YSU ASDS, Statistics, Fall 2019 Lecture 27

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- ► Likelihood Ratio Tests
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- p-Values

Last Lecture ReCap

▶ What are the two Sample *Z*- or *t*-Test about?

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- ▶ Describe the two Sample *t*-test.



Paired *t*-Test for the Difference of two Normals Means Model: $X_1, X_2, ..., X_n \stackrel{IID}{\sim} \mathcal{N}(\mu_X, \sigma_X^2)$,

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Asymptotic Significance Level: $\alpha \in (0,1)$;

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Test Statistics: $t = \frac{\overline{D} - \mu_0}{S_D/\sqrt{n}}$, where S_D is the Sample Deviation of D.

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$$\mu_X - \mu_Y \neq \mu_0 \quad |t| > t_{n-1,1-\frac{\alpha}{2}}$$
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Note: This Test is called the **Paired** *t*-**Test**

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Here we have paired results, since, the first elements in each line, 0 and 1, are the results of running Algorithms on the same Test case. Now, we want to do a Test - paired t-Test, to see if the Probability of correct Classification is the same for each Algorithm.

Example, cont'd

Here is the code for some hypothetical results from two hypothetical Algorithms:

```
res.algo1 <- rbinom(300, size = 1, prob = 0.4)
res.algo2 <- rbinom(300, size = 1, prob = 0.45)
t.test(res.algo1,res.algo2,paired = T)</pre>
```

```
##
##
   Paired t-test
##
## data: res.algo1 and res.algo2
## t = -1.0952, df = 299, p-value = 0.2743
## alternative hypothesis: true difference in means is not
## 95 percent confidence interval:
  -0.1212004 0.0345337
## sample estimates:
## mean of the differences
##
               -0.043333333
```

Likelihood Ratio Test

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where $\mathcal{L}(\theta)$ is our GOF *Likelihood Function*:

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$$\sup_{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mathcal{L}(\boldsymbol{\theta}) = \mathcal{L}\left(\hat{\boldsymbol{\theta}}^{\textit{MLE}}\right).$$

And similarly, $\sup_{\theta \in \Theta_0} \mathcal{L}(\theta)$ can be thought as a *restricted MLE for* θ *over* Θ_0 .

Assume $X_1,...,X_n\sim \mathcal{F}_{\theta}$, and we want to Test, at α -level, a Simple Hypothesis vs Simple Hypothesis:

$$\mathcal{H}_0: \ \theta = \theta_0 \qquad \textit{vs} \qquad \mathcal{H}_1: \ \theta = \theta_1.$$

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$$\mathbb{P}(LR < c \mid \mathcal{H}_0 \text{ is true }) = \alpha.$$

Then this Test has the highest Power among all other Tests of Significance Level α .

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 vs $\mathcal{H}_1: \mu \neq \mu_0$.

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i.e.

$$\mathcal{L}(\mu) = \left(\frac{1}{\sqrt{2\pi\sigma^2}}\right)^n \cdot e^{-\frac{1}{2\sigma^2} \cdot \sum_{k=1}^n (X_k - \mu)^2}$$

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We can write

$$\sup_{\mu \in \mathbb{R}} \mathcal{L}(\mu) = \mathcal{L}\Big(\hat{\mu}^{\textit{MLE}}\Big),$$

since, by the very definition, $\hat{\mu}^{MLE}$ is the global max point of $\mathcal{L}(\mu)$.

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Log-Likelihood Ratio, its Distribution

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$$-2 \cdot \ln LR = -2 \cdot \Big(\ln \big(\sup_{\theta \in \Theta_0} \mathcal{L}(\theta) \big) - \ln \big(\sup_{\theta \in \Theta} \mathcal{L}(\theta) \big) \Big).$$

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$$\nu = \dim \Theta - \dim \Theta_0,$$

and dim Θ and dim Θ_0 are the numbers of Free Parameters under Θ and Θ_0 , respectively.

For example, if we are Testing for the $\mathcal{N}(\mu, \sigma^2)$ Model

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And having this Distribution, we can find the Rejection Region using

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Note: When doing Tests, say, with t.test, \mathbf{R} is calculating the p-Value, and sometimes also the CI. So, to decide whether to Reject Null or Not, using \mathbf{R} , you can use the 2nd and 3rd Methods.

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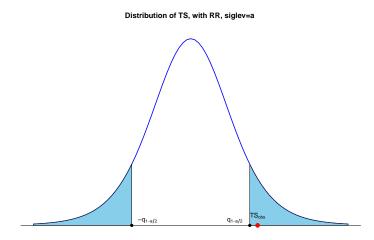
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based on the Test Statistics TS. Assume we already have Observations, and we calculate the value of TS, let us denote that by TS_{obs} (this is just a number). We know that, for a given Significance Level α , we will Reject \mathcal{H}_0 , iff TS_{obs} will be in the RR.

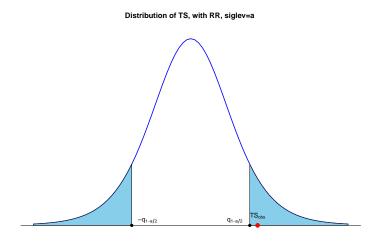
Now, assume the Distribution of TS, our Test Statistics, **under** \mathcal{H}_0 , is given like this (I am drawing for Z- or t-Statistics, for Two Tailed Test, the other cases can be considered in a similar way):

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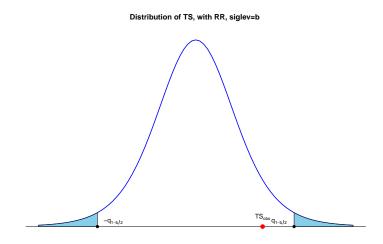
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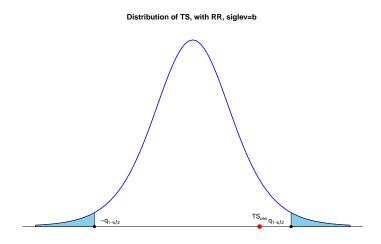
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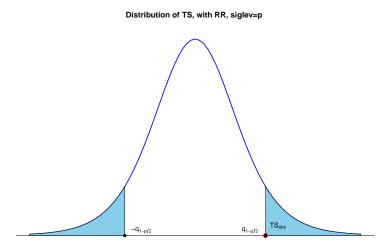
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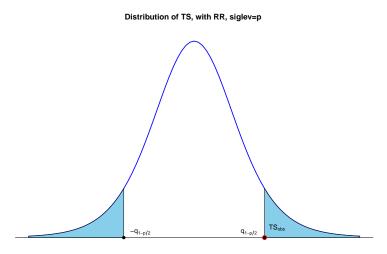
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Now, we denote $p = \alpha^*$ and call it the p-Value of the Test:

p-Value = p = inf{ α : we Reject \mathcal{H}_0 at level α }.





 $\emph{p}\text{-Value},$ the inf value of α at which we Reject \mathcal{H}_0

It is clear from the Figure above that

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To Remember:

- ▶ If p-Value< α , then we Reject \mathcal{H}_0
- ▶ If p-Value $\geq \alpha$, then we Fail to Reject \mathcal{H}_0

```
R Code for the Graphics
   df <- 8;
   x \leftarrow seq(-4,4,0.1); y \leftarrow dt(x, df = df)
   plot.new()
   plot.window(xlim = c(-4, 4), ylim = c(-0.05, 0.4))
   plot(x,y, type="l",col="blue",lwd=2,xaxt="n",yaxt="n",
         bty="n",xlab="",ylab="")
   abline(h=0)
   title("Distribution of TS, with RR, siglev=a ")
   gpoint <- 1.5; tspoint <- 1.7</pre>
    cord.x \leftarrow c(qpoint, seq(qpoint, 4, 0.01), 4)
    cord.y \leftarrow c(0,dt(seq(qpoint,4,0.01), df=df),0)
   polygon(cord.x,cord.y,col='skyblue')
   points(c(qpoint), c(0), pch=20, cex=1.4)
    text(c(qpoint-0.38), c(0.01), labels=expression("q"[1-a/2]))
   cord.x1 \leftarrow c(-4, seq(-4, -qpoint, 0.01), -qpoint)
    cord.y1 \leftarrow c(0,dt(seq(-4,-qpoint,0.01), df=df),0)
   polygon(cord.x1,cord.y1,col='skyblue')
   points(c(-qpoint), c(0), pch=20, cex=1.4)
   text(c(-qpoint+0.4), c(0.01), labels=expression("-q"[1-a/2]))
   points(c(tspoint), c(0), col="red", pch=19, cex=1.4)
```

text(c(tspoint), c(0.02), labels = expression("TS"[obs]))

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and the value of $p ext{-Value}$ is $\ddot{-}$

[1] 0.08543244