

ORIGIN = 0

### One Sample t-Test

This test and associated descriptive statistics is designed to test hypotheses about the mean of a population with unknown variance.

#### Two Tail Case:

ZAR := READPRN("c:/DATA/Biostatistics/ZarEX7.2.txt") < Zar Example 7.2

X := ZAR<sup><1></sup>  
 n := length(X)      n = 12  
 X<sub>bar</sub> := mean(X)      X<sub>bar</sub> = -0.65  
 s := √Var(X)      s = 1.2523      s<sup>2</sup> = 1.5682

	0	1
0	1	1.7
1	2	0.7
2	3	-0.4
3	4	-1.8
4	5	0.2
5	6	0.9
6	7	-1.2
7	8	-0.9
8	9	-1.8
9	10	-1.4
10	11	-1.8
11	12	-2

ZAR =

#### Assumptions:

- Observed values X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub>, ... X<sub>n</sub> are a random sample from ~N(μ,σ<sup>2</sup>).
- Variance σ<sup>2</sup> of the population is *unknown*.      ^ Note: this test is reasonably robust for deviations from ~N(μ,σ<sup>2</sup>).

#### Hypotheses:

μ<sub>0</sub> := 0      < Let μ<sub>0</sub> = 0  
 H<sub>0</sub>: μ = μ<sub>0</sub>      < μ<sub>0</sub> is a specified value for μ  
 H<sub>1</sub>: μ ≠ μ<sub>0</sub>      < Two sided test

#### Test Statistic:

t :=  $\frac{X_{bar} - \mu_0}{\frac{s}{\sqrt{n}}}$       t = -1.7981      < t is the normalized distance between means X<sub>bar</sub> and μ<sub>0</sub>=0

#### Sampling Distribution:

If Assumptions hold and H<sub>0</sub> is true, then t ~t<sub>(n-1)</sub>

#### Critical Value of the Test:

α := 0.05      < Probability of Type I error must be explicitly set  
 C<sub>1</sub> := qt( $\frac{\alpha}{2}, n - 1$ )      C<sub>1</sub> = -2.201      < Critical values found by the inverse cumulative t function.  
 C<sub>2</sub> := qt( $1 - \frac{\alpha}{2}, n - 1$ )      C<sub>2</sub> = 2.201      Note: two critical values are found one for each tail of the distribution. Because the t distribution is symmetrical, the Critical values are of equal absolute magnitude.

#### Decision Rule:

IF t < C<sub>1</sub> or IF t > C<sub>2</sub>, THEN REJECT H<sub>0</sub>, OTHERWISE ACCEPT H<sub>0</sub>  
 IF |t| > |C|, THEN REJECT H<sub>0</sub>, OTHERWISE ACCEPT H<sub>0</sub>  
 t = -1.7981      C<sub>1</sub> = -2.201      C<sub>2</sub> = 2.201      < Therefore DO NOT Reject H<sub>0</sub>

#### Probability Value:

IF t ≤ 0 then: P := 2 · pt(t, n - 1)      P = 0.099637      < this value      < Rosner 2006 Eq. 7.11 p. 241  
 IF t > 0 then: P := 2 · (1 - pt(t, n - 1))      P = 1.9004

#### Confidence Interval for the Mean:

C := |C<sub>1</sub>|      C = 2.201  
 CI :=  $(X_{bar} - C \cdot \frac{s}{\sqrt{n}} \quad X_{bar} + C \cdot \frac{s}{\sqrt{n}})$       CI = (-1.4457 0.1457)      X<sub>bar</sub> = -0.65

**Two Tail Case:**

ZAR := READPRN("c:/DATA/Biostatistics/ZarEX7.1.txt")

< Zar Example 7.1

$$X := ZAR^{(1)}$$

$$n := \text{length}(X) \quad n = 25$$

$$X_{\text{bar}} := \text{mean}(X) \quad X_{\text{bar}} = 25.028$$

$$s := \sqrt{\text{Var}(X)} \quad s = 1.3418 \quad s^2 = 1.8004$$

	0	1
0	1	25.8
1	2	24.6
2	3	26.1
3	4	22.9
4	5	25.1
5	6	27.3
6	7	24
7	8	24.5
8	9	23.9
9	10	26.2
10	11	24.3
11	12	24.6
12	13	23.3
13	14	25.5

ZAR =

**Assumptions:**

- Observed values  $X_1, X_2, X_3, \dots, X_n$  are a random sample from  $\sim N(\mu, \sigma^2)$ .

- Variance  $\sigma^2$  of the population is *unknown*.

**Hypotheses:**

$$\mu_0 := 24.3 \quad < \text{Let } \mu_0 = 24.3$$

$$H_0: \mu = \mu_0 \quad < \mu_0 \text{ is a specified value for } \mu$$

$$H_1: \mu \neq \mu_0 \quad < \text{Two sided test}$$

**Test Statistic:**

$$t := \frac{X_{\text{bar}} - \mu_0}{\frac{s}{\sqrt{n}}} \quad t = 2.7128 \quad < t \text{ is the normalized distance between means } X_{\text{bar}} \text{ and } \mu_0$$

**Sampling Distribution:**

If Assumptions hold and  $H_0$  is true, then  $t \sim t_{(n-1)}$

**Critical Value of the Test:**

$$\alpha := 0.05 \quad < \text{Probability of Type I error must be explicitly set}$$

$$C_1 := \text{qt}\left(\frac{\alpha}{2}, n - 1\right) \quad C_1 = -2.0639 \quad < \text{Critical values found by the inverse cumulative t function.}$$

$$C_2 := \text{qt}\left(1 - \frac{\alpha}{2}, n - 1\right) \quad C_2 = 2.0639 \quad \text{Note: two critical values are found one for each tail of the distribution. Because the t distribution is symmetrical, the Critical values are of equal absolute magnitude.}$$

**Decision Rule:**

IF  $t < C_1$  or IF  $t > C_2$ , THEN REJECT  $H_0$ , OTHERWISE ACCEPT  $H_0$

IF  $|t| > |C|$ , THEN REJECT  $H_0$ , OTHERWISE ACCEPT  $H_0$

$$t = 2.7128 \quad C_1 = -2.0639 \quad C_2 = 2.0639 \quad < \text{Therefore REJECT } H_0$$

**Probability Value:**

$$\text{IF } t \leq 0 \text{ then: } P := 2 \cdot \text{pt}(t, n - 1) \quad P = 1.9879$$

$$\text{IF } t > 0 \text{ then: } P := 2 \cdot (1 - \text{pt}(t, n - 1)) \quad P = 0.012146 \quad < \text{this value}$$

< Rosner 2006 Eq. 7.11 p. 241

**Confidence Interval for the Mean:**

$$C := |C_1| \quad C = 2.0639$$

$$CI := \left( X_{\text{bar}} - C \cdot \frac{s}{\sqrt{n}} \quad X_{\text{bar}} + C \cdot \frac{s}{\sqrt{n}} \right) \quad CI = (24.4741 \quad 25.5819) \quad X_{\text{bar}} = 25.028$$



**Other One Tail Case (Upper Tail):**

ZAR := READPRN("c:/DATA/Biostatistics/ZarEX7.4.txt") < Zar Example 7.4

X := ZAR<sup><1></sup>

n := length(X) n = 8

X<sub>bar</sub> := mean(X) X<sub>bar</sub> = 45.2125

s := √Var(X) s = 1.6401 s<sup>2</sup> = 2.6898

ZAR =

	0	1
0	1	42.7
1	2	43.4
2	3	44.6
3	4	45.1
4	5	45.6
5	6	45.9
6	7	46.8
7	8	47.6

**Assumptions:**

- Observed values X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub>, ... X<sub>n</sub> are a random sample from ~N(μ,σ<sup>2</sup>).
- Variance σ<sup>2</sup> of the population is *unknown*.

**Hypotheses:**

- μ<sub>0</sub> := 45 < Let μ<sub>0</sub> = 45
- H<sub>0</sub>: μ ≤ μ<sub>0</sub> < μ<sub>0</sub> is a specified value for μ
- H<sub>1</sub>: μ > μ<sub>0</sub> < One sided test

**Test Statistic:**

t :=  $\frac{X_{bar} - \mu_0}{\frac{s}{\sqrt{n}}}$  t = 0.3665 < t is the normalized distance between means Xbar and μ<sub>0</sub>

**Sampling Distribution:**

If Assumptions hold and H<sub>0</sub> is true, then t ~t<sub>(n-1)</sub>

**Critical Value of the Test:**

- α := 0.05 < Probability of Type I error must be explicitly set
- C := qt(1 - α, n - 1) C = 1.8946 < Critical value is found by the inverse cumulative t function. Only one Critical value is found.

**Decision Rule:**

- IF t > C, THEN REJECT H<sub>0</sub>, OTHERWISE ACCEPT H<sub>0</sub>
- t = 0.3665 C = 1.8946 < Therefore DO NOT reject H<sub>0</sub>

**Probability Value:**

P := 1 - pt(t, n - 1) P = 0.3624 < Rosner 2006 p. 237

**Confidence Interval for the Mean:**

CI := X<sub>bar</sub> - |C| ·  $\frac{s}{\sqrt{n}}$  CI = 44.1139 to infinity X<sub>bar</sub> = 45.2125  
 ^ lower limit

## Prototype in R:

```
#t.test IN R
#TO FIND INFORMATION ON t.test()
?t.test

#ZAR EXAMPLE 7.2
ZAR=read.table("c:/DATA/Biostatistics/ZarEX7.2.txt")
ZAR
attach(ZAR)
X=RATwtchg
t.test(X,alternative="two.sided", mu=0,conf.level=0.95)
```

```
#ZAR EXAMPLE 7.1
ZAR=read.table("c:/DATA/Biostatistics/ZarEX7.1.txt")
ZAR
attach(ZAR)
X=crabtemp
t.test(X,alternative="two.sided", mu=24.3,conf.level=0.95)
```

```
#ZAR EXAMPLE 7.3
ZAR=read.table("c:/DATA/Biostatistics/ZarEX7.3.txt")
ZAR
attach(ZAR)
X=wtchg
t.test(X,alternative="less", mu=0,conf.level=0.95)
#NOTE: R REPORTS A ONE-WAY CI
```

```
#ZAR EXAMPLE 7.4
ZAR=read.table("c:/DATA/Biostatistics/ZarEX7.4.txt")
ZAR
attach(ZAR)
X=distime
t.test(X,alternative="greater", mu=45,conf.level=0.95)
#NOTE: R REPORTS A ONE-WAY CI
```

One Sample t-test

```
data: X
t = -1.7981, df = 11, p-value = 0.09964
alternative hypothesis: true mean is not equal to 0
95 percent confidence interval:
-1.4456548 0.1456548
sample estimates:
mean of x
-0.65
```

One Sample t-test

```
data: X
t = 2.7128, df = 24, p-value = 0.01215
alternative hypothesis: true mean is not equal to 24.3
95 percent confidence interval:
24.47413 25.58187
sample estimates:
mean of x
25.028
```

One Sample t-test

```
data: X
t = -3.3285, df = 11, p-value = 0.003364
alternative hypothesis: true mean is less than 0
95 percent confidence interval:
-Inf -0.2801098
sample estimates:
mean of x
-0.6083333
```

One Sample t-test

```
data: X
t = 0.3665, df = 7, p-value = 0.3624
alternative hypothesis: true mean is greater than 45
95 percent confidence interval:
44.11393 Inf
sample estimates:
mean of x
45.2125
```