

## MAT2005 - Lab Assessment 5

---

Q1)

R code

```
> print("21BBS0059")
[1] "21BBS0059"
> field = c(rep("field1",1),rep("field2",1),rep("field3",1),rep("field4",1))
> variety = c(rep("varietyA",4),rep("varietyB",4),rep("varietyC",4),rep("varietyD",4))
> seed = c("C","A","B","D","B","D","A","C","A","C","D","B","D","B","C","A")
> freq = c(35,29,29,27,33,29,24,30,30,31,27,31,30,28,30,25)
> mydata = data.frame(variety,field,seed,freq)
> myfit = lm(freq ~ field + variety + seed, mydata)
> anova(myfit)
Analysis of Variance Table

Response: freq
  Df Sum Sq Mean Sq F value Pr(>F)
field      3   46.5  15.500  8.8571 0.01269 *
variety    3     7.5   2.500  1.4286 0.32411
seed       3   48.5  16.167  9.2381 0.01148 *
Residuals  6   10.5   1.750
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
>
```

Q2)  
R code

```
> print("21BBS0059")
[1] "21BBS0059"
> Input = c(
+ Before After
+ 7 7
+ 5 7
+ 6 5
+ 8 8
+ 7 7
+ 5 6
+ 4 6
+ 8 10
+ 9 10
+ 5 8
+ 7 9
+ 6 8
+ ")
> Data = read.table(textConnection(Input), header = TRUE)
> Data
   Before After
1      7     7
2      5     7
3      6     5
4      8     8
5      7     7
6      5     6
7      4     6
8      8    10
9      9    10
10     5     8
11     7     9
12     6     8

-- 
> wilcox.test(Data$Before, Data$After, pired = TRUE)
Wilcoxon rank sum test with continuity correction

data: Data$Before and Data$After
W = 43.5, p-value = 0.1001
alternative hypothesis: true location shift is not equal to 0

Warning message:
In wilcox.test.default(Data$Before, Data$After, pired = TRUE) :
  cannot compute exact p-value with ties
> wilcox.test(Data$Before, Data$After, pired = TRUE, conf.int = TRUE, conf.level = 0.95)

Wilcoxon rank sum test with continuity correction

data: Data$Before and Data$After
W = 43.5, p-value = 0.1001
alternative hypothesis: true location shift is not equal to 0
95 percent confidence interval:
-2.999952e+00  3.009914e-05
sample estimates:
difference in location
-1.000003

Warning messages:
1: In wilcox.test.default(Data$Before, Data$After, pired = TRUE, conf.int = TRUE,  :
  cannot compute exact p-value with ties
2: In wilcox.test.default(Data$Before, Data$After, pired = TRUE, conf.int = TRUE,  :
  cannot compute exact confidence intervals with ties
>
```

## Notebook -

Date \_\_\_\_\_  
Page \_\_\_\_\_

Pranav Murthy  
21BBS0059  
Pranav

MAT2005 DAB ASSESSMENT 5

Concerned code:-

(Q1) > field = c(rep("field1", 1), rep("field2", 1), rep("field3", 1), rep("field4", 1))  
> variety = c(rep("varietyA", 4), rep("varietyB", 4), rep("varietyC", 4),  
rep("varietyD", 4))  
> seed = c("C", "A", "D", "B", "D", "A", "C", "A", "C", "D", "B",  
"D", "B", "C", "A")  
> freq = c(35, 29, 29, 27, 33, 29, 24, 30, 30, 31, 27, 31, 30, 28,  
30, 25)  
> mydata = data.frame(\*variety\*, \*field\*, \*seed, freq)  
> myfit = lm(freq ~ field + variety + seed, mydata)  
> ~~anova~~ anova(myfit)

(Q2) > Input = ("  
+ Before After  
+ 7 7  
+ 5 7  
+ 6 5  
+ 8 8  
+ 7 7  
+ 5 6  
+ 4 6  
+ 8 10  
+ 9 10  
+ 5 8  
+ 7 9  
+ 6 8  
+ ")

Date \_\_\_\_\_  
Page \_\_\_\_\_

Pranav Murthy  
21BBS0059  
Pranav

> Data = read.table(textConnection(Input), header = TRUE)  
> Data  
> wilcox.test(Data\$Before, Data\$After, paired = TRUE)  
> wilcox.test(Data\$Before, Data\$After, paired = TRUE, conf.int = TRUE,  
+ conf.level = 0.95)

## MAT2005 Assessment 4

Q1

```

>
> print("21BBS0059 || Pranav Murthy")
[1] "21BBS0059 || Pranav Murthy"
> df1 = read.csv("/Users/pranav/Desktop/q1.csv")
> df1
   Program1 Program2 Program3 Program4
1         9       10      12       9
2        12       6      14       8
3        14       9      11      11
4        11       9      13       7
5        13      10      11       8
> r=c(t(as.matrix(df1)))
> r
[1]  9 10 12  9 12  6 14  8 14  9 11 11 11  9 13  7 13 10 11  8
> f=c("Program 1","Program 2","Program 3","Program 4")
> f
[1] "Program 1" "Program 2" "Program 3" "Program 4"
> k=4
> n=5
> tm=gl(k,1,n*k,factor(f))
> tm
[1] Program 1 Program 2 Program 3 Program 4 Program 1 Program 2 Program 3 Program 4
[9] Program 1 Program 2 Program 3 Program 4 Program 1 Program 2 Program 3 Program 4
[17] Program 1 Program 2 Program 3 Program 4
Levels: Program 1 Program 2 Program 3 Program 4
> crdfit=aov(r~tm)
> summary(crdfit)
    Df Sum Sq Mean Sq F value Pr(>F)
tm      3  54.95   18.32   7.045 0.00311 ***
Residuals 16  41.60    2.60
---
Signif. codes:  0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1
>
>
```

Q2

```

>
>
> data=c(24.6,20.5,27.6,16.1,16.1,24.8,27.2,28.7,22.6,14.9,16.9,22.4,38.4,39.4,36.7,19.5,15.3,26.2,28.4,30.9,34.8,14.0,17.6,22.5)
> blocks=gl(4,6)
> treatments=gl(6,1,24)
> rbdfit=aov(data~blocks+treatments)
> rbdfit
Call:
aov(formula = data ~ blocks + treatments)

Terms:
blocks treatments Residuals
Sum of Squares 219.4279  901.1921 229.6396
Deg. of Freedom     3          5         15

Residual standard error: 3.912711
Estimated effects may be unbalanced
> summary.aov(rbdfit)
    Df Sum Sq Mean Sq F value Pr(>F)
blocks      3  219.4   73.14   4.778  0.0157 *
treatments  5  901.2  180.24  11.773 9.28e-05 ***
Residuals   15  229.6   15.31
---
Signif. codes:  0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1
> print("Pranav Murthy || 21BBS0059")
[1] "Pranav Murthy || 21BBS0059"
> |
```

Q1)

$H_0$ : Mean =14.5

$H_1$ : Mean  $\neq$  14.5

R Code

```
> print("21BBS0059 || Pranav Murthy")
[1] "21BBS0059 || Pranav Murthy"
>
> m0=14.5
> xbar=13.6
> sigma=2.5
> n=40
> z=(xbar-m0)/(sigma/sqrt(n))
> z
[1] -2.27684
> alpha=0.05
> z.alpha=qnorm(1-alpha)
> z.alpha
[1] 1.644854
>
```

---

Interpretation

Since

$|z| > |z.alpha|$

hence  $H_0$  is rejected.

Mean is not 14.5kg

Q2)

$H_0$ : The proportion of rotten bananas in harvest stays below 11% a year

$H_1$ : The proportion of rotten bananas in harvest is above 11% a year

Code

```
> prop.test(33,213,0.11,alt="less",correct=FALSE)
```

1-sample proportions test without continuity correction

```
data: 33 out of 213, null probability 0.11
X-squared = 4.392, df = 1, p-value = 0.9819
alternative hypothesis: true p is less than 0.11
95 percent confidence interval:
 0.000000 0.200012
sample estimates:
      p 
0.1549296
```

>

---

Q3)

Code

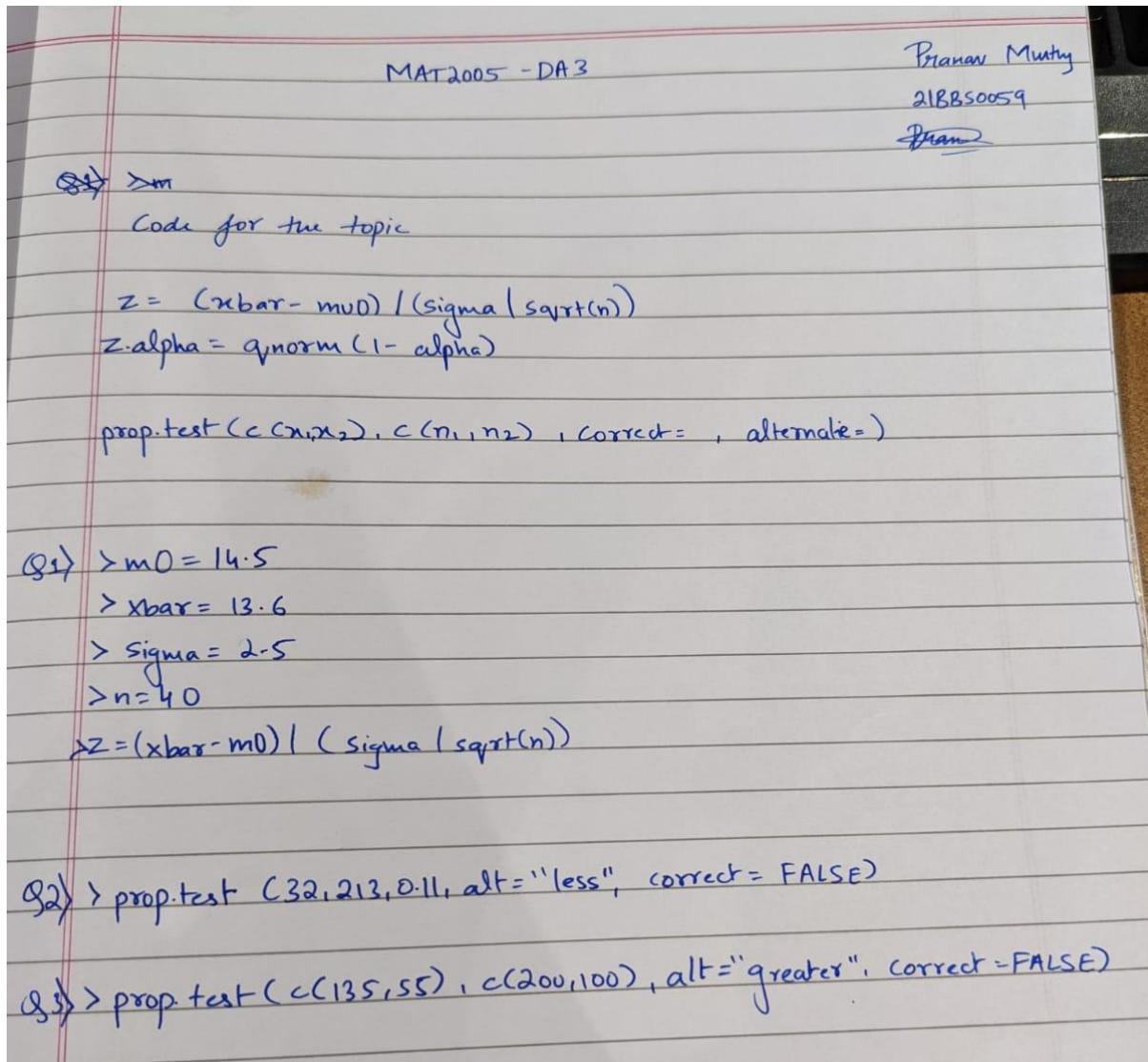
```
> prop.test(c(135,55),c(200,100),alt="greater",correct=FALSE)
```

2-sample test for equality of proportions without continuity correction

```
data: c(135, 55) out of c(200, 100)
X-squared = 4.4856, df = 1, p-value = 0.01709
alternative hypothesis: greater
95 percent confidence interval:
 0.02669505 1.00000000
sample estimates:
prop 1 prop 2
 0.675  0.550
```

&gt;

Notebook:



## MAT2005 – DA2

Code-

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Q1) Interpretation.

Let Price = Y, Quantity = X

iv)  $Y = 8.66359 - 0.02731X$

v)  $X = 205.14 - 14.74Y$

vi) Price when quantity = 160  
 $\Rightarrow X = 160$   
 $\Rightarrow Y = 8.66359 - 0.02731(160)$   
 $\therefore Y = 4.2940$

vii) Quantity when price = 7  
 $\Rightarrow Y = 7$   
 $\Rightarrow X = 205.14 - 14.74(7)$   
 $\therefore X = 101.9600$

Q2. Interpretation

$Y = -700.07 + 21.54^*X_1 - 14.45^*X_2$

When  $X_1 = 100$  &  $X_2 = 90$   
 $Y = -700.07 + 21.54(100) - 14.45(90)$   
 $\therefore Y = 153.4300$

## Interpretation-

Date \_\_\_\_\_  
Page \_\_\_\_\_

MAT2005 DA-2

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Q1. ~~Code~~ Code

```
> price = c(4.5, 5.5, 4.5, 4.5, 4.0, 5.5, 5.5, 6.5, 5.0, 5.5, 6.0, 4.5)
> quantity = c(125, 115, 140, 140, 150, 150, 130, 120, 130, 100, 105, 150)
> cor.test(price, quantity, method = "pearson")
> cor.test(price, quantity, method = "Kendall")
> cor.test(price, quantity, method = "spearman")
> fit = lm(quantity ~ price)
> fit
> fit = lm(quantity ~ price)
> fit
```

Q2) Code

```
> Y1 = c(1110, 180, 170, 1120, 1150, 190, 170, 1120, 1130, 1140)
> X1 = c(130, 140, 120, 150, 160, 140, 120, 160, 150, 135)
> X2 = c(111, 110, 117, 115, 119, 112, 118, 114, 99, 77)
> input_data = data.frame(Y1, X1, X2)
> input_data
> RegModel = lm(Y1 ~ X1 + X2)
> RegModel
> summary(RegModel)
```

## R console Code –

1.

```
> print("PRANAV MURTHY || 21BBS0059")
[1] "PRANAV MURTHY || 21BBS0059"
> price=c(4.5,5.5,4.5,4.5,4.0,5.5,5.5,6.5,5.0,5.5,6.0,4.5)
> quantity=c(125,115,140,140,150,150,130,120,130,100,105,150)
> cor.test(price,quantity,method="pearson")

  Pearson's product-moment correlation

data: price and quantity
t = -2.5959, df = 10, p-value = 0.02668
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
-0.88582895 -0.09527992
sample estimates:
cor
-0.634487

> cor.test(price,quantity,method="kendall")

  Kendall's rank correlation tau

data: price and quantity
z = -2.173, p-value = 0.02978
alternative hypothesis: true tau is not equal to 0
sample estimates:
tau
-0.5227084

Warning message:
In cor.test.default(price, quantity, method = "kendall") :
  Cannot compute exact p-value with ties
> cor.test(price,quantity,method="spearman")

  Spearman's rank correlation rho

data: price and quantity
S = 472.01, p-value = 0.02202
alternative hypothesis: true rho is not equal to 0
sample estimates:
rho
-0.650397

Warning message:
In cor.test.default(price, quantity, method = "spearman") :
  Cannot compute exact p-value with ties
> fit =lm(price~quantity)
> fit

Call:
lm(formula = price ~ quantity)

Coefficients:
(Intercept)     quantity
8.66359      -0.02731

> fit=lm(quantity~price)
> fit

Call:
lm(formula = quantity ~ price)

Coefficients:
(Intercept)       price
205.14          -14.74

> |
```

Interpretation –

Let Price = Y, Quantity = X

$$Y = 8.66359 - 0.02731X$$

$$X = 205.14 - 14.74Y$$

Price when Quantity = 160

$$\Rightarrow \text{Price} = 8.66359 - 0.02731 * 160$$

$$\Rightarrow \text{Price} = 4.2940$$

Quantity when Price = 7

$$\Rightarrow \text{Quantity} = 205.14 - 14.74 * 7$$

$$\Rightarrow \text{Quantity} = 101.9600$$

2.

```
> print("PRANAV MURTHY || 21BBS0059")
[1] "PRANAV MURTHY || 21BBS0059"
> Y=c(1110,180,170,1120,1150,190,170,1120,1130,1140)
> X1=c(130,140,120,150,160,140,120,160,150,135)
> X2=c(111,110,117,115,119,112,118,114,99,77)
> inout_data=data.frame(Y,X1,X2)
> input_data=data.frame(Y,X1,X2)
> input_data
   Y  X1  X2
1 1110 130 111
2  180 140 110
3  170 120 117
4 1120 150 115
5 1150 160 119
6  190 140 112
7  170 120 118
8 1120 160 114
9 1130 150  99
10 1140 135  77
> RegModel=lm(Y~X1 + X2, data=input_data)
> RegModel

Call:
lm(formula = Y ~ X1 + X2, data = input_data)

Coefficients:
            (Intercept)          X1           X2
              -700.07        21.54       -14.45

> summary(RegModel)

Call:
lm(formula = Y ~ X1 + X2, data = input_data)

Residuals:
    Min      1Q  Median      3Q     Max 
-545.67 -20.05  25.63 103.96 614.20 

Coefficients:
            Estimate Std. Error t value Pr(>|t|)    
(Intercept) -700.067   1614.841 -0.434   0.6777    
X1           21.541     8.687   2.480   0.0422 *  
X2          -14.455    10.024  -1.442   0.1925    
---
Signif. codes:  0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

Residual standard error: 380.6 on 7 degrees of freedom
Multiple R-squared:  0.5329, Adjusted R-squared:  0.3994 
F-statistic: 3.993 on 2 and 7 DF,  p-value: 0.06966
```

Interpretation –

$$Y = -700.07 + 21.54 * X1 - 14.45 * X2$$

When X1=100 and X2=90

$$\Rightarrow Y = -700.07 + 21.54 * 100 - 14.45 * 90$$

$$\Rightarrow Y = 153.4300$$

## MAT2005 DA1

1. Write all the basics of R-LAB in your note book .

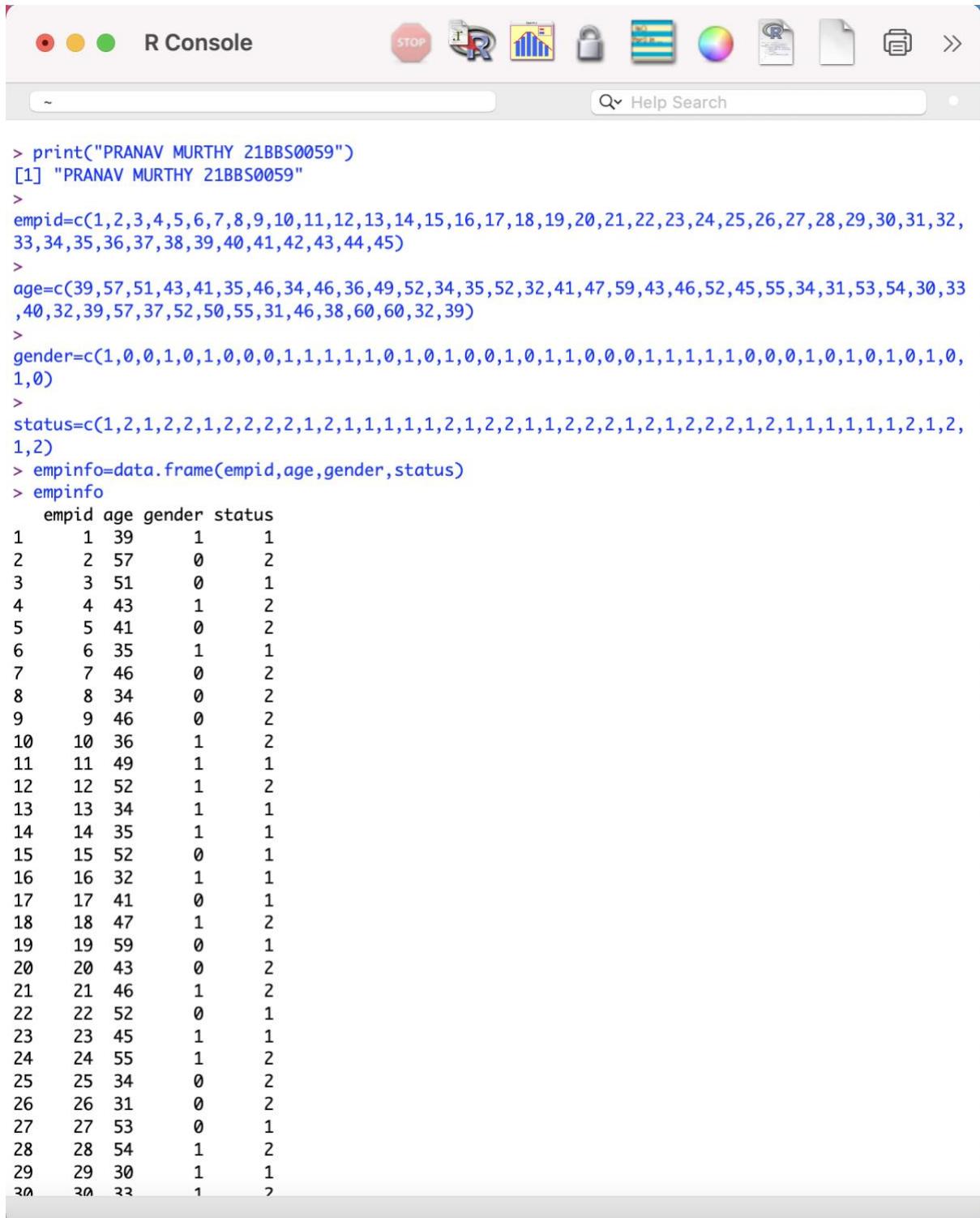
**CODE**

```
> print("21BBS0059 Pranav Murthy")
[1] "21BBS0059 Pranav Murthy"
> 31*78
[1] 2418
> 697/41
[1] 17
> x=39
> x
[1] 39
> y=22
> y
[1] 22
> z=x-y
> z
[1] 17
> log2(sqrt(2345))
[1] 5.597686
> x=10
> y=20
> x*y
[1] 200
> z=x*y
> z
[1] 200
> x=c(1,2,3,4,5)
> x
[1] 1 2 3 4 5
> 2+3
[1] 5
> 2-3
[1] -1
> 2*3
[1] 6
> 2/3
[1] 0.6666667
> 2*3-4+5/6
[1] 2.833333
> 2^3
[1] 8
> 2**0.5
[1] 1.414214
> 2**-0.5
[1] 0.7071068
> 2**3**2
[1] 512
> 2^(3^2)
[1] 512
> c(2,3,5,7)**2
[1] 4 9 25 49
> c(1,2,3,4,5,6,7)**c(2,3,4)
```

```
[1] 1 8 81 16 125 1296 49
Warning message:
In c(1, 2, 3, 4, 5, 6, 7)^c(2, 3, 4) :
  longer object length is not a multiple of shorter object length
> c(1,2,3,4,5,6)**c(2,3,4)
[1] 1 8 81 16 125 1296
> c(1,2,3,4,5)%%2
[1] 1 0 1 0 1
> c(1,2,3,4,5)%%2
[1] 0 1 1 2 2
> max(1.2,3.4,-7.8)
[1] 3.4
> min(1.2,3.4,-7.8)
[1] -7.8
> abs(-4)
[1] 4
> sqrt(4)
[1] 2
> prod(c(2,3,4,5,6))
[1] 720
> round(1.83)
[1] 2
> round(1.23)
[1] 1
> log(10)
[1] 2.302585
> log10(10)
[1] 1
> log(9,base=3)
[1]
```

2. Create a data frame for 45 employees consisting the details for (i) empid(ii)age (iii) genderand (iv) status and perform thesummary of statistics which includes (i) tables for the data (ii) bar diagram and (iii) pie chart (iv) scatter plot for age .

Code



The screenshot shows the R Console interface. The title bar says "R Console". Below the title bar is a toolbar with various icons: STOP, PREFERENCES, HISTOGRAM, LOCK, COLOR, HELP, SEARCH, and PRINT. A search bar at the bottom right contains the text "Help Search". The main area of the console displays the following R code and its output:

```

> print("PRANAV MURTHY 21BBS0059")
[1] "PRANAV MURTHY 21BBS0059"
>
empid=c(1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,
33,34,35,36,37,38,39,40,41,42,43,44,45)
>
age=c(39,57,51,43,41,35,46,34,46,36,49,52,34,35,52,32,41,47,59,43,46,52,45,55,34,31,53,54,30,33
,40,32,39,57,37,52,50,55,31,46,38,60,60,32,39)
>
gender=c(1,0,0,1,0,1,0,0,0,1,1,1,1,0,1,0,1,0,1,0,1,1,1,1,1,0,0,0,1,1,1,1,1,1,0,0,0,1,0,1,0,1,0,
1,0)
>
status=c(1,2,1,2,2,1,2,2,2,1,2,1,1,1,1,1,2,1,2,2,1,1,2,2,2,1,2,1,2,2,2,1,2,1,1,1,1,1,1,2,1,2,
1,2)
> empinfo=data.frame(empid,age,gender,status)
> empinfo
  empid age gender status
1      1  39      1     1
2      2  57      0     2
3      3  51      0     1
4      4  43      1     2
5      5  41      0     2
6      6  35      1     1
7      7  46      0     2
8      8  34      0     2
9      9  46      0     2
10    10  36      1     2
11    11  49      1     1
12    12  52      1     2
13    13  34      1     1
14    14  35      1     1
15    15  52      0     1
16    16  32      1     1
17    17  41      0     1
18    18  47      1     2
19    19  59      0     1
20    20  43      0     2
21    21  46      1     2
22    22  52      0     1
23    23  45      1     1
24    24  55      1     2
25    25  34      0     2
26    26  31      0     2
27    27  53      0     1
28    28  54      1     2
29    29  30      1     1
30    30  22      1     2

```

R Console

STOP QR Help Search

```
> empinfo
  empid age gender status
1      1  39      1      1
2      2  57      0      2
3      3  51      0      1
4      4  43      1      2
5      5  41      0      2
6      6  35      1      1
7      7  46      0      2
8      8  34      0      2
9      9  46      0      2
10     10 36      1      2
11     11 49      1      1
12     12 52      1      2
13     13 34      1      1
14     14 35      1      1
15     15 52      0      1
16     16 32      1      1
17     17 41      0      1
18     18 47      1      2
19     19 59      0      1
20     20 43      0      2
21     21 46      1      2
22     22 52      0      1
23     23 45      1      1
24     24 55      1      2
25     25 34      0      2
26     26 31      0      2
27     27 53      0      1
28     28 54      1      2
29     29 30      1      1
30     30 33      1      2
31     31 40      1      2
32     32 32      1      2
33     33 39      0      1
34     34 57      0      2
35     35 37      0      1
36     36 52      1      1
37     37 50      0      1
38     38 55      1      1
39     39 31      0      1
40     40 46      1      1
41     41 38      0      2
42     42 60      1      1
43     43 60      0      2
44     44 32      1      1
45     45 39      0      2
> summary(empinfo)
```

R Console

```

45 45 39 0 2
> summary(empinfo)
   empid      age      gender      status
  Min.   : 1   Min.   :30.00   Min.   :0.0000   Min.   :1.000
  1st Qu.:12  1st Qu.:35.00  1st Qu.:0.0000  1st Qu.:1.000
  Median :23  Median :43.00  Median :1.0000  Median :1.000
  Mean   :23  Mean   :43.84  Mean   :0.5111  Mean   :1.489
  3rd Qu.:34  3rd Qu.:52.00  3rd Qu.:1.0000  3rd Qu.:2.000
  Max.   :45  Max.   :60.00  Max.   :1.0000  Max.   :2.000
> bargraph=table(empinfo$gender,empinfo$status)
> barplot(table5,beside=T,xlim=c(1,45),ylim=c(0,5),c
ol=c("blue","red"),legend=rownames(bargraph))
Error: unexpected symbol in "barplot(table5,beside=T,xlim=c(1,45),ylim=c(0,5),c ol"
> piechart=table(empinfo$gender)
> pie(piechart)
> summary(empinfo$age)
   Min. 1st Qu. Median Mean 3rd Qu. Max.
  30.00 35.00 43.00 43.84 52.00 60.00
> empinfo$gender=factor(empinfo$gender,labels=c("male","female"))
> pie(piechart)
> pie(piechart)
> piechart=table(empinfo$gender)
> pie(piechart)
> empinfo$status=factor(empinfo$status,labels=c("staff" ,"faculty"))
> empinfo
   empid age gender status
1       1 39 female staff
2       2 57 male faculty
3       3 51 male staff
4       4 43 female faculty
5       5 41 male faculty
6       6 35 female staff
7       7 46 male faculty
8       8 34 male faculty
9       9 46 male faculty
10      10 36 female faculty
11      11 49 female staff
12      12 52 female faculty
13      13 34 female staff
14      14 35 female staff
15      15 52 male staff
16      16 32 female staff
17      17 41 male staff
18      18 47 female faculty
19      19 59 male staff
20      20 43 male faculty
21      21 46 female faculty
22      22 52 male faculty

```

R Console

```

45 45 39 0 2
> summary(empinfo)
   empid      age      gender      status
  Min.   : 1   Min.   :30.00   Min.   :0.0000   Min.   :1.000
  1st Qu.:12  1st Qu.:35.00  1st Qu.:0.0000  1st Qu.:1.000
  Median :23  Median :43.00  Median :1.0000  Median :1.000
  Mean   :23  Mean   :43.84  Mean   :0.5111  Mean   :1.489
  3rd Qu.:34  3rd Qu.:52.00  3rd Qu.:1.0000  3rd Qu.:2.000
  Max.   :45  Max.   :60.00  Max.   :1.0000  Max.   :2.000
> bargraph=table(empinfo$gender,empinfo$status)
> barplot(table5,beside=T,xlim=c(1,45),ylim=c(0,5),c
ol=c("blue","red"),legend=rownames(bargraph))
Error: unexpected symbol in "barplot(table5,beside=T,xlim=c(1,45),ylim=c(0,5),c ol"
> piechart=table(empinfo$gender)
> pie(piechart)
> summary(empinfo$age)
   Min. 1st Qu. Median Mean 3rd Qu. Max.
  30.00 35.00 43.00 43.84 52.00 60.00
> empinfo$gender=factor(empinfo$gender,labels=c("male","female"))
> pie(piechart)
> pie(piechart)
> piechart=table(empinfo$gender)
> pie(piechart)
> empinfo$status=factor(empinfo$status,labels=c("staff" ,"faculty"))
> empinfo
   empid age gender status
1       1 39 female staff
2       2 57 male faculty
3       3 51 male staff
4       4 43 female faculty
5       5 41 male faculty
6       6 35 female staff
7       7 46 male faculty
8       8 34 male faculty
9       9 46 male faculty
10      10 36 female faculty
11      11 49 female staff
12      12 52 female faculty
13      13 34 female staff
14      14 35 female staff
15      15 52 male staff
16      16 32 female staff
17      17 41 male staff
18      18 47 female faculty
19      19 59 male staff
20      20 43 male faculty
21      21 46 female faculty
22      22 52 male faculty

```

The screenshot shows an R console window titled "R Console". The menu bar includes "File", "Edit", "View", "Insert", "Plot", "Tools", "Help", and "Search". Below the menu is a toolbar with icons for Stop, Help, Search, and others. The main area displays the following R session:

```
> empinfo
   empid age gender status
1      1  39 female staff
2      2  57 male faculty
3      3  51 male staff
4      4  43 female faculty
5      5  41 male faculty
6      6  35 female staff
7      7  46 male faculty
8      8  34 male faculty
9      9  46 male faculty
10     10 36 female faculty
11     11 49 female staff
12     12 52 female faculty
13     13 34 female staff
14     14 35 female staff
15     15 52 male staff
16     16 32 female staff
17     17 41 male staff
18     18 47 female faculty
19     19 59 male staff
20     20 43 male faculty
21     21 46 female faculty
22     22 52 male staff
23     23 45 female staff
24     24 55 female faculty
25     25 34 male faculty
26     26 31 male faculty
27     27 53 male staff
28     28 54 female faculty
29     29 30 female staff
30     30 33 female faculty
31     31 40 female faculty
32     32 32 female faculty
33     33 39 male staff
34     34 57 male faculty
35     35 37 male staff
36     36 52 female staff
37     37 50 male staff
38     38 55 female staff
39     39 31 male staff
40     40 46 female staff
41     41 38 male faculty
42     42 60 female staff
43     43 60 male faculty
44     44 32 female staff
45     45 39 male faculty
> statuspie=table(empinfo$status)
```

R Console

```

37 37 50 male staff
38 38 55 female staff
39 39 31 male staff
40 40 46 female staff
41 41 38 male faculty
42 42 60 female staff
43 43 60 male faculty
44 44 32 female staff
45 45 39 male faculty
> statuspie=table(empinfo$status)
> pie(statuspie)
> plot(empinfo$age,empinfo$, main="Age vs gender", xlab="age", ylab="gender", col="red")
Error: unexpected ',' in "plot(empinfo$age,empinfo$,"
> plot(empinfo$age,empinfo$gender, main="Age vs gender", xlab="age", ylab="gender", col="red")
> plot(empinfo$age,y, main="Age vs gender", xlab="age", ylab="gender", col="red")
Error in xy.coords(x, y, xlabel, ylabel, log) :
  'x' and 'y' lengths differ
> plot(empinfo$age,empinfo$empid, main="Age vs id", xlab="age", ylab="empid", col="red")
> tablebar=table(empinfo$gender,empinfo$status)
> barplot(tablebar,beside=T, xlim=c(1,15), ylim=c(0,15),col=c("blue","red"),
legend=rownames(tablebar))
Error: unexpected ',' in "barplot(tablebar,beside=T, xlim=c(1,15), ylim=c(0,15),col=c("blue",
> barplot(tablebar,beside=T, xlim=c(1,15), ylim=c(0,15),col=c("blue","red"),
legend=rownames(tablebar))
> barplot(tablebar,beside=T, xlim=c(1,45), ylim=c(0,15),col=c("blue","red"),
legend=rownames(tablebar))
> barplot(tablebar,beside=T, xlim=c(1,15), ylim=c(0,15),col=c("blue","red"),
legend=rownames(tablebar))
> boxplot(empinfo$age~empinfo$status,col=c("red","blue"))
> plot(empinfo$age,type="l",main="age of subjects",xlab="empid",ylab="age in years",col="blue")
> onewaytable=table(empinfo$gender)
> onewaytable

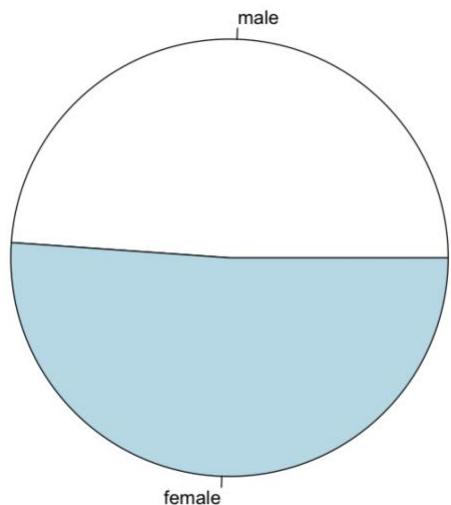
male female
 22    23
> onewaytablestat=table(empinfo$status)
> onewaytablestat

staff faculty
 23     22
> twowaytable=table(empinfo$gender,empinfo$status)
> twowaytable

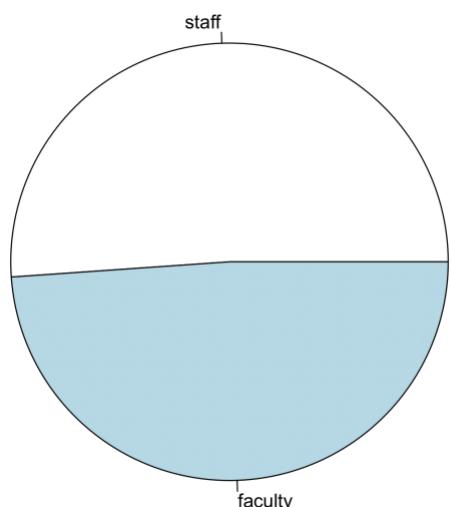
      staff faculty
male      10      12
female    13      10
>

```

Pie Chart  
Male and Female

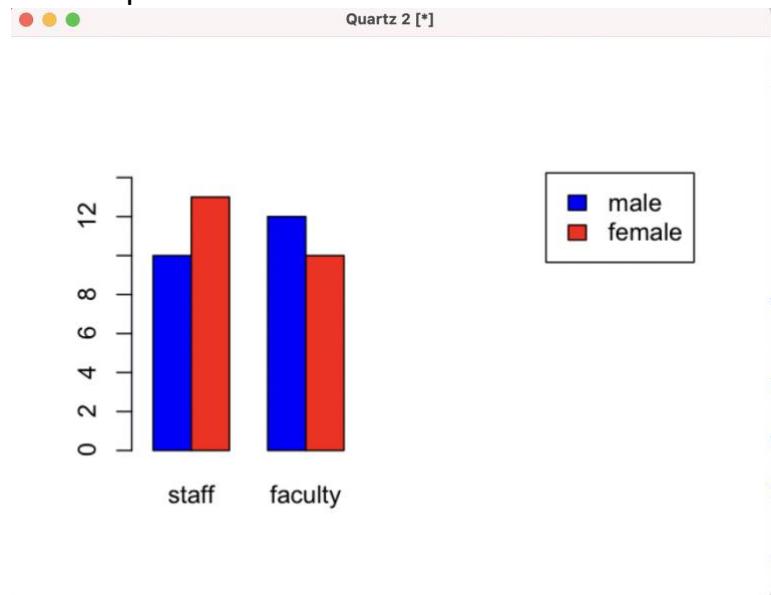
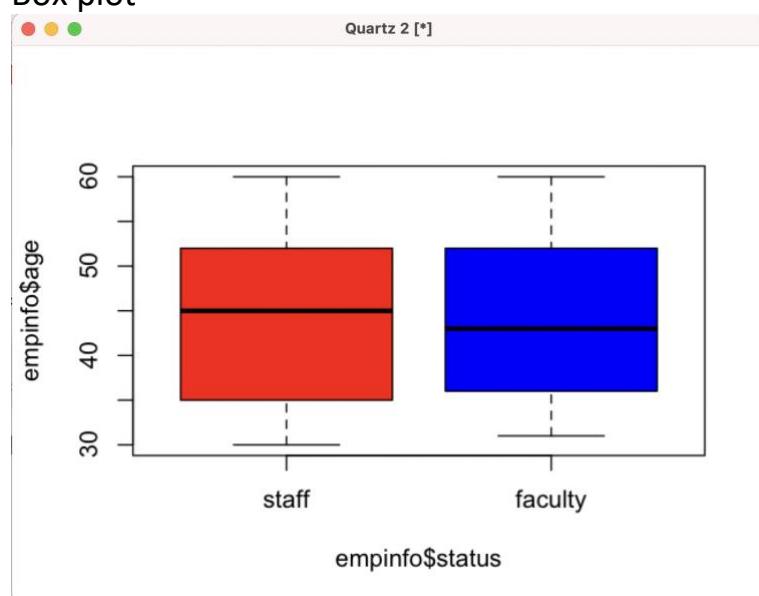


Status



Scatter plot  
Age vs empid



**Bar Graph****Box plot****Line Graph**