



INSTITUTE OF ACTUARIAL
& QUANTITATIVE STUDIES

Subject: App of IT-Basics of R

Chapter: Unit 1

Category: Practice Question Solution

Hypothesis Tests:**##Answer1**

```
sd1 = 8.4
n1 = 10
x1 = 162.8
```

##a]

```
##H0 : mu = 165
##Ha : mu < 165
```

```
mu1a = 165
Y1a <- c(rnorm(n1,x1,sd1))
t.test(Y1a,mu = mu1a,alternative = "less")
##Since, p-value > 0.05, we fail to reject our null hypothesis
```

##b]

```
##H0 : mu = 158
##Ha : mu != 158
mu1b = 158
Y1b <- c(rnorm(n1,x1,sd1))
t.test(Y1b,mu = mu1b)
##Since, p-bvalue > 0.05, we fail to reject our null hypothesis
```

##Answer2

```
sd2 = 12.1
n2 = 6
mu2 = 64.2
```

```
data <- c(64.30,68.93,74.91,52.71,59.70,73.60)
```

##b] Hypothesis Test

```
##H0: Mean weight of women in the US is 64.2kg
##Ha: Mean weight of women in the US is not 64.2kg
t.test(data,mu = mu2)
##Since, p-value > 0.05, we fail to reject our null hypothesis
##The mean weight of women in the US is 64.2kg
```

##Answer3

```

n3 = 10
u3 = 22
##a] Storing values in variable rain
rain <- c(17.2,28.1,25.3,26.2,30.7,19.2,23.4,27.5,29.5,31.6)

##b] Hypothesis Test
##H0: Mean annual rainfall is equal to 22cm
##Ha: Mean annual rainfall has increased from 22cm
t.test(rain,mu = u3,alternative = "greater")
##p-value = 0.01495
##Since, p-value < 0.05, we reject our null hypothesis.
##The average rainfall has increased from it's longer value of 22cm.

```

##Answer4

```

control <- c(27.0,32.2,30.4,28.0,26.5,25.5,29.6,27.2)
additive <- c(31.4,29.9,33.2,34.4,32.0,28.7,26.1,30.3)
var(control)
var(additive)
##Output
##> var(control)
##[1] 5.054286
##> var(additive)
##[1] 6.865714
##Thus, the variances are not equal, i.e. they are not homogeneous.

##CONducting t-test
t.test(control,additive,var.equal = FALSE)
##Since,p-value > 0.05, we fail to reject our null hypothesis.

```

##Answer5

```

n5a = 50
n5b = 40
x5a = 132
x5b = 128
sd5a = 20
sd5b = 25

Y5a <- c(rnorm(n5a,x5a,sd5a))
Y5b <- c(rnorm(n5b,x5b,sd5b))
var(Y5a)
var(Y5b)
##variances are not equal
##H0: Mean(IQs of University A) = Mean(IQs of University B)
##H1: Mean(IQs of University A) > Mean(IQs of University B)
t.test(Y5a,Y5b,var.equal = FALSE,alternative = "greater")
##Since,p-value > 0.05, we fail to reject our null hypothesis.

```

PROBABILITY

1

##a]

##Let A be the event that the first card was red and B be the event that the second card is a face card

$$##n(A) = 26$$

$$##n(B) = 12 \text{ (Jack, Queen and King)}$$

##Let PA be P(A), PB be P(B) and PAB be P(AnB)

$$PA = 26/52$$

$$PB = 12/52$$

$$PAB = 6/52$$

$$##P(B|A) = P(AnB)/P(A)$$

$$\text{conPBA} = PAB/PA$$

$$\text{conPBA}$$

$$##\text{Answer: } 0.2307692$$

##b]

##Let C be the event that the first card was face card and D be the event that the second card is an ace card.

##Using similar kind of notations

$$PC = 12/52$$

$$PD = 4/52$$

##Since, both events are independent $P(CnD) = P(C) * P(D)$

$$PCD = PC * PD$$

$$\text{conPDC} = PCD/PC$$

$$\text{conPDC}$$

$$##\text{Answer: } 0.07692308$$

##c]

##Let E be the event that the first card was red ace and F be the event that the second card is a black jack

##using similar notations,

$$PE = 2/52$$

$$PF = 2/52$$

##Since, both events are independent,

$$PEF = PE * PF$$

$$\text{conPFE} = PEF/PE$$

$$\text{conPFE}$$

$$##\text{Answer: } 0.03846154$$

##Answer2:

##Let PA, PB, PC and PX denote the probabilities of the respective events.

$$PA = 0.35$$

$$PB = 0.45$$

$$PC = 0.2$$

$$\text{conXA} = 0.8$$

$$\text{conXB} = 0.65$$

$$\text{conXC} = 0.3$$

$$PX = \text{conXA} \cdot PA + \text{conXB} \cdot PB + \text{conXC} \cdot PC$$

##Let conAX, conBX, and conCX be $P(A|X)$, $P(B|X)$ and $P(C|X)$ respectively.

$$\text{conAX} = (\text{conXA} \cdot PA) / PX$$

$$\text{conAX}$$

$$\text{conBX} = (\text{conXB} \cdot PB) / PX$$

$$\text{conBX}$$

$$\text{conCX} = (\text{conXC} \cdot PC) / PX$$

$$\text{conCX}$$

##Answer: $P(A|X) = 0.4426877$, $P(B|X) = 0.4624506$, $P(C|X) = 0.09486166$

##Answer3:

##Let H be the event of getting a heart attack.

##Let $P(H) = PH$

##Let DA and DB be the events of giving Drug A and Drug B respectively

$$PDA = 50/200$$

$$PDB = 50/200$$

$$PDADB = 100/200$$

$$PH = 0.80$$

$$\text{conHDA} = (1 - 0.35) \cdot PH \quad \text{##} P(H|DA)$$

$$\text{conHDB} = (1 - 0.20) \cdot PH \quad \text{##} P(H|DB)$$

$$\text{conHDADB} = (1 - 0.35) \cdot (1 - 0.20) \cdot PH \quad \text{##} P(H|DA \cap DB)$$

##To find: $P(DA \cap DB|H)$

$$\text{conDADBH} = (\text{conHDADB} \cdot PDADB) / (\text{conHDA} \cdot PDA + \text{conHDB} \cdot PDB + \text{conHDADB} \cdot PDADB)$$

$$\text{conDADBH}$$

##Answer: 0.4176707

##Answer4:

##a)

$$a4 = \text{ppois}(2, 4.2)$$

$$a4$$

$$\text{##} P(X \leq 2) = 0.210238$$

##b)

$P(X \geq 5) = 1 - P(X < 5)$ or $1 - P(X \leq 4)$

b4 = 1-ppois(4,4.2)

b4

$P(X \geq 5) = 0.410173$

##c)

c4 = dpois(8,4.2)

c4

$P(X=8) = 0.03601115$

##Answer5:

n_binomial = 30

p_binomial = 0.04

##a)

a5 = dbinom(25,n_binomial,p_binomial)

a5

$P(r=25) = 1.308245e-30$

##b)

b5 = dbinom(3,n_binomial,p_binomial)

b5

$P(r=3) = 0.08630368$

##c)

c5 = dbinom(5,n_binomial,p_binomial)

c5

$P(r=5) = 0.00525913$

CHI-SQUARE TESTS

##Answer1:

##Loading Dataset

```
data_1 <- read.csv(file.choose())
```

```
data_1
```

##EDA

```
head(data_1)
```

```
print(str(data_1))
```

```
View(data_1)
```

##Chi-square test

##Considering NE and NW

##H0: Mean(NE) = Mean(NW)

##Ha: Mean(NE) != Mean(NW)

```
chisq.test(data_1$NE,data_1$NW)
```

##Since, p-value > 0.05, we accept our null hypothesis.

##Considering NE and SE

##H0: Mean(NE) = Mean(SE)

##Ha: Mean(NE) != Mean(SE)

```
chisq.test(data_1$NE,data_1$SE)
```

##Since, p-value > 0.05, we accept our null hypothesis.

##Considering NE and SW

##H0: Mean(NE) = Mean(SW)

##Ha: Mean(NE) != Mean(SW)

```
chisq.test(data_1$NE,data_1$SW)
```

##Since, p-value > 0.05, we accept our null hypothesis.

##Considering NW and SE

##H0: Mean(NW) = Mean(SE)

##Ha: Mean(NW) != Mean(SE)

```
chisq.test(data_1$NW,data_1$SE)
```

##Since, p-value > 0.05, we accept our null hypothesis.

##Considering NW and SW

```
##H0: Mean(NW) = Mean(SW)
##Ha: Mean(NW) != Mean(SW)
chisq.test(data_1$NW,data_1$SW)
##Since, p-value > 0.05, we accept our null hypothesis.
```

```
##Considering SE and SW
```

```
##H0: Mean(SE) = Mean(SW)
##Ha: Mean(SE) != Mean(SW)
chisq.test(data_1$SE,data_1$SW)
##Since, p-value > 0.05, we accept our null hypothesis.
##Since for all regions, p-value > 0.05, the brand's share is evenly distributed
```

Answer2:

```
##Loading dataset survey
data()
View(survey)
data("survey")
print(str(survey))
```

```
##EDA
head(survey)
```

```
##Dataframe
survey_1 <- data.frame(survey)
survey_2 <- data.frame(survey_1$Smoke,survey_1$Exer)
survey_table <- table(survey_2)
survey_table
```

```
##Chi-square
##H0: Mean(smoke) = Mean(exer)
##Ha: Mean(smoke) != Mean(exer)
chisq.test(survey_table)
##since,p-value >0.05, we accept the null hypothesis.
```

```
##Correlation
cor(survey_table)
##They have strong correlation, and thus are dependent.
```

##Answer3:

```
##Loading dataset
data()
View(ChickWeight)
data("ChickWeight")
print(str(ChickWeight))
```

```
##EDA
head(ChickWeight)

##Dataframe
ChickWeight_1 <- data.frame(ChickWeight)
ChickWeight_2 <- data.frame(ChickWeight_1$Diet,ChickWeight_1$weight)
ChickWeight_2$weightcategory <-
ifelse(ChickWeight_2$ChickWeight_1.weight<median(ChickWeight_2$ChickWeight_1.weight),
"Low","High")
ChickWeight_table <-
table(ChickWeight_2$weightcategory,ChickWeight_2$ChickWeight_1.Diet)
ChickWeight_table

##Chi-square test
##H0: Mean(Diet) = Mean(Weight)
##Ha: Mean(Diet) != Mean(weight)
chisq.test(ChickWeight_table)
##Since, p-value < 0.05, we reject our null hypothesis

##Correlation
cor(ChickWeight_table)
##They are correlated, i.e. dependent.
```

##Answer4:

```
##Loading data
View(Arthritis)
data("Arthritis")
print(str(Arthritis))

##EDA
head(Arthritis)

##Dataframe
last1 <- data.frame(Arthritis)
last2 <- data.frame(last1$Treatment,last1$Improved)
last_table <- table(last2)
last_table

##Chi-Square test
##H0: Mean(Treatment) = Mean(Improved)
##H1: Mean(Treatment) != Mean(Improved)
chisq.test(last_table)
##X-squared = 13.055, p-value = 0.001463
##Since, p-value < 0.05, we reject our null hypothesis.
```