# **Hypothesis Testing**

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Hypothesis testing is an act in statistics whereby an analyst tests an assumption regarding a population parameter. The methodology
employed by the analyst depends on the nature of the data used and the reason for the analysis.

#### **KEY TAKEAWAYS**

- Hypothesis testing is used to assess the plausibility of a hypothesis by using sample data.
- The test provides evidence concerning the plausibility of the hypothesis, given the data.
- Statistical analysts test a hypothesis by measuring and examining a random sample of the population being analyzed.

## **How Hypothesis Testing Works**

- In hypothesis testing, an analyst tests a statistical sample, with the goal of providing evidence on the plausibility of the null hypothesis.
- Statistical analysts test a hypothesis by measuring and examining a random sample of the population being analyzed. All analysts use a random population sample to test two different hypotheses: the null hypothesis and the alternative hypothesis.
- The null hypothesis is usually a hypothesis of equality between population parameters; e.g., a null hypothesis may state that the population mean return is equal to zero.
- The alternative hypothesis is effectively the opposite of a null hypothesis (e.g., the population mean return is not equal to zero). Thus, they are mutually exclusive, and only one can be true. However, one of the two hypotheses will always be true.

```
import matplotlib.pyplot as plt
import numpy as np

from scipy.stats import norm
from scipy.stats import t

In [9]:

def tscore(sample_size, sample_mean, pop_mean, sample_std):
    nume=sample_mean-pop_mean
    denom=sample_std/(sample_size)**0.5
    return nume/denom
```

# Understanding the Hypothesis Testing

#### Step - 1:

Alternate Hypothesis(Bold Claim):  $$H_1 $ \sim $\star , \$ Null Hypothesis (status Quo):  $$H_0 $ \sim \$ Null Hypothesis (status Quo):  $$H_0 $ \sim \$ 

Step - 2:

- Collect a sample of size n
- Calculate the Mean of the sample (\$\bar{x}\$)

Step - 3: Compute test statistic:

- If standard deviation is known: \$\$ z=\frac{\bar{x}-\mu}{\frac{\sigma}{\sgrt[2]{n}}} \$\$
- If standard deviation is not known:

\$ t=\frac{\bar{x}-\mu}{\frac{s}{\sqrt[2]{n}}}\$\$

step - 4: Decide Significance level \$\alpha\$. If the alpha is lower remember we need stronger evidence to reject Null hypothesis.

step - 5.1: Apply Decision Rule:

- If test statistic is Z score:
  - Two Tailed Z-test:

 $|z| > z_{\alpha}, \$  Tailed Z-Test:  $z > z_{\alpha}, \$  Tailed Z-Test:  $z > z_{\alpha}, \$  Tailed Z-Test:  $z > z_{\alpha}, \$ 

■ Left Tailed Z-Test:

\$ z < -z\_\frac{\alpha}{2}\, \implies \, Accept H\_1 or Reject H\_0 \$\$

- If test statistic is t-score:
  - Two tailed test:
    - \$  $|t| > t_{n-1,\alpha/2} \in Accept H_1 or Reject H_0 $$$
  - Right Tailed t-test:

\$ t > t\_{n-1,\alpha} \implies Accept H\_1 or Reject H\_0 \$\$

■ Left tailed t-test:

 $t < t_{n-1,\alpha}$  \implies Accept H\_1 or Reject H\_0 \$\$

Step - 5.2 Compute P-Value \$\$ P(test statistics | H\_0) \$\$

• For Two tailed test:

\$\$ pvalue=2\*(1.0-cdf(test statistic)) \$\$

• For one tailed test:

\$\$ pvalue=(1.0-cdf(test statistic))\$\$ Now,

\$\$ if(pvalue<\alpha)\implies Accept H\_1 or Reject H\_0 \$\$

#### Practical Example-1

Q-1: Pista House selling Hyderabadi Chicken Dum biryani claims that each parcel packet has 500 grams of biryani (also mentioned on the label of packet). You are sceptic of their claims and believe that on average each packet does not contain 500 grams of biryani. How do you prove your claim?

#### Step 1:

Alternate Hypothesis (Bold Claim):

\$\$ H\_1: On \,an\, Average\, each\, Chicken\, dum\, biryani\, packet \,does\, not\, weigh\, 500 \,grams\$\$ \$\$\mu \neq 500\$\$ Null Hypothesis (status quo):

\$\$ H\_0: Pista\, House\, claims\, that\, each\, chickendum\, biryani \,packet \,weighs\, 500 grams \$\$ \$\$\mu = 500\$\$ step - 2:

Collect a sample of size n=12

\$\$ [500,510,505,498,476,520,521,480,510,504,550,469]\$\$

Compute sample mean \$\$\bar{x}=497.0 \$\$

Step - 3: compute Test Statistic:

since standard deviation is not provide we compute t score where,

\$\$ t=\frac{\left(\frac{s}{\sqrt{2}{n}}\right)=-0.5725114387470406\$\$}

Step - 4: Decide \$\alpha\$ or significance level

\$\$\alpha=0.95\$\$ step - 5.1: Two Tailed t-test:

\$\$ Reject\, H\_0\, if \, tscore\, >\, t\, critical \$\$ \$\$From\, the \,above\,condition \,we \,failed\,to\,reject\,Null\,Hypothesis\$\$

step - 5.2:Calculate P-value

\$\$Calculated \,P- value \,is\, :\, 0.20641847943818092 \$\$ \$\$if \,P -value > alpha\,: \,reject \,null\, hypothesis\$\$ \$\$ Hence,\,we\, failed\, to\, reject\, null\, hypothesis\$\$\$

```
sample mean=np.array(samples).mean()
print("sample mean:",sample_mean)
def sample std (samples, sample_mean):
    summation=0
    for i in range(len(samples)):
        summation=summation+(samples[i]-sample mean)**2
    return (summation/9)**0.5
sample std=sample std (samples, sample mean)
print("sample standard deviation:",sample_std)
sample mean: 465.3
sample standard deviation: 86.84859110991827
                                                                                                        In [4]:
confidence_level=0.95
alpha=1-confidence level
t_critical=t.ppf(1-alpha/2,df=9)
print(t critical)
2.2621571627409915
                                                                                                        In [5]:
sample_size=10
pop mean=500
                                                                                                        In [6]:
t =tscore(sample size, sample mean, pop mean, sample std)
print(t_)
-1.2634751284446715
                                                                                                        In [7]:
x min = 200
x max = 800
# Defining the sampling distribution mean and sampling distribution std
mean = pop mean
std = sample std / sample size**0.5
# Ploting the graph and setting the x limits
x = np.linspace(x min, x max, 100)
y = norm.pdf(x, mean, std)
plt.xlim(x min, x max)
plt.plot(x, y)
# Computing the left and right critical values (Two tailed Test)
t_critical_left = pop_mean + (-t_critical * std)
t_critical_right = pop_mean + (t_critical * std)
# Shading the left rejection region
x1 = np.linspace(x min, t critical left, 100)
y1 = norm.pdf(x1, mean, std)
plt.fill_between(x1, y1, color='orange')
# Shading the right rejection region
x2 = np.linspace(t_critical_right, x_max, 100)
y2 = norm.pdf(x2, mean, std)
plt.fill between(x2, y2, color='orange')
# Ploting the sample mean and concluding the results
plt.scatter(sample mean, 0)
plt.annotate("x_bar", (sample_mean, 0.0007))
```

```
Out[7]:
Text(465.3, 0.0007, 'x bar')
 0.014
 0.012
 0.010
 0.008
 0.006
 0.004
 0.002
                          x bar
 0.000
            300
                    400
                           500
                                   600
                                           700
                                                  800
    200
                                                                                                                   In [8]:
if(abs(t ) > t critical):
     print("Reject Null Hypothesis")
else:
     print ("Fail to reject Null Hypothesis")
Fail to reject Null Hypothesis
                                                                                                                   In [9]:
p \text{ value} = 2 * (1.0 - norm.cdf(np.abs(t)))
print("p value = ", p value)
if (p_value < alpha):</pre>
     print("Reject Null Hypothesis")
else:
     print("Fail to reject Null Hypothesis")
p value = 0.20641847943818092
Fail to reject Null Hypothesis
```

# Practical Example - 2

Q-2: You have developed a new Natural Language Processing Algorithms and done a user study. You claim that the average rating given by the users is greater than 4 on a scale of 1 to 5. How do you prove this to your client?

#### Step 1:

Alternate Hypothesis (Bold Claim):

 $$$ H_1: Average\, rating\, given\, by\, the \,users\, is\, greater\, than \,4 $$ $\mu>4$$ Null Hypothesis (status quo): $$ H_0: Average\, rating\, given\, by \,the \,users\, i, less thanorequalto\, 4 $$ $\mu\eq 4$$ step - 2:$ 

Collect a sample of size n=12

\$\$ [4,3,5,4,5,3,5,5,4,2,4,5,5,4,4,5,4,5,4,5] \$\$

Compute sample mean \$\$\bar{x}=4.25 \$\$

Step - 3: compute Test Statistic:

since standard deviation is not provide we compute t score where,  $\$  t=\frac{\bar{x}-\mu}{\frac{s}{\sqrt[2]{n}}}=1.3483997249264843 \$\$ Step - 4: Decide \$\alpha\$ or significance level \$\$\alpha=0.95\$\$ step - 5.1: one Tailed t-test:

• Right Tailed t-test: \$\$ t > t\_{n-1,\alpha} \implies Accept H\_1 or Reject H\_0 \$\$

 $\$From\, the \,above\,condition\,we \,failed\,to\,reject\,Null\,Hypothesis\,$ 

```
step - 5.2:Calculate P-value
```

 $\cline{Color:} $$Calculated \,P- value \,is\, :\, 0.17752985241215358 $$ $$if \,P-value > alpha\,: \,reject \,null\, hypothesis$$ $$Hence,\,we\, failed\, to\, reject\, null\, hypothesis$$$ 

In [10]:

```
# One Tail - Calculating the z-critical value
confidence_level = 0.95
```

```
alpha = 1 - confidence level
t critical = t.ppf(1 - alpha, df=19)
print(t critical)
1.729132811521367
                                                                                                        In [11]:
sample_size = 20
sample mean = 4.25
{\tt pop\_mean=4}
                                                                                                        In [12]:
samples=[4,3,5,4,5,3,5,5,4,2,4,5,5,4,4,5,4,5,4,5]
sample mean=np.array(samples).mean()
print("sample mean:", sample mean)
def sample_std_(samples, sample_mean):
    summation=0
    for i in range(len(samples)):
        summation=summation+(samples[i]-sample_mean)**2
    return (summation/len(samples))**0.5
sample_std=sample_std_(samples, sample_mean)
print("sample standard deviation:", sample std)
sample mean: 4.25
sample standard deviation: 0.82915619758885
                                                                                                        In [13]:
t =tscore(sample size, sample mean, pop mean, sample std)
print(t)
1.3483997249264843
                                                                                                        In [14]:
#plotting
x min = 2
x max = 6
# Defining the sampling distribution mean and sampling distribution std
mean = pop mean
std = sample_std / sample_size**0.5
# Ploting the graph and setting the x limits
x = np.linspace(x min, x max, 100)
y = norm.pdf(x, mean, std)
plt.xlim(x_min, x_max)
plt.plot(x, y)
# Computing the left and right critical values (Two tailed Test)
t critical left = pop mean + (-t critical * std)
t_critical_right = pop_mean + (t_critical * std)
# Shading the right rejection region
x2 = np.linspace(t_critical_right, x_max, 100)
y2 = norm.pdf(x2, mean, std)
plt.fill_between(x2, y2, color='orange')
# Ploting the sample mean and concluding the results
plt.scatter(sample mean, 0)
plt.annotate("x bar", (sample mean, 0.0007))
```

```
Out[14]:
Text(4.25, 0.0007, 'x bar')
 2.0
 1.0
 0.5
 0.0
        2.5
              3.0
                    3.5
                         4.0
                               4.5
                                     5.0
                                           5.5
   2.0
                                                6.0
                                                                                                                 In [15]:
if(abs(t ) > t critical):
     print("Reject Null Hypothesis")
else:
     print ("Fail to reject Null Hypothesis")
Fail to reject Null Hypothesis
                                                                                                                 In [16]:
p \text{ value} = 2 * (1.0 - norm.cdf(np.abs(t)))
print("p value = ", p value)
if (p_value < alpha):</pre>
     print("Reject Null Hypothesis")
else:
     print("Fail to reject Null Hypothesis")
p value = 0.17752985241215358
Fail to reject Null Hypothesis
```

## Practical Example - 3

Q-3: TATA has developed a better fuel management system for the SUV segment. They claim that with this system, on average the SUV's mileage is at least 15 km/litre?

#### Step 1:

Alternate Hypothesis (Bold Claim):

\$\$ H\_0: Average SUV's mileage is greater than or equal to 15km/hr \$\$ \$\$\mu\geq15km/hr\$\$ step - 2:

• Collect a sample of size n=10

\$\$ [14.08, 14.13, 15.65, 13.78, 16.26, 14.97, 15.36, 15.81, 14.53, 16.79, 15.78, 16.98, 13.23, 15.43, 15.46, 13.88, 14.31, 14.41, 15.76, 15.38] \$\$

• Compute sample mean \$\$\bar{x}=15.1 \$\$

Step - 3: compute Test Statistic:

since standard deviation is not provide we compute t score where,  $$$ t=\frac{x}-\max_{x}-\max_{\frac{2}{n}}=0.44748014931892083 $$ Step - 4: Decide $\alpha$ or significance level$  $$$\alpha=0.95$$ step - 5.1: one Tailed t-test:$ 

Right Tailed t-test:
 \$\$ t > t\_{n-1,\alpha} \implies Accept H\_1 or Reject H\_0 \$\$

 $\rm \from\, the \, above\, condition\, we \, failed\, to\, reject\, Null\, Hypothesis \$ 

```
step - 5.2:Calculate P-value
```

 $\$  value \,p- value \,is\, :\, 0.6545284174132713 \$\$ \$\$ if \,p -value > alpha\,: \,reject \,null\, hypothesis\$\$ \$\$ Hence,\,we\, failed\, to\, reject\, null\, hypothesis\$\$

In [17]:

```
confidence level = 0.95
alpha = 1 - confidence level
t_critical = t.ppf(1 - alpha,df=19)
print(t critical)
1.729132811521367
                                                                                                       In [18]:
samples=[14.08, 14.13, 15.65, 13.78, 16.26, 14.97, 15.36, 15.81, 14.53, 16.79, 15.78, 16.98, 13.23, 15.43,
sample_mean=np.array(samples).mean()
print("sample mean:", sample mean)
def sample_std_(samples, sample_mean):
    summation=0
    for i in range(len(samples)):
        summation=summation+(samples[i]-sample mean)**2
    return (summation/len(samples))**0.5
sample_std=sample_std_(samples, sample_mean)
print("sample standard deviation:", sample std)
sample mean: 15.099
sample standard deviation: 0.9994043225842081
                                                                                                       In [19]:
sample size = 20
sample mean = 15.1
pop mean = 15
pop std = 1
                                                                                                       In [20]:
t_=tscore(sample_size,sample_mean,pop_mean,sample_std)
print(t )
0.44748014931892083
                                                                                                       In [21]:
x \min = 13
x max = 17
mean = pop mean
std = pop std / (sample size**0.5)
x = np.linspace(x min, x max, 100)
y = norm.pdf(x, mean, std)
plt.xlim(x min, x max)
# plt.ylim(0, 0.03)
plt.plot(x, y)
t critical right = pop mean + (t critical * std)
x1 = np.linspace(t_critical_right, x_max, 100)
y1 = norm.pdf(x1, mean, std)
plt.fill_between(x1, y1, color='orange')
plt.scatter(sample mean, 0)
plt.annotate("x bar", (sample mean, 0.1))
```

```
Out[21]:
Text(15.1, 0.1, 'x bar')
 1.75
 1.50
 1.25
 1.00
 0.75
 0.50
 0.25
                             x_ba
 0.00
         13.5
              14.0
                    14.5
                          15.0
                                15.5
                                     16.0
                                           16.5
                                                 17.0
   13.0
                                                                                                                 In [22]:
if(abs(t ) > t critical):
     print("Reject Null Hypothesis")
else:
     print ("Fail to reject Null Hypothesis")
Fail to reject Null Hypothesis
                                                                                                                 In [23]:
p value = 2 * (1.0 - norm.cdf(abs(t)))
print("p value = ", p value)
if (p_value < alpha):</pre>
     print("Reject Null Hypothesis")
else:
     print("Fail to reject Null Hypothesis")
p value = 0.6545284174132713
Fail to reject Null Hypothesis
```

### Practical Example - 4

Q-4: You have developed a new Machine Learning Application and claim that on average it takes less than 100 ms to predict for any future datapoint. How do you convince your client about this claim?

```
Step 1:
```

Alternate Hypothesis (Bold Claim):

 $$$ H_1: Average\,MachineLearning\,Model\,takes \,lessthan\,100ms\,to\,predict \,any\,future \,data point 4 $$$ 

\$\$\mu < 100\$\$ Null Hypothesis (status quo):

 $$$ H_0: Average\,\$  Model\,takes \,greater than or equal to\,100ms\,to\,predict \,any\,future \,data point \$\$ \\$\mu\geq100\$\$ step - 2:

• Collect a sample of size n=10

\$\$[120,100,98,88,115,88,78,99,121,102] \$\$

• Compute sample mean \$\$\bar{x}=100.9 \$\$

Step - 3: compute Test Statistic:

since standard deviation is not provide we compute t score where, \$\$ t=\frac{\bar{x}-\mu}{\frac{s}{\sqrt[2]{n}}}= 0.20987632600435635\$\$ Step - 4: Decide \$\alpha\$ or significance level \$\$\alpha=0.99\$\$ step - 5.1: one Tailed t-test:

• Left tailed t-test:

\$ t < t\_{n-1,\lambda} \implies Accept H\_1 or Reject H\_0 \$ \$From\, the \,above\,condition \,we \,failed\,to\,reject\,Null\,Hypothesis\$\$

Step - 5.2:Calculate P-value

 $\C$  alculated \,P- value \,is\, :\, 0.8337641998328718 \$\$ \$\$if \,P -value > alpha\,: \,reject \,null\, hypothesis\$\$ \$\$ Hence,\,we\, failed\, to\, reject\, null\, hypothesis\$\$\$

In [ ]:

```
alpha = 1 - 0.99
t critical = t.ppf(1 - alpha, df=9)
print(t_critical)
2.8214379233005493
                                                                                                         In [5]:
samples=[120,100,98,88,115,88,78,99,121,102]
sample mean=np.array(samples).mean()
print("sample mean:",sample mean)
def sample_std_(samples, sample_mean):
    summation=0
    for i in range(len(samples)):
        summation=summation+(samples[i]-sample_mean)**2
    return (summation/len(samples))**0.5
sample_std=sample_std_(samples, sample_mean)
print("sample standard deviation:",sample std)
sample mean: 100.9
sample standard deviation: 13.560604706280618
                                                                                                        In [12]:
sample_size =10
sample mean=100.9
pop mean=100
pop_std=15
                                                                                                        In [13]:
t =tscore(sample size, sample mean, pop mean, sample std)
print(t)
0.20987632600435635
                                                                                                        In [17]:
x \min = 70
x max = 130
mean = pop_mean
std = pop_std / (sample_size**0.5)
x = np.linspace(x_min, x_max, 100)
y = norm.pdf(x, mean, std)
plt.xlim(x_min, x_max)
# plt.ylim(0, 0.03)
plt.plot(x, y)
t critical left = pop mean + (-t critical * std)
x1 = np.linspace(x_min, t_critical_left, 100)
y1 = norm.pdf(x1, mean, std)
plt.fill_between(x1, y1, color='orange')
plt.scatter(sample_mean, 0)
plt.annotate("x_bar", (sample_mean, 0.002))
```

```
Out[17]:
Text(100.9, 0.002, 'x_bar')
 0.08
 0.06
 0.04
 0.02
                           <u>x</u>_bar
 0.00
    70
           80
                  90
                          100
                                 110
                                        120
                                                130
                                                                                                               In [15]:
if(abs(t_) > t_critical):
    print("Reject Null Hypothesis")
else:
     print("Fail to reject Null Hypothesis")
Fail to reject Null Hypothesis
                                                                                                               In [16]:
p_{value} = 2 * (1.0 - norm.cdf(abs(t_)))
print("p_value = ", p_value)
if (p_value < alpha):</pre>
     print("Reject Null Hypothesis")
else:
     print("Fail to reject Null Hypothesis")
p value = 0.8337641998328718
```

In [ ]:

Fail to reject Null Hypothesis