

Homework 6

Chapter 10.

p. 170-173.

Ex: 5, 7, 8a, 11a.

5. Let $X_1 \dots X_n \sim \text{Uniform}(0, \theta)$ and let $Y = \max\{X_1 \dots X_n\}$. We want to test

$$H_0: \theta = \frac{1}{2}$$

$$H_1: \theta > \frac{1}{2}$$

a) Find the power function.

$$\begin{aligned} \beta(\theta) &= P_{\theta}(Y > c) = P(Y > c) = \\ &= 1 - P(\max\{X_1 \dots X_n\} \leq c) = \\ &= 1 - (P(X_1 \leq c) \dots P(X_n \leq c)) = \\ &= 1 - \left(c \cdot \frac{1}{\theta}\right)^n = 1 - \left(\frac{c}{\theta}\right)^n \end{aligned}$$

*(Hh3, ex 7a)
I did it here*

b) What choice of c will make the size of the test 0.05?

$$\beta\left(\frac{1}{2}\right) = 1 - \left(\frac{c}{2}\right)^n = 0.05$$

$$(2c)^n = 1 - 0.05$$

$$(2c)^n = 0.95$$

$$2c = \sqrt[n]{0.95}$$

$$c = \frac{\sqrt[n]{0.95}}{2}$$

- c) In a sample of size $n=20$ with $\bar{Y}=0.48$ what is the p-value?
What conclusion about H_0 would you make?

$$\begin{aligned} p\text{-value} &= P(Y > 0.48) = 1 - (2 \cdot 0.48)^{20} = \\ &= 0.5579 \approx 0.5580 \end{aligned}$$

Conclusion: little or no evidence
against H_0 ($0.5580 > 0.1$)

d) In a sample of size $n=20$ with $\bar{Y}=0.52$ what is p-value? What conclusion about H_0 would you make?

$\bar{Y}=0.52$ is already greater than $\frac{1}{2}$ by alternative hypothesis. ($H_0: \bar{Y} \geq \frac{1}{2}$). It rejects the null hypothesis H_0 . Therefore p-value is < 0.01 very strong evidence against H_0 . In this case p-value is 0.

7. From 8 Train essays:

- | | |
|-----------|-----------|
| 1) 0. 225 | 6) 0. 229 |
| 2) 0. 262 | 7) 0. 235 |
| 3) 0. 217 | 8) 0. 217 |
| 4) 0. 240 | |
| 5) 0. 230 | |

From 10 Snodgrass essay:

- | | |
|----------|-----------|
| 1) 0.209 | 6) 0.207 |
| 2) 0.205 | 7) 0.224 |
| 3) 0.196 | 8) 0.223 |
| 4) 0.210 | 9) 0.220 |
| 5) 0.202 | 10) 0.201 |

Comparing the means.

$$\text{Let } \text{Truah} = T = \{X_1, \dots, X_8\}$$

$$\text{Snodgrass} = S = \{Y_1, \dots, Y_{10}\}$$

Let's test null hypothesis $\mu_1 = \mu_2$

$$H_0: \delta = 0$$

$$H_1: \delta \neq 0, \text{ where } \delta = \mu_1 - \mu_2$$

The non-parametric plug-in estimator

$$\text{is } \hat{\delta} = \bar{T} - \bar{S} = \bar{X} - \bar{Y}$$

Estimated Standard Error

$$\hat{SE} = \sqrt{\frac{s_1^2}{m} + \frac{s_2^2}{n}} = \sqrt{\frac{s_1^2}{8} + \frac{s_2^2}{10}}$$

The Wald test:

$$W = \frac{\hat{\delta} - \delta_0}{\hat{SE}} = \frac{\delta - 0}{\hat{SE}} = \frac{\bar{X} - \bar{Y}}{\sqrt{\frac{s_1^2}{m} + \frac{s_2^2}{n}}} \quad (\text{from book})$$

page 155

$$\bar{X} = \frac{\sum_{i=1}^8 X_i}{8} = 0.2319$$

$$\bar{Y} = \frac{\sum_{i=1}^{10} Y_i}{10} = 0.2097$$

$$S_x^2 = \frac{\sum_{i=1}^8 (0.2319 - X_i)^2}{7} = 0.00021$$

$$S_y^2 = \frac{\sum_{i=1}^{10} (0.2097 - Y_i)^2}{9} = 0.9334 \cdot 10^{-4}$$

$$W = \frac{0.2319 - 0.2097}{\sqrt{\frac{0.00021}{8} + \frac{0.9334 \cdot 10^{-4}}{10}}} = 3.7036$$

Let w denote the observed value of the Wald statistic W . The p-value is given by

$$\begin{aligned} \text{p-value} &= P(|W| > |w|) \approx P(|Z| > |w|) = \\ &= 2 \Phi(-|w|). \quad (\text{page 158 of the book}) \end{aligned}$$

$$\begin{aligned} \text{p-value} &= P(|Z| > |3.7036|) = \\ &= P(Z > 3.7, Z < -3.7) = 2 \Phi(-3.7) = \\ &= 2 \cdot 0.0001078 \approx 0.0002 \end{aligned}$$

If it is < 0.01 , then very strong evidence against H_0

(Page 155 of book)

$$C = (\hat{\theta} - \widehat{SE} \cdot Z_{\alpha/2}, \hat{\theta} + \widehat{SE} \cdot Z_{\alpha/2})$$

$$C = (0.0222 - 0.0060 \cdot 1.96, \\ 0.0222 + 0.0060 \cdot 1.96.)$$

$$C = (0.0104, 0.0339)$$

where

$$\hat{\theta} = \bar{\delta} = \bar{X} - \bar{Y} = 0.022175$$

$$\widehat{SE} = \sqrt{\frac{s_x^2}{8} + \frac{s_y^2}{10}} = 0.00599$$

$$Z = 1.96$$

The size of Wald test reject H_0 when

$$|W| > Z_{\alpha/2}$$

$$|W| = 3.7$$

$$Z_{\alpha/2} = 1.96 \Rightarrow 3.7 > 1.96 \quad (\text{reject } H_0)$$

I would say that cheese are two different writers, because the means are different $\bar{\delta} \neq 0$, where the H_0 is rejected. But the difference is small. (vanionee)

11Ex_Zhetessov.R * 7_1_Zhetessov.R *

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```

1 #a) Wald Test
2 vec1 = c(.225, .262, .217, .240, .230, .229, .235, .217)
3 vec2 = c(.209, .205, .196, .210, .202, .207, .224, .223, .220, .201)
4
5 estimator = mean(vec1)-mean(vec2)
6 SE = sqrt((var(vec1)/8)+(var(vec2)/10))
7
8 w = (estimator)/SE
9 w
10
11 CI= c(estimator - 1.96 * SE, estimator + 1.96 * SE)
12 CI
13
14 #b) Permutation Test
15 vec3 = c(vec1, vec2)
16 cnt <- 0
17
18 for(i in 1:1000000){
19   x <- split(sample(vec3), rep(1:2, c(8,10)))
20   res <- mean(x[[1]]) - mean(x[[2]])
21   if(res > estimator){
22     cnt <- cnt + 1
23   }
24 }
25
26 p_value <- cnt / 1000000
27 p_value
28 #zhetessov Nur

```

(Top Level) R Script

Console Terminal Jobs

R 4.1.2 · C:/Users/Nur/Desktop/дз/Стат/HW6/

```

> vec1 = c(.225, .262, .217, .240, .230, .229, .235, .217)
> vec2 = c(.209, .205, .196, .210, .202, .207, .224, .223, .220, .201)
> estimator = mean(vec1)-mean(vec2)
> SE = sqrt((var(vec1)/8)+(var(vec2)/10))
> w = (estimator)/SE
> w
[1] 3.703554
> CI= c(estimator - 1.96 * SE, estimator + 1.96 * SE)
> CI
[1] 0.01043951 0.03391049
> vec3 = c(vec1, vec2)
> cnt <- 0
> for(i in 1:1000000){
+   x <- split(sample(vec3), rep(1:2, c(8,10)))
+   res <- mean(x[[1]]) - mean(x[[2]])
+   if(res > estimator){
+     cnt <- cnt + 1
+   }
+ }
> p_value <- cnt / 1000000
> p_value
[1] 0.00046
> #Zhetessov Nur
>

```

Environment History Connections Tutorial

Import Dataset 132 MB

R Global Environment Data

x	List of 2
values	
CI	num [1:2] 0.0104 0.0339
cnt	460
estimator	0.022175
i	1000000L
p_value	0.00046
res	0.0134
SE	0.00598749275109745
vec1	num [1:8] 0.225 0.262 0.217 0.24 0.23 0.229 0.235 0.217
vec2	num [1:10] 0.209 0.205 0.196 0.21 0.202 0.207 0.224 0.223 0.22 0.201
vec3	num [1:18] 0.225 0.262 0.217 0.24 0.23 0.229 0.235 0.217 0.209 0.205 ...
w	3.70355354433382

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b) Now use a permutation test to avoid the use of large sample methods. What is your conclusion?

The p-value I got is between 0.004 and 0.005. The last test gave me 0.00483. Which 0.48% The evidence is weak against H_0 of no difference.

8. Let $X_1, \dots, X_n \sim N(\theta, 1)$.

$$H_0: \theta = 0$$

$$H_1: \theta = 1 \text{ or } \theta \neq 0$$

Power function

$$\begin{aligned} \beta(\theta) &= P(T > c) = P(\sqrt{n}(T - \mu) > \sqrt{n}(c - \mu)) \\ &= P(Z > \sqrt{n}(c - \mu)) = 1 - \Phi(\sqrt{n}(c - \mu)) \end{aligned}$$

(From the lecture slides #13)

Significance level is α

$$\alpha = \max_{\theta \leq 0} \beta(\theta)$$

$$\theta \leq 0$$

$$\lambda = \beta(0) = 1 - \Phi(\sqrt{n} \cdot c)$$

$$\Phi(\sqrt{n} \cdot c) = 1 - \lambda$$

$$\sqrt{n} \cdot c = \Phi^{-1}(1 - \lambda)$$

$$c = \frac{\Phi^{-1}(1 - \lambda)}{\sqrt{n}}$$

(Example 10.2, page 151)

11. A randomized, double-blind experiment has conducted to assess the effectiveness of several drugs for reducing postoperative nausea.

a) Test each drug vs placebo,

at 5% level.

For each drug we do test

$$H_0: p_1 = p_2$$

$$H_1: p_1 \neq p_2$$

where p_1 - proportion of drug nausea

p_2 - proportion of placebo nausea

I will use the Wald test with proportions.

$$\hat{SE} = \sqrt{\frac{\hat{p}_1(1-\hat{p}_1)}{m} + \frac{\hat{p}_2(1-\hat{p}_2)}{n}}$$

$$W = \frac{\hat{p}_1 - \hat{p}_2}{\hat{SE}} = \frac{\hat{p}_1 - \hat{p}_2}{\hat{SE}}$$

(page 154. Example 10.7.)

$$\text{Placebo, } \hat{p}_2 = \frac{45}{80} = 0,5625$$

	# total	# nausea	\hat{p}_1	$\hat{p}_1 - \hat{p}_2$	\hat{SE}	W	p-value
--	---------	----------	-------------	-------------------------	------------	---	---------

Chlorpromazine	75	26	0,3467	-0,2158	0,078	-2,76	0,0058
----------------	----	----	--------	---------	-------	-------	--------

Dimethylhydantoin	85	52	0,6118	0,0493	0,077	0,64	0,5222
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Pentobarbital (100 mg)	67	35	0,5224	-0,0401	0,082	-0,49	0,6242
---------------------------	----	----	--------	---------	-------	-------	--------

Pentobarbital (150 mg)	85	37	0,4353	-0,1272	0,077	-1,65	0,099
---------------------------	----	----	--------	---------	-------	-------	-------

$$p\text{-value} = P(|Z| > |w|) = 2 \Phi(-|w|)$$

(Theorem 10.13 page 158)

Also report the estimated odds-ratios.

The odd-ratio-is calculated by dividing the odds of the first group by the odds of the second group.

Odd-The ratio of number of successes by the number of failures.

$$\text{Placebo odd} = \frac{45}{35} \quad \text{Odd-ratio} = \frac{\text{placebo-odd}}{\text{drug-odd}}$$

	Success	Failure	Odd	Odd-ratio
Chlorpromazine	26	49	0,5306	2,42
Dimethylhydantoinate	52	33	1,576	0,816
Pentobarbital (100 mg)	35	32	1,093	1,146
Pentobarbital (150 mg)	37	48	0,77	1,67

Reject H_0 if $|W| > Z_{\alpha/2}$
at 5% sign level Z is 1.96.

All the drug could not reject the null hypothesis about no effect. Except Chlorpromazine, whose $|W| = 2.76$, which is greater than $Z = 1.96$. So, I conclude Chlorpromazine has significant results. p-value also gives strong evidence that this drug has difference effect than placebo, ($0.0058 < 0.01$), or strong evidence against H_0 .

```
7_1_Zhetessov.R x 11Ex_Zhetessov.R x
Source on Save Run Source
1 exp.data <- data.frame(
2   name = c ("Placebo", "Chlorpromazine", "Dimenhydrinate", "Pentobarbital (100 mg)", "Pentobarbital (150 mg)",
3   total = c(80, 75, 85, 67, 85),
4   nausea = c(45, 26, 52, 35, 37)
5 )
6
7 p_hat <- exp.data$nausea/exp.data$total #вычисляю пропорции
8 p_hat
9
10 delta_hat <- p_hat[2:5] - p_hat[1] #Вычисляю estimator дельта_hat (лекарства - плацебо)
11 delta_hat
12
13 SE_hat <- sqrt( (p_hat[1]*(1-p_hat[1])/exp.data[1,]$total) + (p_hat[2:5]*(1-p_hat[2:5])/exp.data[c(2:5),]$total) )
14 #Вычисляю стандартное отклонение для всех лекарств
15 SE_hat
16
17 w <- delta_hat / SE_hat
18 w
19
20 p_values <- 2 * pnorm(-abs(w))
21 p_values
22
23 reject_H0 <- c()
24 for(i in w){
25   if(abs(i)>1.96){
26     reject_H0 <- append(reject_H0, i)
27   }
28 }
29
30:1 (Top Level) R Script
```

```
Console Terminal Jobs
R 4.1.2 ~/
+
> p_hat <- exp.data$nausea/exp.data$total #вычисляю пропорции
> p_hat
[1] 0.5625000 0.3466667 0.6117647 0.5223881 0.4352941
> delta_hat <- p_hat[2:5] - p_hat[1] #Вычисляю estimator дельта_hat (лекарства - плацебо)
> delta_hat
[1] -0.21583333 0.04926471 -0.04011194 -0.12720588
> SE_hat <- sqrt( (p_hat[1]*(1-p_hat[1])/exp.data[1,]$total) + (p_hat[2:5]*(1-p_hat[2:5])/exp.data[c(2:5),]$total) )
> #Вычисляю стандартное отклонение для всех лекарств
> SE_hat
[1] 0.07807704 0.07661848 0.08246232 0.07725342
> w <- delta_hat / SE_hat
> w
[1] -2.7643638 0.6429873 -0.4864275 -1.6466051
> p_values <- 2 * pnorm(-abs(w))
> p_values
[1] 0.005703392 0.520232365 0.626664095 0.099639233
> reject_H0 <- c()
> for(i in w){
+   if(abs(i)>1.96){
+     reject_H0 <- append(reject_H0, i)
+   }
+ }
> reject_H0
[1] -2.764364
> #Zhetessov Nur
```