MAT 5030 Chapter 4: Descriptive Statistics and Graphics

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Tips

A list of datasets included by default in R package:

http://stat.ethz.ch/R-manual/R-patched/library/datasets/html/00Index.html

Summary Statistics for a Single Group

The 'summary' function summarize a data frame.

```
> Loan
  Inst Rate YTM Amount
       5.6
             15 100000
    BΑ
 Citi 4.7
             15 150000
  Citi 7.5
             30 310000
    BA 5.1
             30 510000
 Chase 4.2
                90000
    BA 2.9
              5 190000
 Chase 6.6 30 450000
> summary(Loan)
```

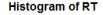
Inst	Rate	YTM	Amount
BA :3	Min. :2.900	Min. : 5.00	Min. : 90000
Chase:2	1st Qu.:4.450	1st Qu.:11.00	1st Qu.:125000
Citi :2	Median :5.100	Median :15.00	Median :190000
	Mean :5.229	Mean :18.86	Mean :257143
	3rd Qu.:6.100	3rd Qu.:30.00	3rd Qu.:380000
	Max. :7.500	Max. :30.00	Max. :510000

Histogram

- Use the 'breaks' option to customize break points of a histogram.
- Use the 'freq=F' option to make a density histogram. With this, a histogram can be overlaid with density curves.

```
RT < - rt(1000, df=5)
## Figure 1
hist(RT)
## Figure 2
B <- seq(floor(min(RT)), ceiling(max(RT)), by =0.5)
  # break points
hist(RT, breaks=B)
## Figure 3
hist(RT, breaks=B,
ylim=c(0,0.5), freq=F, xlab="",ylab="")
par(new=T)
X < -0.02*(-100:100) # x-coordinates of density curve
plot(X. dt(X.df=5).
xlim = range(B), ylim=c(0,0.5), type="l",
xlab="",ylab="Relative Frequency")
par(new=F)
```

Figure 1



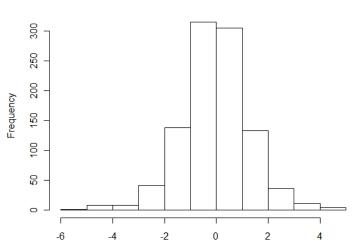
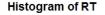


Figure 2



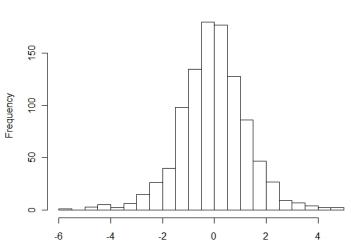
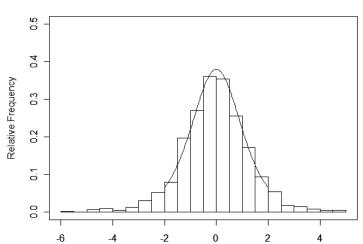


Figure 3

Histogram of RT



Empirical Cumulative Distribution function

Let \boldsymbol{X} be a random variable. The cumulative distribution function (CDF) \boldsymbol{F} is:

$$F(x)=P(X\leq x).$$

When x_1, \dots, x_n are observed, the **empirical CDF** \hat{F} is defined by:

$$\hat{F}(x) = \frac{\text{Number of observations with } (x_i \leq x)}{n}$$

The empirical CDF is a right-continuous function which is flat except for the points x_1, \dots, x_n .

Example 1:

Suppose we have 4 observations $\{-1, 0.1, 0.5.1.3\}$,

$$\hat{F}(X) = 0 \quad (if \quad x < -1)$$
 $\hat{F}(X) = 1/4 \quad (if \quad -1 \le x < 0.1)$
 $\hat{F}(X) = 2/4 \quad (if \quad 0.1 \le x < 0.5)$
 $\hat{F}(X) = 3/4 \quad (if \quad 0.5 \le x < 1.3)$
 $\hat{F}(X) = 1 \quad (if \quad 1.3 \le x)$

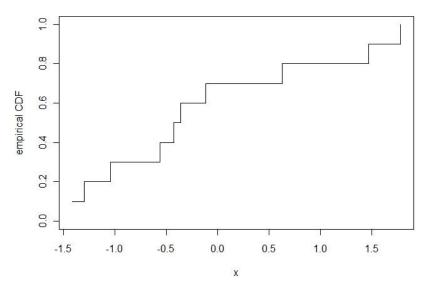
Example 2:

Generate 10 random numbers which follows standard normal, and draw an empirical CDF.

```
X <- sort(rnorm(10)) # 10 r.v.s, increasing order
n <- length(X)
# Figure 1
plot(X, (1:n)/n, type="s", ylim=c(0,1),xlab="x", ylab= "empirical CDF")</pre>
```

Note: type="s" generates a step function.

Figure 1



 $\hat{F}(x)$ must be 0 when x is very small, 1 when x is very large.

```
# Figure 2
X1 <- c(min(X)-1, sort(X),max(X)+1)
Oldpar <- par()  # store current graphic parameters as 'Oldpar'

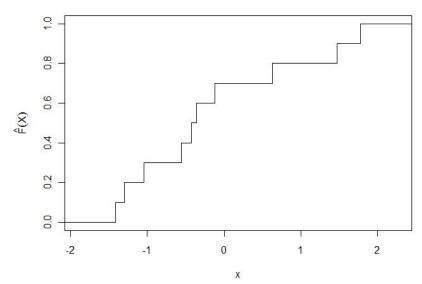
par(mai= c(1.02, 1, 0.82, 0.42))
    # set margin size (bottom, left, top, right)
plot(X1, 0:(n+1)/n,
type="s", ylim=c(0,1),xlim=c(min(X)-0.5,max(X)+0.5),
xlab="x", ylab=expression(paste(hat(F), (X)))
)

par(Oldpar) # return to the old parameters</pre>
```

Note:

- 'par()' represents all graphic parameters. See 'help(par)' for details.
- 'expression' and 'paste' functions are used for mathematical annotation. see 'help(plotmath)' for details.

Figure 2



Q-Q Plot

Let x_1, \dots, x_n be ordered observations, and F be the hypothetical CDF for the observations. The plot $(F^{-1}(\frac{i}{n+1}), x_i)$ $(i = 1, \dots, n)$ is called a Q-Q plot.)

If the observations come from the hypothetical distribution, approximately $F(x_i) = \frac{i}{n+1}$ (or equivalently $x_i = F^{-1}(\frac{i}{n+1})$). Hence, the Q-Q plot is roughly on the straight line y = x. So the Q-Q plot is used to see if the observations fit well to a hypothetical (e.g., normal) distribution.

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Example 1:

To check if $\{-1, 0.1, 0.5.1.3\}$ come from standard normal, we have to check if

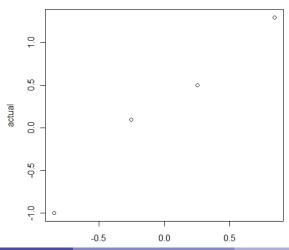
$$\left(F^{-1}\left(\frac{1}{5}\right), -1\right), \left(F^{-1}\left(\frac{2}{5}\right), 0.1\right), \left(F^{-1}\left(\frac{3}{5}\right), 0.5\right), \left(F^{-1}\left(\frac{4}{5}\right), 1.3\right)$$
 (1)

are close to y = x.

Code for Example 1:

```
Y <- c(-1, 0.1, 0.5, 1.3)
X <- qnorm((1:4)/5)
plot(X,Y,xlab="theoretical",ylab="actual")
```

Normal Q-Q plot for 4 observations





Example 2:

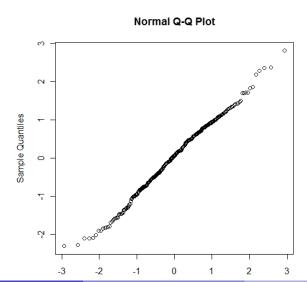
Generate 300 random numbers from a standard normal distribution, then create a normal Q-Q plot.

```
Y <- rnorm(300) > qqnorm(Y)
```

Since the observations are actually from a standard normal distribution, points should be approximately on the line y = x.

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normal Q-Q plot for normal observations



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Example 2:

Generate 300 random numbers from a normal distribution $N(10,5^2)$, then create a normal Q-Q plot.

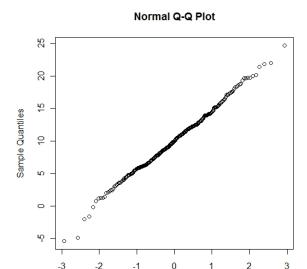
```
Y <- rnorm(300, mean=10, sd=5) qqnorm(Y)
```

The observations are almost on the straight line: y = 5x + 10.

Fact:

Observations follow normal \Leftrightarrow all points are roughly on a straight line.

Normal Q-Q plot for normal observations



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Exercise 1:

The *t*-distribution is known as a heavy-tailed distribution, which have more extreme values than normal distributions.

Generate 300 random numbers from t_3 distribution, and create normal Q-Q plot.

Exercise 2:

A (continuous) uniform distribution (f(x) = 1/(b-a) on [a, b]) have upper/lower limits, so there are no extreme values.

Generate 300 random numbers from a uniform distribution on (0, 1), and create normal Q-Q plot.

Summary Statistics by groups

In addition to the 'apply' family in Chapter 1 slides, 'aggregate' and 'by' functions are useful to summarize grouped data.

Summary Statistics by groups

'aggregate'

```
> CO2
 Plant
         Type Treatment conc uptake
   Qn1 Quebec nonchilled
                          95
                               16.0
   Qn1 Quebec nonchilled 175 30.4
   On1 Ouebec nonchilled 250 34.8
... (truncated) ...
83 Mc3 Mississippi chilled
                              675
                                    18.9
84
    Mc3 Mississippi chilled 1000
                                    19.9
> aggregate(CO2, CO2["Type"], mean)
        Type Plant Type Treatment conc uptake
      Ouebec .
               NA NA
                              NA 435 33.54286
2 Mississippi NA NA
                              NA 435 20.88333
Warning messages:
1: In mean.default(X[[1L]], ...):
 argument is not numeric or logical: returning NA
```

Summary Statistics by groups

```
'by'
```

```
> by(CO2, CO2["Type"], mean)
Type: Quebec
   Plant Type Treatment conc uptake
     NΑ
             NΑ
                NA 435.00000 33.54286
Type: Mississippi
   Plant Type Treatment conc uptake
                NA 435.00000 20.88333
     NΑ
             NA
> by(CO2, CO2["Type"], summary)
Type: Ouebec
   Plant
             Type
                           Treatment conc uptake
Qn1 :7 Quebec :42 nonchilled:21 Min. : 95
                                                 Min. : 9.30
Qn2 :7 Mississippi: 0 chilled :21 1st Qu.: 175 1st Qu.: 30.32
Qn3 :7
                                     Median : 350
                                                 Median : 37.15
Qc1 :7
                                     Mean : 435 Mean : 33.54
0c3 :7
                                     3rd Ou.: 675
                                                  3rd Ou.: 40.15
0c2 :7
                                     Max. :1000
                                                 Max. :45.50
(Other):0
<Truncated>
```

Box plot

Suppose we want to describe the sepal length in 'iris' data by species.

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species		
1	5.1	3.5	1.4	0.2	setosa		
2	4.9	3.0	1.4	0.2	setosa		
3	4.7	3.2	1.3	0.2	setosa		
(truncated)							
51	7.0	3.2	4.7	1.4	versicolor		
52	6.4	3.2	4.5	1.5	versicolor		
(truncated)							
149	6.2	3.4	5.4	2.3	virginica		
150	5.9	3.0	5.1	1.8	virginica		

Box plot (Continued)

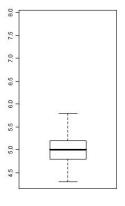
A box plot is a graphical way to describe the 1st quartile, median, 3rd quartile and outliers etc. We may want to lay out 3 box plots to compare these.

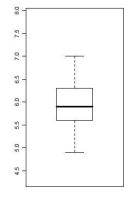
Sample R code 1:

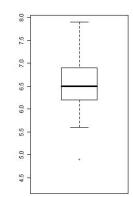
```
D1 <- subset(iris, Species=="setosa")
D2 <- subset(iris, Species=="versicolor")
D3 <- subset(iris, Species=="virginica")

par(mfrow = c(1,3))  # align 3 figures as 1 x 3 by row
R <- range(iris$Sepal.Length)
boxplot(D1$Sepal.Length, ylim = R)
boxplot(D2$Sepal.Length, ylim = R)
boxplot(D3$Sepal.Length, ylim = R)
par(mfrow = c(1,1))  # return to single figure</pre>
```

Sample R code 1:







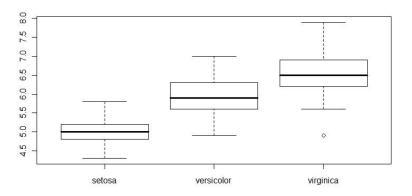
Sample R code 2:

boxplot(iris\$Sepal.Length ~ iris\$Species)

Sample R code 3:

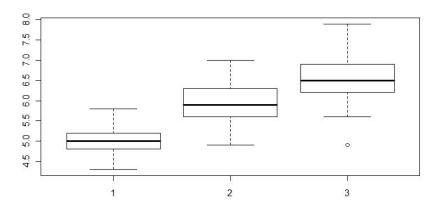
boxplot(D1\$Sepal.Length, D2\$Sepal.Length,D3\$Sepal.Length)

Sample R code 2:



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Sample R code 3:



A contingency table in 'table' format can be transformed to a list of frequencies in 'data frame' format.

A 'table' object in R is basically the same as 'matrix'. When you transform it to a data frame, it has to be a 'table'.

How to make a table?

```
> matrix(c(1,2,3,4),2,2)
     [,1] [,2]
[1,] 1 3
[2.] 2 4
> M \leftarrow matrix(c(1,2,3,4),2,2)
> colnames(M) <- c("A", "B") # assign column names
> rownames(M) <- c("C", "D") # assign row names
> names(dimnames(M)) <- c("Y","X")</pre>
 # assign category names for rows and columns
> M
   X
Y A B
 C 1 3
 D 2 4
```

4 D B

How to transform a table to a data frame?

How to summarize a data frame?

Motor Cars Dataset in R package:

```
> str(mtcars)
               32 obs. of 11 variables:
'data.frame':
 $ mpg : num 21 21 22.8 21.4 18.7 18.1 14.3 24.4 22.8 19.2 ...
 $ cvl : num
            6646868446...
 $ disp: num
             160 160 108 258 360 ...
 $ hp : num
             110 110 93 110 175 105 245 62 95 123 ...
             3.9 3.9 3.85 3.08 3.15 2.76 3.21 3.69 3.92 3.92 ...
 $ drat: num
 $ wt : num
             2.62 2.88 2.32 3.21 3.44 ...
 $ qsec: num
             16.5 17 18.6 19.4 17 ...
             0 0 1 1 0 1 0 1 1 1 ...
      : num
      : num
             1 1 1 0 0 0 0 0 0 0 ...
 $ gear: num
             4 4 4 3 3 3 3 4 4 4 ...
 $ carb: num
             4 4 1 1 2 1 4 2 2 4 ...
```

Tables

How to summarize a data frame? - 1

```
> attach(mtcars)
> table(gear) # summarize by gear
gear
3 4 5
15 12 5
> table(gear, carb) # summarize by gear and carb
    carb
gear 1 2 3 4 6 8
   3 3 4 3 5 0 0
   4 4 4 0 4 0 0
   5 0 2 0 1 1 1
```

Tables

How to summarize a data frame? – 2

```
> xtabs(~ gear + carb, data=mtcars)
    # same as table(gear, carb)
    carb
gear 1 2 3 4 6 8
    3 3 4 3 5 0 0
    4 4 4 0 4 0 0
    5 0 2 0 1 1 1
```

Tables

How to summarize a data frame? - 3

Function	Description
ftable	Similar to 'xtabs'. It makes a flat (2-dimensional)
	table even when there are more than 2 variables.
margin.table	Sum up counts by rows (or columns).
prop.table	Transform a table of counts into a table of proportions.

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Barplot

Dataset:

```
> D <- HairEyeColor[,,1]
  # Hair/Eye color of statistics student (male)
> D
      Eye
Hair
       Brown Blue Hazel Green
 Black
          32
               11
                     10
       53
               50 25 15
 Brown
 Red 10 10
 Blond
           3
               30
                            8
> D.E <- margin.table(D,2) # Students by Eye
> D.E.
Eye
Brown
      Blue Hazel Green
  98
       101
              47
                    33
```

Barplot

```
## barplot 1
barplot(D.H, col="pink") # students by eye

## barplot 2
Col <- c("black","brown","red",colors()[78])
barplot(D,col=Col, main="Students by Eye Color")
legend(3, 100,
c("Hair:Black","Hair:Brown","Hair:Red","Hair:Blond"),
col=Col, pch=15)</pre>
```

Note:

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- Type "colors()" to see all ready-made colors in R. (cf. http://research.stowers-institute.org/efg/R/Color/Chart/).
- "pch" is point character, and "pch=15" is a black square.

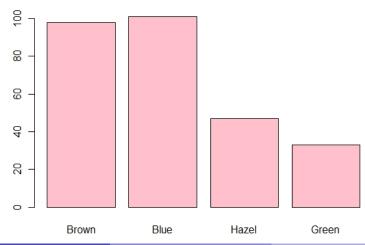
4 ロ ト 4 템 ト 4 분 ト 4 분 ト 9 은 연

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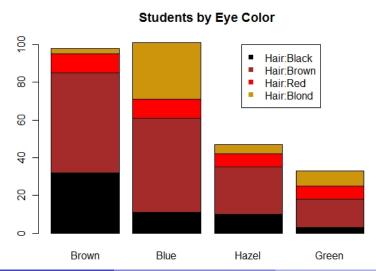
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Example: Barplot 1



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Example: Barplot 2



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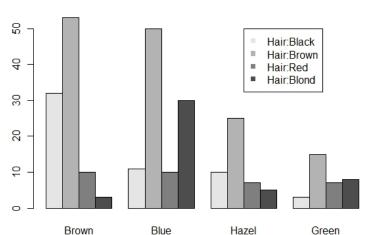
Exercise

The "HairEyeColor[,,2]" includes data for female. Create a barplot similar to the previous page, but interchange the role of eye and hair color.

```
## barplot 3
Col2 <- c("grey90","grey70","grey50","grey30")
barplot(D,col=Col2, xlab="Eye",
beside=T, main="Students by Eye & Hair Color")
legend(13, 50,
c("Hair:Black","Hair:Brown","Hair:Red","Hair:Blond"),
col= Col2, pch=15)</pre>
```

Example: Barplot 3

Students by Eye & Hair Color



Pie Chart

```
pie(D.E,
  col=c("brown","blue", colors()[146],"green"),
  main="Students by Eye Color")
```

Note:

 To start at 12 o'clock and move clockwise, add an option "clockwise=T".

Students by Eye Color

