $\sigma$  is unknown . ?
- ullet You should replace  $\sigma$  by the unbiased SD, S.

$$t = \frac{\bar{x} - \mu}{\frac{S}{\sqrt{N}}}$$

Follows the t-distribution of df = (N-1)





#### Test for mean 2 (population SD is unknown)

• vi) For a, find a rejection region that meets  $P(T(X_1,X_2,...,X_N) \in R) = a.$ 

- •For two-sided test, out of upper/ lower 2.5percentiles(= $\pm t_{N-1}$ (0.025))
- •For one-sided test, above the upper 5-percentile  $(=t_{N-1}(0.05))$  or below the lower 5-percentile  $(=-t_{N-1}(0.05))$

# Example 2

In a certain factory, they make a component of a compter, named A. After we measured the length of 10 samples of this A, the mean and unbiased variance were 7.2[cm] and 0.04, respectively.

If the length of A follows the normal distribution, can we state that the length of A is 7.0cm on average? Conduct the hypothesis test with the significance level of 5%.



#### Flow of hypothesis test

- i) Set the population ( Similar to confidence interval)
- ii) Set the null hypothesis H<sub>0</sub>.
- iii) Extract samples  $x_1, x_2, ..., x_N$  from the population.
- iv) Find a statistics  $T(x_1,x_2,...,x_N)$  from the sample above.
- v) Calculate the probability density of the statistics  $T(X_1, X_2, ..., X_N)$  for r.v.s  $X_1, X_2, ..., X_N$ .





#### Flow of hypothesis test

vi) For a certain significance level a, find a region R.
 where
 DITIV V

$$P(T(X_1,X_2,...,X_N) \in R) = a$$

holds(This region R is called as the critical region)

• vii) If  $T(x_1, x_2, ..., x_N) \in \mathbb{R}$ , reject the null hypothesis  $H_0$  / otherwise,  $H_0$  cannot be rejected.



### Example 2 [Answer]

Flow of hypothesis test:

- Null hypothesis H<sub>0</sub>:"The length of A is 7.0cm."
  - Alternative hypothesis H<sub>1</sub>:"The length of A is not 7.0cm."
- Find the test statistic  $T(x_1, x_2, ..., x_N)$ .
- Since the population variance is unknown, we take

$$t = \frac{x - \mu}{\frac{S}{\sqrt{N}}}$$







• Find the p.d.f. of the test statistic t.



#### Example 2 [Answer]

• If X follows  $N(\mu, \sigma^2)$ , then, the test statistic

$$t = \frac{X - \mu}{\frac{s}{\sqrt{n}}}$$

follows the t-distribution with df=(n-1). Here,

$$s = \sqrt{\frac{(X_1 - \bar{X})^2 + (X_2 - \bar{X})^2 + \dots + (X_n - \bar{X})^2}{n - 1}}$$

# Example 2 [Answer]

- Find the p.d.f. of the test statistic t.
- ⇒t-distribution with df=9.

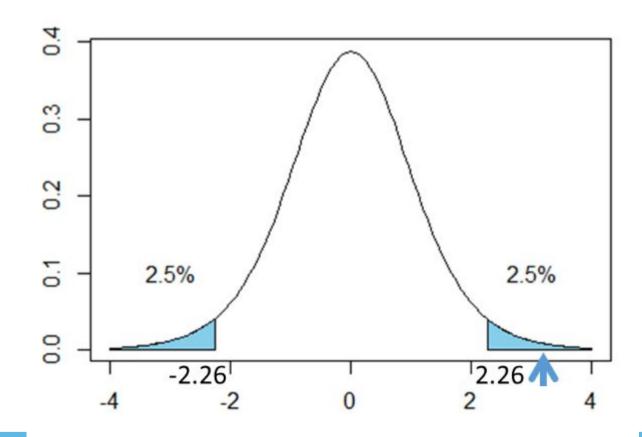


- Determine the rejection region.
  - ⇒ Two-sided test.

The rejection region is outside of upper/lower 2.5-percentiles of the t-distribution.

$$t_9(0.05/2) = 2.26$$

• The region |t| > 2.26 is the rejection region.

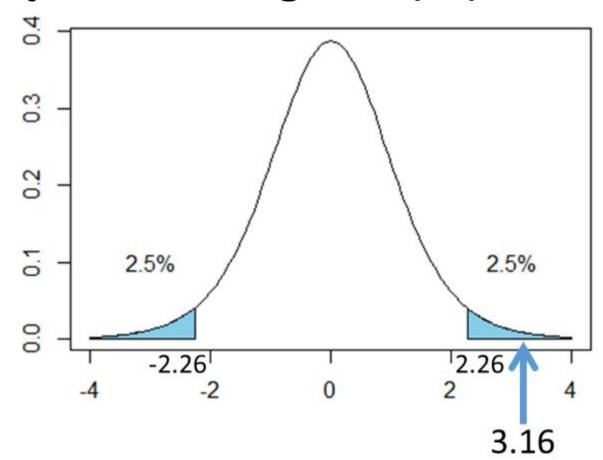




• Find the value of the test statistic  $T(x_1, x_2, ..., x_N)$ :

$$t = \frac{7.2 - 7}{\frac{0.2}{\sqrt{10}}} = 3.16$$

Belongs to the rejection region | t | >2.26 !







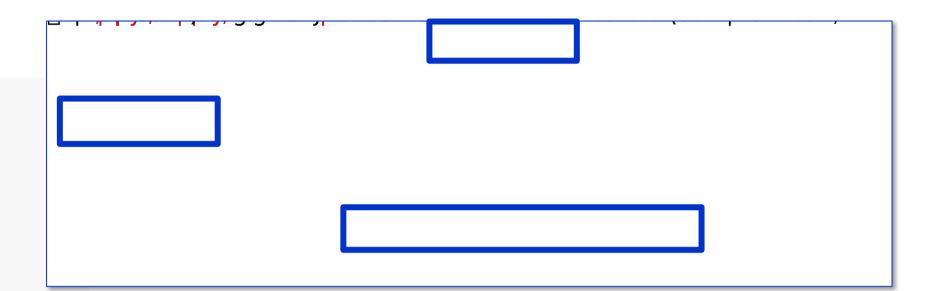
Reject H0 under the significance level of 5%.

"The length of A is not 7.0cm."



# Example 2 [Answer] Using Python

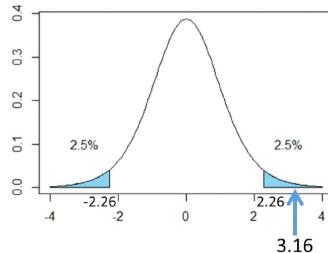
```
import numpy as np
from scipy import stats
from scipy.stats import t
mu 0=7.0
avg = 7.2
std = 0.2#X. std()
N=10 #X. size
#print sample mean.
stat t = (avg - mu \ 0) / (std / np.sqrt(N))
#print z-value.
print(stat t)
p = t.cdf(-np.abs(stat t), df=N-1) * 2
print(p)
```





# Example 2 [Answer] Using Python

```
import numpy as np
   from scipy import stats
   from scipy.stats import t
                                                       Null
  mu_0=7.0
                                                       hypothesis
   avg = 7.2
   std = 0.2#X. std()
   N=10 #X. size
   #print sample mean.
stat_t = (avg -mu_0)/ (std / np.sqrt(N))
                                                        Test statistic
   #print z-value.
```



print(stat\_t)
p = t.cdf(-np.abs(stat\_t), df=N-1) \* 2
print(p)

 To find the p-value, you should make it twice

in case of the two-sided test.

#### Answer

"The diameter is not 7.0cm /significantly different from 7.0cm"

Or

"Reject HO"

\*Assuming you can answer H0 correctly!



Hypothesis test for mean (In case the population S.D. is unknown / one-sided test)

# Example 3

A certain maker states that the lifetime of their light bulb is 2000 hours. To validate this statement, we bought 15 samples and tested their lifetime.

Then, the mean and unbiased S.D. were 1900 and 150 hours, respectively.

If the lifetime follows the normal distribution, can we say that the statement of this maker is correct?

Conduct the hypothesis test with the significance level of 5%.



#### Flow of hypothesis test

- i) Set the population ( Similar to confidence interval)
- ii) Set the null hypothesis H<sub>0.</sub>
- iii) Extract samples  $x_1, x_2, ..., x_N$  from the population.
- iv) Find a statistics  $T(x_1,x_2,...,x_N)$  from the sample above.
- v) Calculate the probability density of the statistics  $T(X_1, X_2, ..., X_N)$  for r.v.s  $X_1, X_2, ..., X_N$ .





#### Flow of hypothesis test

vi) For a certain significance level a, find a region R.
 where
 DITIV V

$$P(T(X_1,X_2,...,X_N) \in R) = a$$

holds(This region R is called as the critical region)

• vii) If  $T(x_1, x_2, ..., x_N) \in \mathbb{R}$ , reject the null hypothesis  $H_0$  / otherwise,  $H_0$  cannot be rejected.



• Null hypothesis H<sub>0</sub>:"The lifetime is 2000 hours."

Alternative hypothesis  $H_1$ :"The lifetime is less than 2000 hours." (No problem if it's longer)

- Find the test statistic  $T(x_1, x_2, ..., x_N)$ .
  - Since the population variance is unknown, we take:

$$t = \frac{\bar{x} - \mu}{\frac{S}{\sqrt{N}}}$$





• Find the p.d.f. of the test statistic t.



• If X follows  $N(\mu, \sigma^2)$ , then the test statistic

$$t = \frac{X - \mu}{\frac{s}{\sqrt{n}}}$$

follows the t-distribution with df=(n-1). Here,

$$s = \sqrt{\frac{(X_1 - \bar{X})^2 + (X_2 - \bar{X})^2 + \dots + (X_n - \bar{X})^2}{n - 1}}$$

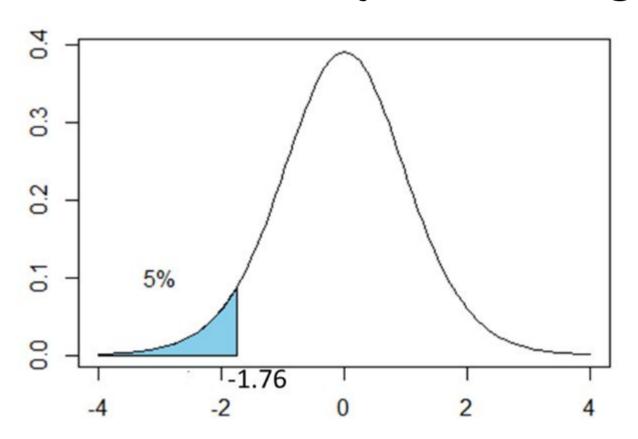
- Find the pd.f. of the test statistic t.
  - $\Rightarrow$  follows the t-distribution with df= 14.

- Determine the rejection region.
  - ⇒one-sided test.

The rejection region is left side of the lower 5-percentile of the ti-distribution.

$$t_{14}(0.05) = -1.76$$

•The region of t<-1.76 is the rejection region.

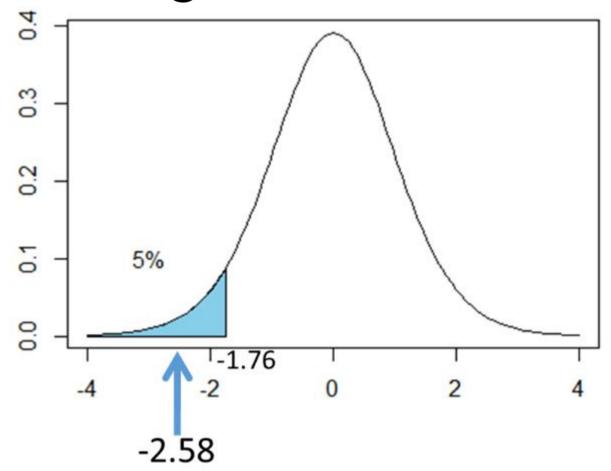




• Find the value of the test statistic  $T(x_1, x_2, ..., x_N)$ :

$$t = \frac{1900 - 2000}{\frac{150}{\sqrt{15}}} = -2.58$$

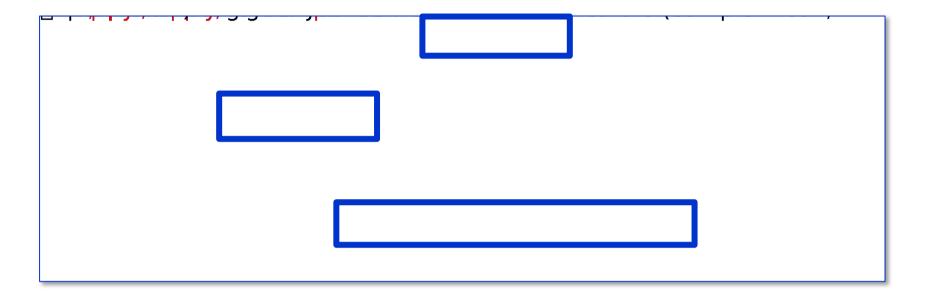
Belong to the rejection region t<-1.76!</li>



We reject H0 under the significance level of 5%.

Thus, "The lifetime of ther light bulb is shorter than 2000 hours. Their statement should be corrected."







# Example 3 [Answer] Using Python

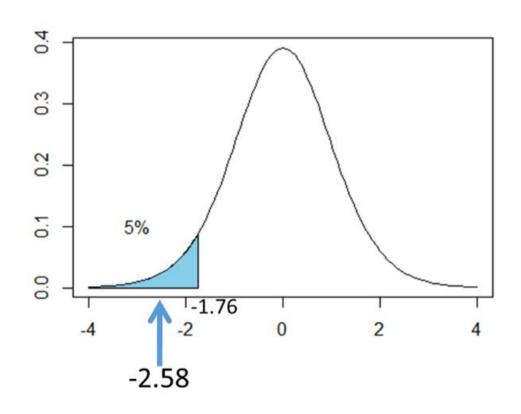
```
import numpy as np
  from scipy import stats
  from scipy.stats import t
                                                              Null
                                                              hypothesis
  avg = 1900
  std = 150 #X. std()
  N=15 #X. size
  #print sample mean.
                                                               Test statistic
stat_t = (avg -mu_0)/ (std / np.sqrt(N))
  #print z-value.
  print(stat_t)
p = t.cdf(-np.abs(stat_t), df=N-1)
                                             •Find the p-value. One-sided test this
  print(p)
                                             case!
                                             (No need to make twice).
```



# Example 3 [Answer] Using Python

[出力]

```
import numpy as np
from scipy import stats
from scipy.stats import t
mu_0 = 2000
avg = 1900
std = 150 #X. std()
N=15 #X. size
#print sample mean.
stat_t = (avg -mu_0) / (std / np.sqrt(N))
#print z-value.
print(stat t)
p = t.cdf(-np.abs(stat t), df=N-1)
print(p)
```



-2.581988897471611 0.01086262159930588

P-value < 5% In the rejection region.



#### Cautions in one-sided test

• In case of the one-sided test, if the sample mean belongs to the reverse side of the null hypothesis, we can promptly stop test by concluding "we cannot reject  $H_0$ ".

• XIn case of the prevous example, if the samples mean is less than 2000, we can promptly terminate the test.



#### Exercises



### Exercise(1)

In a certain farm, they developed a new fertilizer. Then, after we measured the their yield in 6 farms where they used this new fertilizer, we obtained the following data of the yield per unit area.

42.9 43.7 43.2 40.8 42.8 44.2 [kg]

The average yield with the conventional fertilizer per unit area was 41.4kg. Then, can we say that this new fertilizer improves the yield? Do the hypothesis test with known S.D.  $\sigma$ =3.5[kg] under the significance level of 5%.

## Exercise 1 [Answer]

Let us denote the average yield with the new fertilizer as  $\mu$ .

Null hypothesis  $H_0$ :" $\mu$ =41.4[kg]"

Alternative hypothesis H<sub>1</sub>:" µ>41.4[kg]"

Since the population variance is known, we take:

$$t = \frac{\bar{x} - \mu}{\frac{\sigma}{\sqrt{n}}}$$



We have:  $\bar{x} = 42.93$ 

Thus,

$$t = \frac{42.93 - 41.4}{\frac{3.5}{\sqrt{6}}} = 1.071$$

Does not satisfy t>1.64. It does not belong to the significance level. So we can't reject H<sub>0</sub>

⇒ "We can't say that the new fertilizer has a significant effect"



# Exercise 1 [Answer] Using Python

```
#Exercise 4.
import numpy as np
from scipy import stats
from scipy.stats import norm
X = np.array([42.9, 43.7, 43.2, 40.8, 42.8, 44.2])
mu 0=41.4
a \vee g = X.mean()
std = 3.5 #X. std()
N=X.size
#print sample mean.
print (avg)
z = (avg - mu 0) / (std / np.sqrt(N))
#print z-value.
print(z)
p = norm.cdf(-np.abs(z), 0, 1)
print(p)
```





# Exercise 1 [Answer] Using Python

```
#Exercise 4.
import numpy as np
from scipy import stats
from scipy.stats import norm
X = np.array([42.9, 43.7, 43.2, 40.8, 42.8, 44.2])
mu 0=41.4
avg = X.mean()
std = 3.5 #X. std()
N=X.size
#print sample mean.
print (avg)

    Find the p-value. One-sided test this

z = (avg - mu 0) / (std / np.sqrt(N))
                                                     case!
#print z-value.
                                                     (No need to make twice).
print(z)
p = norm.cdf( -np.abs(z), 0, 1)
print(p)
```

# Exercise 1 [Answer] Using Python

#### [Output]

```
#Exercise 4.
import numpy as np
from scipy import stats
from scipy.stats import norm
X = np.array([42.9, 43.7, 43.2, 40.8, 42.8, 44.2])
mu_0=41.4
avg = X.mean()
std = 3.5 #X. std()
N=X.size
#print sample mean.
print (avg)
z = (avg - mu_0) / (std / np.sqrt(N))
#print z-value.
print(z)
p = norm.cdf(-np.abs(z), 0, 1)
print(p)
```

```
42.93333333333334
```

- 1.0731097920764434
- 0.14161092902072725



# Exercise 2

In a certain maker of a part (named as "M") of computer, its diameter is described as 1.54[cm] in its product specification. In a certain sample survey, they extracted 8 samples randomly, and observed the following data of measured diameter [cm]:

1.53 1.57 1.54 1.57 1.53 1.55 1.56 1.53

The population variance is unknown. Then, can you say that this part follow its product specification? Test with the significance level of 5%.

# Exercise 2 [Answer]

Null hypothesis  $H_0$ :" $\mu$ =1.54[cm]"

Alternative hypothesis H₁:" µ≠1.54[cm]"

Since the population variance is known, we take:

$$t = \frac{\bar{x} - \mu}{\frac{S}{\sqrt{N}}}$$

#### Exercise 2 [Answer]

We have  $\bar{x} = 1.547$  and the unbiased S.D. is

$$S^{2} = \frac{(1.53 - 1.5475)^{2} + (1.57 - 1.5475)^{2} + \dots + (1.53 - 1.5475)^{2}}{8 - 1} = 0.000307$$

$$t = \frac{1.5475 - 1.54}{\frac{0.0175}{\sqrt{8}}} = 1.212$$

The upper 2.5-percentile of the t-distribution with df=(8-1=7) is

$$t_7(\frac{0.05}{2}) = 2.365$$

Since | t | < 2.365, it doesn't belong to the rejection region. We cannot reject H0.

"We cannot recognize the different under the significance level of 5%."





# Exercise 2 [Answer] Using Python

```
#Exercise 5.
import numpy as np
from scipy import stats
from scipy.stats import t
X = np.array([1.53, 1.57, 1.54, 1.57, 1.53, 1.55, 1.56, 1.53])
mu 0=1.54
avg = X.mean()
                                                                 To find test statistic, you should make it twice
std = np.std(X, ddof=1)
                                                                 in case of the two-sided test.
N=X.size
#print sample mean.
#print(avg)
stats t = (avg - mu \ 0) / (std / np.sqrt(N))
#print stats_t-value.
print(stats_t)
printistats_t)
p = t.cdf( -np.abs(stats_t), df=N-1) * 2
print(p)
```



# Exercise 2 [Answer] Using Python

```
[Output #Exercise 5.
              import numpy as np
              from scipy import stats
              from scipy.stats import t
              X = np.array([1.53, 1.57, 1.54, 1.57, 1.53, 1.55, 1.56, 1.53])
              mu 0=1.54
              a \vee g = X.mean()
              std = np.std(X, ddof=1)
              N=X.size
              #print sample mean.
              #print(avg)
              stats_t = (avg -mu_0) / (std / np.sqrt(N))
              #print stats_t-value.
              print(stats_t)
              p = t.cdf(-np.abs(stats_t), df=N-1) * 2
              print(p)
                                                                    P-value >5%
```

Cannot reject the null hypothesis.

1.2104198771789023

0.2653980394260665



# Exercise 3

A catalog of a climbing shop I states that the breaking strength of their 15mm rope is 4500kg.

Now, after we conducted the sample measurement of its strength by using 50 samples, the mean and unbiased S.D. were 4450[kg] and 120[kg], respectively.

Then, can we state that the stated strength 4500[kg] is satisfied on average? Conduct the hypothesis test with the significance level of 5%.

## Exercise 3 [Answer]

Let  $\mu$  denote its mean.

Null hypothesis  $H_0$ :" $\mu$ =4500"

Alternative hypothesis H<sub>1</sub>:"µ<4500" (No problem in case it's larger.)

Since the population variance is unknown, we take:

$$t = \frac{\bar{x} - \mu}{\frac{S}{\sqrt{N}}}$$

# Exercise 3 [Answer]

Sample mean and unbiased S.D. were 4450 and 120, respectively. So,

$$t = \frac{4450 - 4500}{\frac{120}{\sqrt{50}}} = -2.946$$

This follows the t-distribution with df= 50-1=49. But its lower 5-percentile is  $-t_{49}(\underline{0.05})_{88}$ .

Now, since t < -1.68, it belongs to the regnificance region, and H0 is rejected. We can say "The mean of samples are different from the one stated in their catalog under the significance level of 5%."



# Exercise 3 [Answer] Using Python

```
#Exercise 6.
import numpy as np
from scipy import stats
from scipy.stats import t
\#X = np. array([1.53, 1.57, 1.54, 1.57, 1.53, 1.55, 1.56, 1.53])
mu 0=4500
avg =4450# X. mean()
std = 120 \# np. std(X, ddof=1)

    Find the p-value. One-sided test this

N=50 #X. size
stats_t = (avg -mu_0)/(std / np.sqrt(N))
                                                      case!
#print stats_t-value.
                                                      (No need to make twice)
print(stats_t)
p = t.cdf( -np.abs(stats_t), df=N-1)
```

-2.946278254943948

0.0024555744280253798



# Exercise 3 [Answer] Using Python

```
[Output]<sub>e 6</sub>
   import numpy as np
   from scipy import stats
   from scipy.stats import t
   \#X = np. array([1.53, 1.57, 1.54, 1.57, 1.53, 1.55, 1.56, 1.53])
   mu 0=4500
   avg =4450# X. mean()
   std = 120 \# np. std(X, ddof=1)
   N=50 #X. size
   stats_t = (avg -mu_0)/(std/np.sqrt(N))
   #print stats_t-value.
   print(stats t)
   p = t.cdf(-np.abs(stats t), df=N-1)
                                                             P-value<5%
   print(p)
                                                            Reject the null
 -2.946278254943948
                                                             hypothesis.
 0.0024555744280253798
```



For a certain product, we measured its diameter of 5 samples, and observed:

36.3, 35.7, 35.9, 37.1, 36.1 [mm].

The spec of this product states that it should be 35.5m. Then, check if we can state that the actual diameter of this product is significantly larger than the spec or not.

Assume the normality of population and apply the hypothesis test for mean under the significance level of 5%.



#### 演習4【解答】

```
import numpy as np
from scipy import stats
from scipy.stats import t
X=np.array([36.3, 35.7, 35.9, 37.1, 36.1])
mu_0=35.5
X_mean=X.mean()
X_sd=np.std(X,ddof=1)
N=X.size
stats_t = (X_mean -mu_0)/(X_sd/np.sqrt(N))
p_val = t.cdf( -np.abs(stats_t), df=N-1)
print("p-value is")
print(p_val)
if p_val<0.05:</pre>
   print("帰無仮説棄却")
else:
   print("帰無仮説棄却できない")
```

p-value is 0.020381913442855257 帰無仮説棄却

