

# Resampling Techniques and their Application

## -Class 2-

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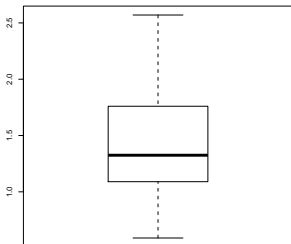
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## Motivation and Examples-II

The diameter of cork of a Champagne bottle is supposed to be 1.5 cm. If the cork is either too large or too small, it will not fit in the bottle. The manufacturer measures the diameter in a random sample of **n=36** bottles and obtains:

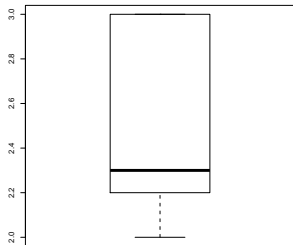


```
x = c (
0.59, 1.23, 1.00, 0.84, 0.88, 1.71,
1.81, 1.84, 2.03, 1.39, 1.30, 1.31,
1.96, 1.33, 2.57, 1.19, 1.01, 2.06,
1.32, 1.55, 1.28, 0.93, 1.63, 1.24,
1.83, 1.81, 0.94, 1.46, 1.25, 1.56,
0.61, 0.83, 1.17, 2.24, 1.68, 1.51)
```

- Data Analysis: Confidence interval and t-test

## Motivation and Examples-III

A researcher measures the reaction time of  $n = 10$  mice to signal pain when a stitch is applied to their tail.



```
x = c(  
  2.4, 3.0, 3.0, 2.2, 2.2,  
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## Example

The diameter of cork of a Champagne bottle is supposed to be 1.5 cm. If the cork is either too large or too small, it will not fit in the bottle. The manufacturer measures the diameter in a random sample of  $n=36$  bottles. Is there evidence at 1% level that the true mean diameter has moved away from the target?

- Hypothesis:  $H_0 : \mu = 1.5$  vs.  $H_1 : \mu \neq 1.5$

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mx <- mean(x)
vx <- var(x)
T <- sqrt(36)*(mx-1.5)/sqrt(vx)
p <- 2*min(pt(T,35),1-pt(T,35))
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- p-value: 2 times the area to the **left** of T under the t curve with 35 df. (Here,  $p=0.27$ )
- Quality of the estimator? Is the method valid? Is  $p = 0.27$  a good estimate?

## Example

A researcher measures the reaction time of  $n = 10$  mice to signal pain when a stitch is applied to their tail. Compute a 95% confidence interval.

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2.2, 2.2, 2.8, 2.0, 3.0)  
mx <- mean(x)  
vx <- var(x)  
crit <- qt(0.975,9)  
lower <- mx-crit/sqrt(10)*sqrt(vx)  
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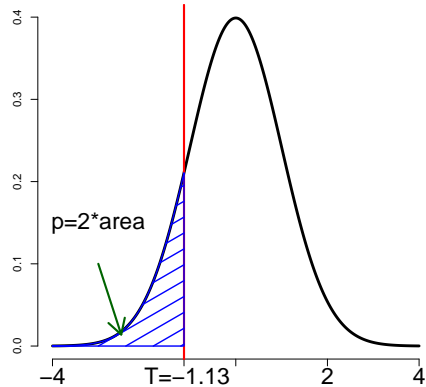
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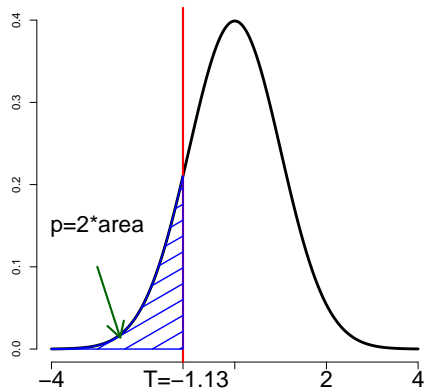
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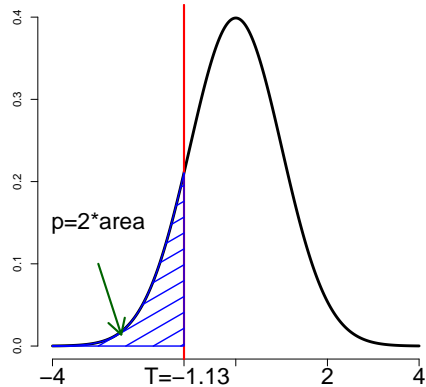
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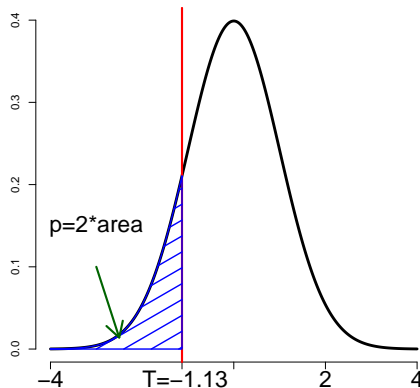
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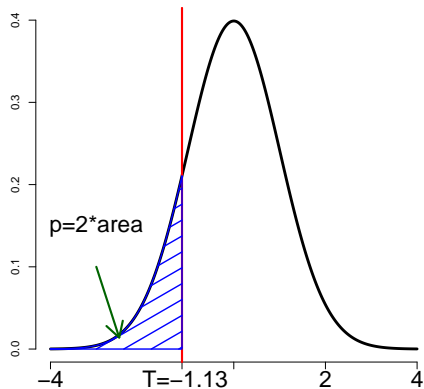
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( $X_1, \dots, X_n \sim N(\mu, \sigma^2)$ )
- **What happens if data come from a different distribution?**



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- Can we verify/visualize the distribution from the boxplot?

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  4. Repeat the above a large number of times
  5. Estimate the type-1 error by averaging the indicators

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- $E(X_k) = 0$  and  $\text{Var}(X_k) = \sigma^2$ 
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  - ...

```
mysimulation <-function(Needed Parameters){
```

```
  Data Generation
```

```
  Compute the Test Statistic  
  Hypothesis Rejected? (0/1)
```

```
  Estimate Type-1 error rate  
  output table}
```

```
mysimulation(...)
```

```
mysimulation <-function(n,s2,nsim){  
  crit= qt(0.975, n-1) # critical value at 5\% level  
  ttest <-c(); set.seed(1)  
  for(i in 1:nsim){ #Begin Simulation Loop
```

```
    x <- rnorm(n)*sqrt(s2) #generate from normal dist.  
    #mu=0 and has variance s2
```

```
    mx <-mean(x); sdx <- sd(x); T<-sqrt(n)*mx/sdx #compute test statistic  
    ttest[i] <- (abs(T)>=crit) #Reject yes/no  
  } #End Simulation Loop
```

```
  result= data.frame(n=n, alpha=0.05, sigma2=s2, tTest=mean(ttest))  
  result}  
  mysimulation(10,1,10000)
```

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```
    x <- (rexp(n)-1)*sqrt(s2) #generate data from exponential  
    #(hypothesis is TRUE! has variance=s2)
```

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    mx <-mean(x); sdx <- sd(x); T<-sqrt(n)*mx/sdx #compute test statistic  
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```
mysimulation <-function(n,s2,nsim, Distribution){ #use Distribution as argument
```

```
crit= qt(0.975, n-1) # critical value at 5\% level  
ttest <-c()  
set.seed(1)  
for(i in 1:nsim){ #Begin Simulation Loop
```

```
  if(Distribution=="Exp"){#generate data from exponential  
    x <- (rexp(n)-1)*sqrt(s2) }  
  if(Distribution=="Normal"){#generate data from normal  
    x <- rnorm(n)*sqrt(s2) }
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  mx <-mean(x); sdx <- sd(x); T<-sqrt(n)*mx/sdx #compute test statistic  
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- Type-I error simulation (10K simulations,  $\alpha = 5\%$ ); Test  $H_0 : \mu = 0$



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- Different distributions
- Fill the table:

| Sample Size | Distribution | Shape     | Emp. Type-I | Accurate (yes/no) |
|-------------|--------------|-----------|-------------|-------------------|
| Small       | Normal       | Symmetric |             |                   |
|             | Exponential  | Skewed    |             |                   |
| Moderate    | Normal       | Symmetric |             |                   |
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- Can we analyze the data sets with the method?

## Impact of the Small Samples

- Type-I error simulation (10K simulations,  $\alpha = 5\%$ ); Test  $H_0 : \mu = 0$
- Sample sizes: Small ( $n=10$ ), moderate ( $n=25$ ) and large ( $n=50$ )
- Different distributions
- Fill the table:

| Sample Size | Distribution | Shape     | Emp. Type-I | Accurate (yes/no) |
|-------------|--------------|-----------|-------------|-------------------|
| Small       | Normal       | Symmetric |             |                   |
|             | Exponential  | Skewed    |             |                   |
| Moderate    | Normal       | Symmetric |             |                   |
|             | Exponential  | Skewed    |             |                   |
| Large       | Normal       | Symmetric |             |                   |
|             | Exponential  | Skewed    |             |                   |

- Can we analyze the data sets with the method?
- Can we take data characteristics into account? ( $\Rightarrow$  Resampling)

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- Try to avoid "for" loops
- Generate a matrix of variables (example next slide)

```
mysimulation <-function(n,s2,Distribution,nsim){  
  crit= qt(0.975, n-1) # critical value at 5\% level  
  set.seed(1)
```

```
  if(Distribution=="Normal"){  
    x <- matrix(rnorm(n=n*nsim)*sqrt(s2),ncol=nsim)}  
    if(Distribution=="Exp"){#(Hypothesis is TRUE!)  
    x <- matrix((rexp(n=n*nsim)-1)*sqrt(s2),ncol=nsim)}
```

```
  mx <-colMeans(x)  
  vx <- (colSums(x^2)-n*mx^2)/(n-1)  
  T <- sqrt(n)*mx/sqrt(vx)  
  ttest <- (abs(T)>=crit)
```

```
  result= data.frame(n=n,Dist=Distribution,sigma2=s2,ttest=mean(ttest))  
  result}  
  mysimulation(10,1,"Normal",10000)
```

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- Which one is better?
- Assess a simulation to compute a) their bias and b) their standard error

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  - Compute estimator and save the value of  $\hat{\theta}_\ell$ ,  $\ell = 1, \dots, n_{sim}$
  - End of simulation loop
  - Assess bias and mean square error as

$$bias = \frac{1}{n_{sim}} \sum_{\ell=1}^{n_{sim}} (\hat{\theta}_\ell - \theta) \qquad MSE = \frac{1}{n_{sim}} \sum_{\ell=1}^{n_{sim}} (\hat{\theta}_\ell - \theta)^2$$

```
mysimulation <-function(Needed Parameters){
```

```
  Data Generation (know true value)
```

```
  Compute the Point Estimator and save the value
```

```
  Compute the bias and MSE  
  output table}
```

```
mysimulation(...)
```

```
mysimulation <-function(n,s2,Distribution,nsim){
```

```
  if(Distribution=="Normal"){  
    x <- matrix(rnorm(n=n*nsim)*sqrt(s2),ncol=nsim)}  
    if(Distribution=="Exp"){  
      x <- matrix((rexp(nsim*n)-1)*sqrt(s2),ncol=nsim)} #(Parameter is known)
```

```
  mx <-colMeans(x)  
  v1 <- (colSums(x^2)-n*mx^2)/(n-1)  
  v2<- (colSums(x^2)-n*mx^2)/n
```

```
  result= data.frame(n=n, sigma2=s2, Distribution= Distribution,  
    bias.v1=mean(v1-s2), MSE.v1=mean((v1-s2)^2), bias.v2=mean(v2-s2),  
    MSE.v2=mean((v2-s2)^2))  
  result}  
  mysimulation(10,1,"Normal",10000)
```

# Project

1. Write a simulation program to assess the quality of 95% confidence interval for mean
2. Let  $X_1, \dots, X_n$  have  $E(X_k) = \mu$ . We want to estimate  $\theta = \mu^2$ . Which of the estimators

$$\hat{\theta}_1 = \bar{X}^2 \quad \text{or} \quad \hat{\theta}_2 = \frac{1}{n(n-1)} \sum_{i \neq j} X_i \cdot X_j$$

would you recommend?