Resampling Techniques and their Application

-Class 6-

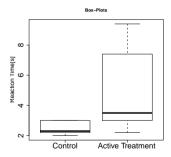
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Motivation and Examples-III

Researchers produce a pain killer using poison from a snake. They investigate the effect of the treatment on n_1 mice in the **control** group and $n_2 = 10$ mice in the **active treatment**. The response variable is the reaction time of the mice to signal pain when a stitch is applied to their tail. Is the treatment effective? (all mice survived the dose)



```
x=c(
2.4, 3.0, 3.0, 2.2, 2.2,
2.2, 2.2, 2.8, 2.0, 3.0)

y=c(
2.8, 2.2, 3.8, 9.4, 8.4,
3.0, 3.2, 4.4, 3.2, 7.4)
```

• Aim: Test H_0 : $\mu_1 = \mu_2$ and confidence interval for $\delta = \mu_1 - \mu_2$

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$$\overline{X}_{i\cdot} = \frac{1}{n_i} \sum_{k=1}^{n_i} X_{ik}$$

• $\hat{\sigma}_1^2$ and $\hat{\sigma}_2^2$: empirical variances per group with

$$\widehat{\sigma}_i^2 = \frac{1}{n_i - 1} \sum_{k=1}^{n_i} (X_{ik} - \overline{X}_{i.})^2$$

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$$\widehat{\sigma}_{p}^{2} = \frac{(n_{1}-1)\widehat{\sigma}_{1}^{2} + (n_{2}-1)\widehat{\sigma}_{2}^{2}}{n_{1}+n_{2}-2} = \frac{1}{N-2}\sum_{i=1}^{2}\sum_{k=1}^{n_{i}}(X_{ik}-\overline{X}_{i\cdot})^{2}$$

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$$T = \frac{\overline{X}_{1.} - \overline{X}_{2.}}{\widehat{\sigma}_P \sqrt{1/n_1 + 1/n_2}}$$

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• Reject H_0 , if $|T| \ge t_{1-\alpha/2}(n_1 + n_2 - 2)$, the $(1 - \alpha/2)$ -quantile from the t-distribution with N-2 degrees of freedom

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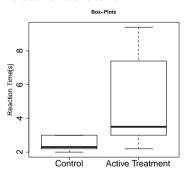
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• Reject H_0 , if $|T| \ge t_{1-\alpha/2}(\nu)$,

$$\nu = \frac{(\frac{\widehat{\sigma}_1^2}{n_1} + \frac{\widehat{\sigma}_2^2}{n_2})^2}{\frac{\widehat{\sigma}_1^4}{n_1^2(n_1 - 1)} + \frac{\widehat{\sigma}_2^4}{n_2^2(n_2 - 1)}}$$

degrees of freedom (Satterthwaite's approximation).

Researchers produce pain killer using poison from a cobra. The investigate the effect of the treatment on n_1 mice in the **control** group and $n_2 = 10$ mice in the **active treatment**. The response variable is the reaction time of the mice to signal pain when a stitch is applied to their tail. Is the treatment effective?



```
react <- data.frame(resp=c(x,y),
grp=factor(c(rep(1,10),rep(2,10))))
t.test(resp~grp,data=react,
var.equal=TRUE)
t.test(resp~grp,data=react,
var.equal=FALSE)</pre>
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- ullet Compute critical and p-values from the permutation distribution of T_{ns}

- Workflow
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- Repeat the above a large number of times (n_{perm}) and estimate the p-value

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- Use matrix technique to multiply a permutation matrix with the sample

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• Arrange the coefficients c_{ℓ}^* in a $n_{perm} \times N$ matrix **P** and multiply with **X**. This operation computes all permuted values T_{ns}^*

```
mypermu<-function(nsim,nperm,n1,n2,s1,s2,Distribution) {
N<-n1+n2;erg<-c()
#-------Permutation Matrices-----#
i1<-sqrt(n1*n2/N)*c(rep(1/n1,n1),rep(-1/n2,n2))
P<-t(apply(matrix(i1,nrow=nperm,ncol=N,byrow=TRUE),1,sample))</pre>
```

```
x1<-matrix(rnorm(n1*nsim)*sqrt(s1),ncol=nperm,nrow=n1)
x2 <-matrix(rnorm(n2*nsim)*sqrt(s2),ncol=nperm,nrow=n2)}
x<-rbind(x1,x2)
mx1<-colMeans(x1); mx2<-colMeans(x2)
Tns <-sqrt(n1*n2/N)*(mx1-mx2)</pre>
```

if (Distribution == "Normal") {

```
for (i in 1:nsim){
  X<-x[,i]
#-------Permutations-----#
  TnsP <- P%*%X #actual samole
  pvalue<-2*min(mean(TnsP<=Tns[i]),mean(TnsP>=Tns[i]))
```

```
\label{eq:continuous} $$ \operatorname{erg}[i] <-(\operatorname{pvalue}<0.05)$ \\ \operatorname{result}<-\operatorname{data.frame}(\operatorname{nsim-nsim},\operatorname{nperm-nperm},\operatorname{n1=n1},\operatorname{n2=n2},\operatorname{s1=s1},\operatorname{s2=s2},\operatorname{Dist-Distribution}, \\ \operatorname{Permu-mean}(\operatorname{erg})) \\ \operatorname{result}$ \\ \operatorname{mypermu}(10000,10000,10,10,10,1,1,"\operatorname{Normal"}) \\ \end{aligned}
```

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- $\overline{X}_{..} = \frac{1}{N} \sum_{i=1}^{2} \sum_{k=1}^{n_i} X_{ik}$
- Only valid, if... $n_1 = n_2$ or $\sigma_1^2 = \sigma_2^2$

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$$E(X_i^* X_j^* | \mathbf{X}) = \sum_{\ell \neq k} X_\ell X_k P(X_i^* = X_\ell, X_j^* = X_k) = \frac{1}{N(N-1)} \sum_{\ell \neq k} X_\ell X_k$$

$$= \overline{X}... \sum_{\ell=1}^N c_\ell = 0$$

$$Cov(X_i^*, X_j^* | \mathbf{X}) = E(X_i^* X_j^* | \mathbf{X}) - E(X_i^* | \mathbf{X}) E(X_j^* | \mathbf{X})$$

$$= -\frac{1}{N(N-1)} \sum_{\ell=1}^N (X_\ell - \overline{X}..)^2$$
Now plug everything together

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$$T = \frac{\overline{X}_{1.} - \overline{X}_{2.}}{\sqrt{\widehat{\sigma}_1^2/n_1 + \widehat{\sigma}_2^2/n_2}}$$

• Instead of using the numerator of the statistic only, investigate the permutation distribution of

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Verify in a simulation study

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Group wise Nonparametric Bootstrap

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...

In R: sample(x1,replace=TRUE)

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- $X_2 = (X_{21}, \dots, X_{2n_2})$ (fixed values)
- Drawing with Replacement: randomly draw n_1 and n_2 observations from X_1 and X_2
- Example $\mathbf{X}_1 = (1, 2, 3, 4, 5) \Rightarrow$ $\mathbf{X}_1^* = (2, 2, 4, 3, 2)$ $\mathbf{X}_1^* = (1, 1, 2, 3, 3)$ $\mathbf{X}_1^* = (2, 5, 5, 3, 3)$

...

- In R: sample(x1,replace=TRUE)
- Also known as Group wise Nonparametric Bootstrap

• Data $X = (X_{11}, \dots, X_{2n_2})$ (fixed values)

- Data $X = (X_{11}, ..., X_{2n_2})$ (fixed values)
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- In R: sample(x)
- Also known as Permutation

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- Also known as Parametric Bootstrap (Why is that not equivalent to the t-approximation?)

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Also known as Wild-Bootstrap

Project

 Which resampling method is better? Permutation, Nonparametric Bootstrap or Parametric Bootstrap?

Project

- Which resampling method is better? Permutation, Nonparametric Bootstrap or Parametric Bootstrap?
- Write your own simulation program