

# **uRProgramming**

**By**

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**Created on Sun Oct 29 22:02:18  
2017**

# Table of Contents

<b>1 Introduction.....</b>	<b>4</b>
<b>2 Environment Setup for R programming Developers.....</b>	<b>4</b>
2.1 Environment setup.....	4
2.2 Using Version Control System.....	4
2.3 Using IDE to develop R scripts.....	4
2.4 Writing functions.....	4
2.5 Create R package System.....	4
<b>3 Simple Data Types and Objects.....</b>	<b>5</b>
<b>4 Control Structures.....</b>	<b>7</b>
4.1 If statements.....	7
4.2 If then Else statements.....	9
4.3 Switch statements.....	9
<b>5 Loop Structures.....</b>	<b>11</b>
5.1 For loop.....	11
5.2 While loop.....	11
<b>6 Data Structures.....</b>	<b>13</b>
6.1 Data Frame.....	15
6.2 Vector.....	16
6.3 Matrix.....	17
6.4 List.....	17
<b>7 Simple Statistical Analysis.....</b>	<b>18</b>
<b>8 Graphic Outputs.....</b>	<b>20</b>
<b>9 Simple Table Outputs.....</b>	<b>20</b>
<b>10 Reference.....</b>	<b>20</b>

# **Acknowledgements**

I would love thank my wife Sevtap Otles putting up with me while I was working on this book. I would not meet with R programming, if I was not hired by Dr. Paul Carbone who has shown how to be dedicated and sincere for his colleagues.

It would have been impossible to reach wherever I am at without my mom's trust and support. She (Zulfinaz Otles) influenced me and others to reach our education goals to be successful in life. She constantly worries the lack of educational opportunities for the people and children of Turkey, she is so concerned about education cost which is financially stressing the families.

My grand father (Ethem Demirkol) has taught me to work hard, be honest and treat all the people equal.

Finally, I would love thank to both of my sons (Erkin and Arel) for their sincerity, honesty, life choices and support to my wife and myself.

Thanks to all supporters, readers of this book and contributors to society.

# 1 Introduction

The idea is to create simple book, it will flow well and also easy to understand. Many people don't have much time to start a R programming and don't have experience other software engineering tools. Here in this book my aim is to start simple and built on using some software technologies.

## 2 Environment Setup for R programming Developers

### 2.1 *Environment setup*

	Package	Version
1	uRProgramming	1.1.1
2	tcltk	3.2.3
3	readr	1.1.1
4	log4r	0.2
5	odfWeave	0.8.4
6	XML	3.98-1.9
7	lattice	0.20-33
8	chron	2.3-51
9	RMySQL	0.10.13
10	DBI	0.7
11	stats	3.2.3
12	graphics	3.2.3
13	grDevices	3.2.3
14	utils	3.2.3
15	datasets	3.2.3

	<b>Package</b>	<b>Version</b>
16	methods	3.2.3

	Package	Version
17	base	3.2.3

```
> environment("ls")
```

```
usage of ls()
funcNameusage of ls(all=TRUE)
funcName
> cat("Current Working Dir \n", getwd())
Current Working Dir
```

/home/zekai/R\_workspace/odfWeaveTmp



## 2.2 ***Using Version Control System***

Talk about the version control, and why it is so important.

Explain why you decide to use git and demonstrate the git.

Talk about git, demonstrate add, commit checkout the difference in the files.  
Binary files in the git,  
how to branch.

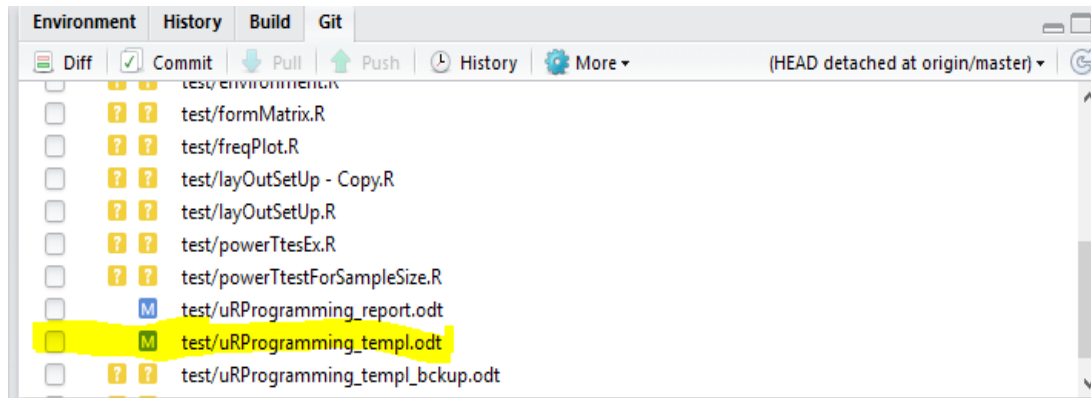




Figure 1 : Git from Rstudio shows which files are modified recently



Figure 3 : Source tree Git shows the created branches



Figure 4 : Git from Rstudio shows the file from the branch

The git commit example from Rstudio, after committed with following message “Modified after adding bar plot example to the document.” we also push our code to the repository.

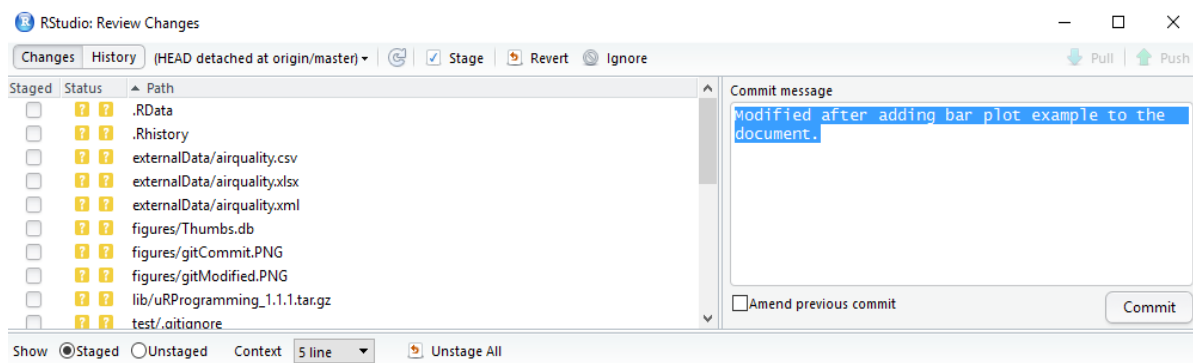




Figure 2 : Git from Rstudio shows how to commit the modified file



Git branching, create either using command line or git GUI like source tree can be used for creating branch and other git functionality.

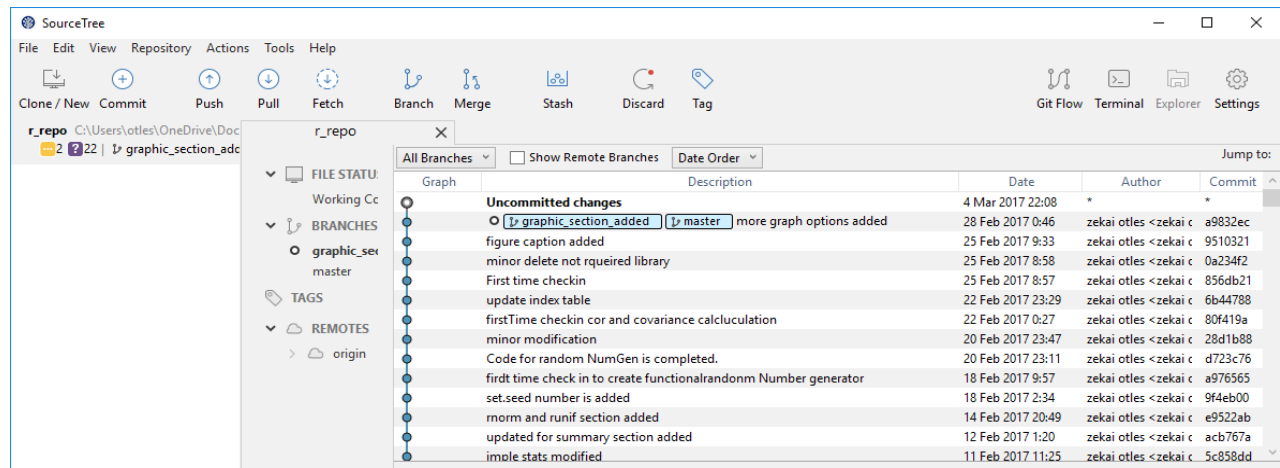




Figure 3 : Source tree Git shows the created branches

When another branches checked out from R studio, it shows what files are checkout from the branch.

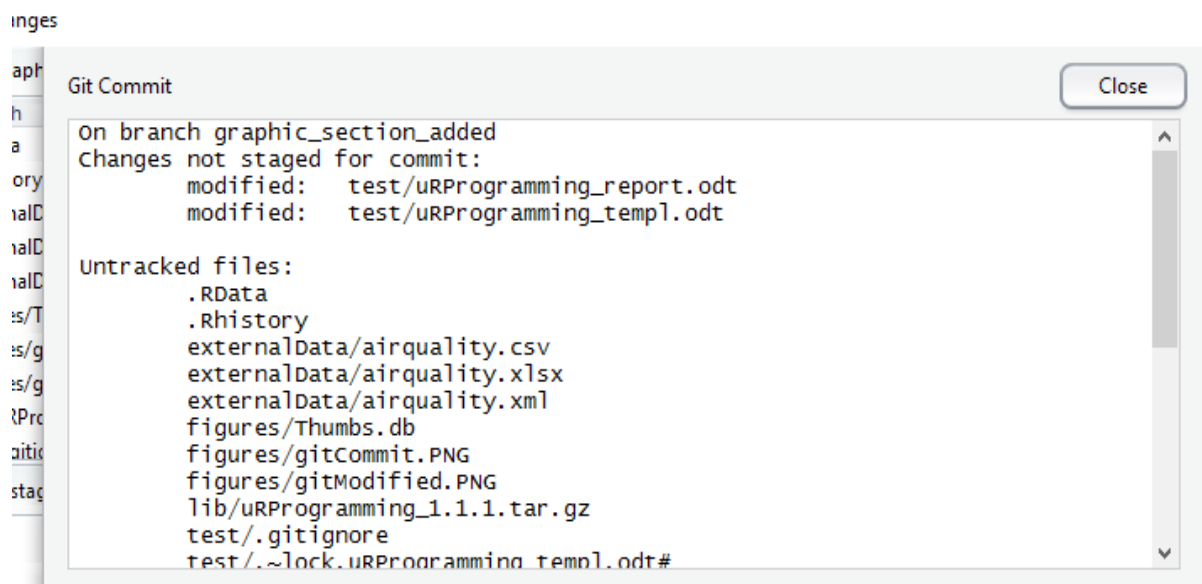




Figure 4 : Git from Rstudio shows the file from the branch

## 2.3 **Using IDE to develop R scripts**

R studio or other tools to develop application,

## 2.4 **Writing functions**

Simple example

## 2.5 **Create R package System**

Build R package Rtools needs to be installed (Refence) for windows and Linux.

Windows insallation and linux instalation. Windows installation is requires downloading Rtools.exe for the R version. Version of R can be found by *version* to the R shell window.

R CMD build uRProgramming

Show them how to build pacjkage from GUI

R CMD BATCH odfWeave.R

## 3 **Simple Data Types and Objects**

R language has following defined data types, Double, Character, Logical and Complex. Numeric variable as a default defined as double,simple text string defined as Character, single character [T and F] is defined as Logical,and complex numbers are defined as Complex.

Integer objects are created with integer, as.integer keywords. Similarly numeric objects are created with numeric and as.numeric keywords. Boolean, Double, Character, Logical, Complex objects are created by their types [boolean, double,character,..] and as.types [boolean.double,..], The variables or vector also can be casted from one type to another type using similar keywords.

```
library(uRProgramming)
dataTypes(24)
dataTypes('ali')
dataTypes(T)
a<-as.integer(24)
dataTypes(a)
standard output from above lines
[1] "double"
[1] "character"
[1] "logical"
[1] "integer"
```

In the following examples show how casting works. Following snippet it, can be copied and saved as testDataTypes.R. This R script should be sourced in the R shell in order to have function testDataTypes(), and run the program. The x vector created as sequence of variables between 1 and 100 increment by 10.011. The x is created as default double variable. The x vector is casted to integer y vector. The later part of the example demonstrates, casting x vector into z character vector.

```
testDataTypes<-function(){
#create logical variable
iTest<-T
is.logical(iTest)
if(iTest == is.logical(iTest)) cat('Logical ')
#create simple double vector
x<-seq(1,100,by=10.011)
is.double(x)
if(iTest == is.double(x)) cat('x is double Vector ')
#cast double vector to integer
y<-as.integer(x)
is.integer(y)
if(iTest == is.integer(y)) cat('y is integer Vector ')
#cast to character vector
z<-as.character(x)
is.character(z)
if(iTest == is.character(z)) cat('z is character Vector ')
}
```

Source this above program, and run the program (testDataTypes())

```
TRUE
Logical
1 11.011 21.022 31.033 41.044 51.055 61.066 71.077 81.088 91.099
x is double Vector
1 11 21 31 41 51 61 71 81 91
y is integer Vector
1 11.011 21.022 31.033 41.044 51.055 61.066 71.077 81.088 91.099
z is character Vector
```

## 4 Control Structures

Control structures are used to change a flow or follow a logic. The control structures are easily could make any program more followable for maintenance and understanding, they could also turn the most unfollowable code. Control structures are very straight forward when used in a modular way in a less number

of statements or calling other reusable functions.

In the following sections, the usage of *if* block will be explained with examples. The samples should help reader to solidify the understanding of the control structures.

## 4.1 If statements

Let's use simple vector which contains only numbers, check if numbers are equal to specific value assign *iFound* as *True*.

*If* ( Statements) { Assignment}

```
x<-c(0,3,6,9,12,15,18,21,24)
if(x[which(x==6)]==6){iFound<-TRUE}
print(iFound)
[1] TRUE
```

In the above example, which finds the index number for the value. The usage of *which* in vector is pretty powerful as shown in the example. The *which* statement is pretty handy determining the index of vector or array. When the vector with right index with exact value found, *iFound* assigned to be *true*.

Logical operators in R are

### Operator Description

==	Equal to
!=	Not equal to
<	Less than
<=	Less or equal to
>	Greater than
>=	Greater or equal to
&	And
	Or

Following code snippet is simple example of how to use logical operators. The first example is the exact *equal* to operator. The second example demonstrates *less than or equal* to operator and assigning the value based on the result. The third example shows *greater than* comparison. The last example is combination *greater than, and, less than or equal* to.

```
Ex. 1
if(demDat[indx,1] ==1 ) {demDat[indx,1]<-'M'}
```

```
Ex.2
if(demDat[indx,2] <=25 )demDat[indx,2]<-'Y'
```

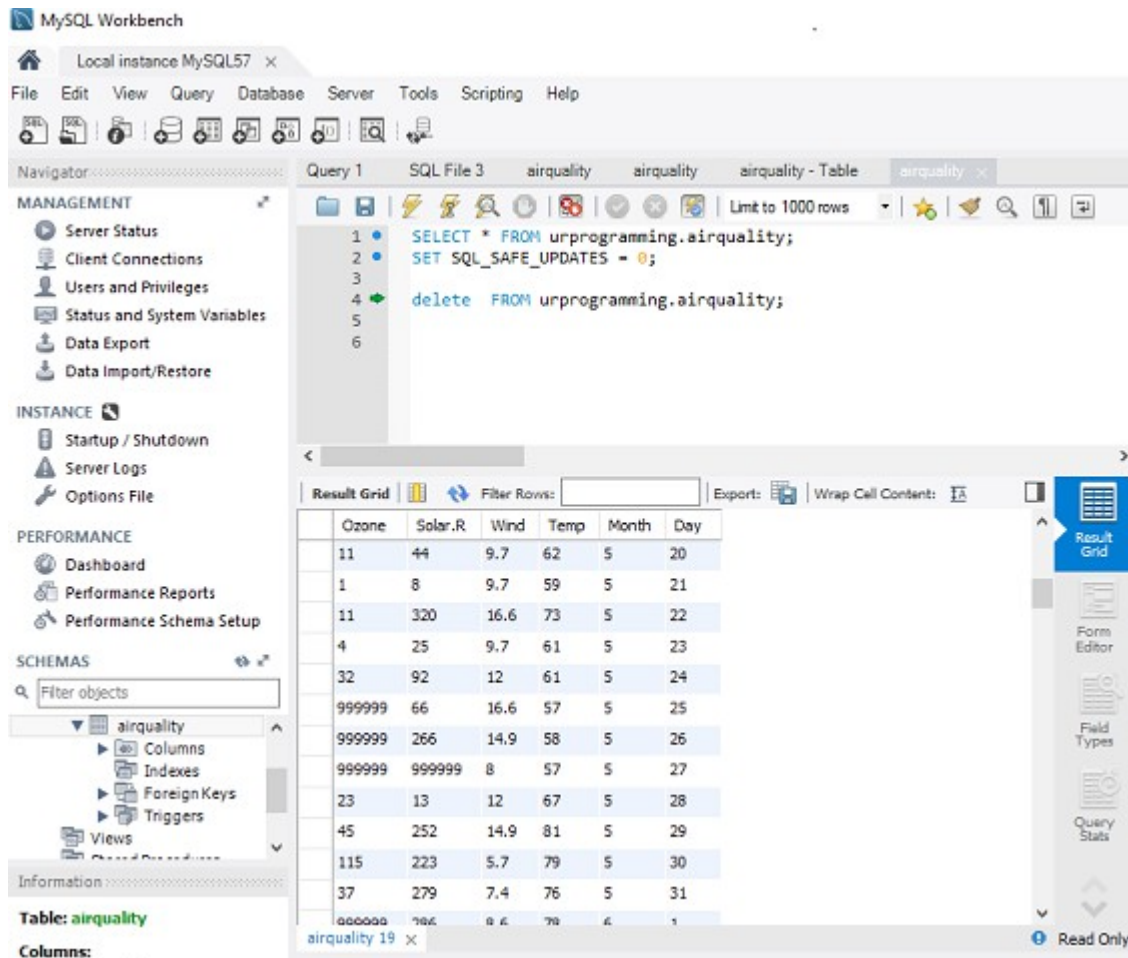
```
Ex.3
if(demDat[indx,2] > 50 ) {demDat[indx,2]<-'O'}
```

```
Ex.4
if(demDat[indx,2] > 25 & demDat[indx,2] <=50 )demDat[indx,2]<-'M'
```

## 4.2 If then Else statements

If then else block is used when comparison expected to yield result either true or false. It gives the programmer, the power to control logical flow. In R programming, if then else blocks are in the following format

*If ( Statements) { Assignment}*



*else (statements) { Assignment}*

## 4.3 Switch statements

One can use switch statement to modular conditional statements. The switch statement evaluates according to the first argument.

*switch(Expression,List)*

The first argument determines how the results will be evaluated. The first line





Figure 5 : MySQL screen shot showing the table and queries

shows a number in the first argument will select the corresponding list element, in the second line of code snippet shows the the corresponding character strings of list element.

```
> switch(3, 1.5, 2, 300, 4)
[1] 300
> switch("iki", bir = "bir", iki = "Two", uc = "uc")
[1] "Two"
```

Following function demonstrates how simple if then else block could be replaced by modular switch statements, this type of example can be extended to more modular functionality.

```
#Simple example for data types casting using swith statement
usingSwitch<-function(x,dataTypes){

  switch (dataTypes,
    integer = {x<-as.integer(x)},
    double = {x<-as.double(x)},
    character = {x<-as.character(x)}

  )

}
> x <- seq(1, 100, by = 10.011)
> x
[1] 1.000 11.011 21.022 31.033 41.044 51.055 61.066 71.077 81.088 91.099
> xC <- usingSwitch(x, "character")
> xC
[1] "1" "11.011" "21.022" "31.033" "41.044" "51.055" "61.066" "71.077"
"81.088" "91.099"
> xI <- usingSwitch(x, "integer")
> xI
[1] 1 11 21 31 41 51 61 71 81 91
```

## 5 Loop Structures

The loop structures are needed to access elements of vectors, matrix, list and data frame. The access of the data is important to reach or manipulate the elements. The *for* and *while* loop are used to iterate elements.

### 5.1 For loop

The for loop is used for sequentially access the data elements. Each element of data structures can be reached, used or manipulated within the *for* loop. The *for* loop is in the following format.

```
for (values in sequence) {  
    statement  
}
```

The following code snippets how *for* loop is used. In the following example, 100 number with specified seed assign to vector. This vector has only integers for demonstration purpose.

```
set.seed(312)  
#create 100 random number between 0 and 1000  
xInt<-floor(runif(100,0,1000))  
for(indx in 0:length(xInt) ){  
  if(indx %%10 ==0){print(xInt[indx])}  
}  
numeric(0)  
[1] 450  
[1] 261  
[1] 405  
[1] 295  
[1] 763  
[1] 356  
[1] 833  
[1] 329  
[1] 419  
[1] 948
```

### 5.2 While loop

The access of certain elements could be controlled by using while loop, the expression determines executing statements or not.

```
while(expression){  
    statement  
}
```

In the following example, the elements between two indices are assigned as either *even* or *odd*. Execution will stop when expression is false, otherwise statements within the block is executed.

```
set.seed(312)
#create 100 random number between 0 and 1000
xInt<-floor(runif(100,0,1000))
index<-80
while(index<=length(xInt)){
  xInt[index]<-ifelse(as.integer(xInt[index]) %%2==0, "Even", "Odd")
  if(index %% 4 == 0 ){print(xInt[index])}
  index<-index+1
}
[1] "Odd"
[1] "Even"
[1] "Odd"
[1] "Odd"
[1] "Odd"
[1] "Even"
```

## 6 Data Structures

In my experience as software engineer and data analyst, introduction to database and utilizing with the data structures would be valuable experience in terms of programming, Specifically storing and retrieving data from database is a great experience for the structured programming. Thus, we introduce the database in this section. The MySQL was chosen because of of simplicity to installation and free to use. The community version is freely available to use, it can be downloaded from oracle. The RMySQL package allows us utilize some of database functionality within R scripts. Available datasets from basic R installation will be loaded to the urprogramming schema into MySql database. Later those dataset tables will be used to demonstrate to basic data structures like vector, list and matrix.

Insert air quality data from R datasets into airquality table into the schema. Later on selected data from mysql table will be converted into vector and list, etc.

```
#insert data into "airquality" table
insertData<-function(){
library("RMySQL")
#if(con)dbDisconnect(con)

numR<-dim(airquality)[1]
#con <- dbConnect(RMySQL::MySQL(), dbname =
"urprogramming",password="urprogramming")

con <- dbConnect(RMySQL::MySQL(), user = 'root', password = 'erzurum', host =
'127.0.0.1' , dbname = 'urprogramming')
results<-dbSendQuery(con, "SELECT * FROM airquality")
dataFrm<-dbFetch(results)

#dbFetch(results)
print(dim(dataFrm)[1])
#print(dataFrm[1:5,])
iSkip<-FALSE
if(dim(dataFrm)[[1]] > 1){iSkip<-FALSE
}else {
  iSkip<-TRUE
}

if(iSkip){
  for (iRow in 1:numR) {

# print(iRow)
data<-airquality[iRow,]

data[is.na(data)]<-9999999
statement<-paste("INSERT INTO ", "airquality")

# print(statement)
insertQuery <- paste(paste(statement,"VALUES("), paste(data, collapse = ", "),
")")
#
print(insertQuery)
```

```

dbGetQuery(con,insertQuery)
}
}#end of iSkip
dbDisconnect(con)

}

```

In order to load airquality data to the table in MySql database, connection to database will open with following statement.

```

#con <- dbConnect(RMySQL::MySQL(), dbname =
"urprogramming",password="urprogramming")

```

```

con <- dbConnect(RMySQL::MySQL(), user = 'root', password = 'erzurum', host =
'127.0.0.1' , dbname = 'urprogramming')

```

The dbConnect statement in RMySQL library requires name of database, schema, username and password. Once the connection is established, the connection object is used to populate the rows in table using the statements starting with "INSERT INTO airquality VALUES(,,,,)".

```

> insertData()
[1] 153
[1] TRUE
> results <- dbSendQuery(con, "SELECT * FROM airquality")
> dataFrm <- dbFetch(results, )
> print(dim(dataFrm))
[1] 153 6
> print(dataFrm[1:5, ])
  Ozone Solar.R Wind Temp Month Day
1    41    190  7.4  67    5    1
2    36    118  8.0  72    5    2
3    12    149 12.6  74    5    3
4    18    313 11.5  62    5    4
5 999999 999999 14.3  56    5    5

```

The following statement will select each row and form a result set.

```

results<-dbSendQuery(con, "SELECT * FROM airquality")

```

The result set will be loaded to dataframe using *dbFetch(results,)* function from the RMySQL library. The *insertData* function is configured so the records will only inserted when there is no data in the table. Here, we introduced the basic functionality of database in R. Similar to *insert and select* , *update* , *truncate* could be used to manipulate data in tables. There are many options for SQL

statements to create complex joins or selecting the subsets of the data . This kind of SQL statements can be found in MySQL documentations, that's why we did not want to go beyond the basic of SQL statements.



Figure 5 : MySQL screen shot showing the table and queries



Figure 5 : MySQL screen shot showing the table and queries



# MySQL Workbench

Local instance MySQL57 x

File Edit View Query Database Server Tools Scripting Help



Navigator

## MANAGEMENT

- Server Status
- Client Connections
- Users and Privileges
- Status and System Variables
- Data Export
- Data Import/Restore

## INSTANCE

- Startup / Shutdown
- Server Logs
- Options File

## PERFORMANCE

- Dashboard
- Performance Reports
- Performance Schema Setup

## SCHEMAS

Filter objects

- airquality
  - Columns
  - Indexes
  - Foreign Keys
  - Triggers
  - Views
  - Stored Procedures

Information

Table: airquality

Columns:

Query 1

SQL File 3

airquality

airquality

airquality - Table

airquality



```
1 SELECT * FROM unprogramming.airquality;
2 SET SQL_SAFE_UPDATES = 0;
3
4 delete FROM unprogramming.airquality;
5
6
```

Result Grid Filter Rows: Export: Wrap Cell Content:

	Ozone	Solar.R	Wind	Temp	Month	Day
11	44	9.7	62	5	20	
1	8	9.7	59	5	21	
11	320	16.6	73	5	22	
4	25	9.7	61	5	23	
32	92	12	61	5	24	
999999	66	16.6	57	5	25	
999999	266	14.9	58	5	26	
999999	999999	8	57	5	27	
23	13	12	67	5	28	
45	252	14.9	81	5	29	
115	223	5.7	79	5	30	
37	279	7.4	76	5	31	
999999	796	8.6	79	6	1	

airquality 19 x

Read Only

Result Grid

Form Editor

Field Types

Query Stats



Figure 5 : MySQL screen shot showing the table and queries

Similar data stored in in data bases, one could easily have data stored so called as flat data asci files. Those are created from either either simple editor or spread sheet programs. R has very easy interfaces to read flat files or spreadsheet data. The flat file stored in `"/home/zekai/R_workspace/r_repo/externalData/airquality.csv"` can be read with following statement

```
read.csv(file="/home/zekai/R_workspace/r_repo/externalData/airquality.csv",header=TRUE,sep=","  
)
```

If data is in spreadsheet, it can be saved as flat file from spreadsheet then read data from flat file using similar staement. The data file could be read using `read.csv` funtion from R util pacakage. There are other packages avialable for reading spreadsheet data to R.

## 6.1 Data Frame

Data frame is one of the available data structures. The data frame is similar to the spreadsheet data, one can easily see the structure of the data. Once the data saved in data frame structure, the structure of data, dimensions and contents of data is readily available in R. The structure of data is determined from **`str( name of data frame object)`** as shown in the the following example

**`str(dataFrm).`**

The statement gives the structure of data, name of data variables and also type of variable stored. As seen in the example, air quality data store Ozone, Solar.R, Temp, Month, Day variables as **`int`** and wind as **`num`**. These variables can be changed using casting as we shown earlier in Simple Data Type section. We can figure out the data stored in a data frame using existing function **`dim`** in R base functions, **`colnames`** (again from R base functions) refers the variable names stored in data structures. The **`View`** function R util functions can display the data frame contents so we can have some idea about the data. The following section

**`View(dataFrm[1:4,])`**

is displaying the data from the first through fourth row in dataFrm.

```
results<-dbSendQuery(con, "SELECT * FROM airquality")  
dataFrm<-dbFetch(results)
```

the results from select assigned to dataFrm. The dataFrm has following structure  
`str(dataFrm)`  
'data.frame': 153 obs. of 6 variables:

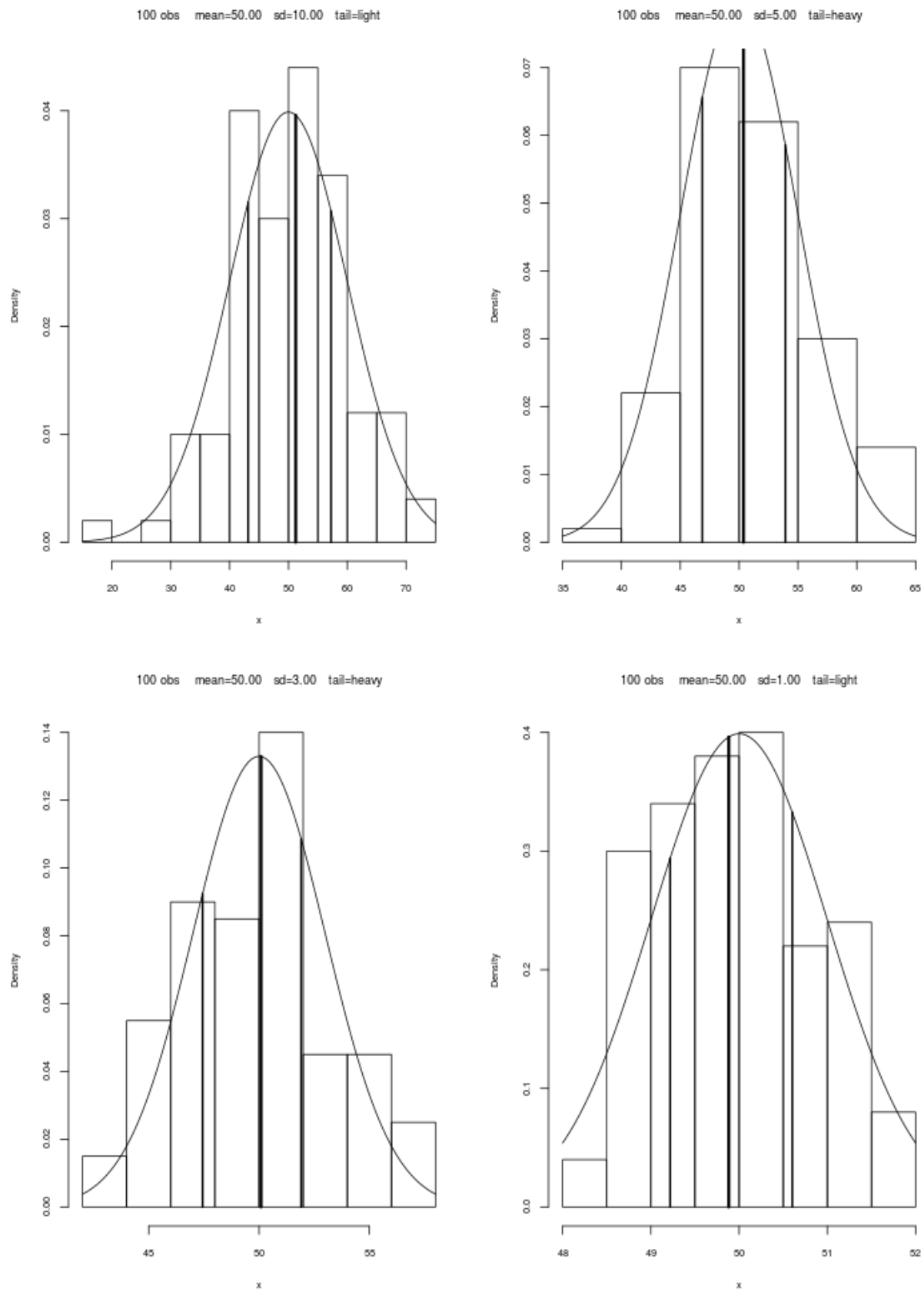


Figure 6 : Example Histogram

```

$ Ozone   : int   41 36 12 18 999999 28 23 19 8 999999 ...
$ Solar.R: int   190 118 149 313 999999 999999 299 99 19 194 ...
$ Wind    : num   7.4 8 12.6 11.5 14.3 14.9 8.6 13.8 20.1 8.6 ...
$ Temp    : int   67 72 74 62 56 66 65 59 61 69 ...
$ Month   : int    5 5 5 5 5 5 5 5 5 5 ...
$ Day     : int    1 2 3 4 5 6 7 8 9 10 ...
dim(dataFrm)
[1] 153    6
colnames(dataFrm)
[1] "Ozone" "Solar.R" "Wind" "Temp" "Month" "Day"
View(dataFrm[1:4,])
  Ozone Solar.R Wind Temp Month Day
1    41    190  7.4   67     5    1
2    36    118  8.0   72     5    2
3    12    149 12.6   74     5    3
4    18    313 11.5   62     5    4

```

## 6.2 Vector

Vector is used for atomic data types (integer, numeric, complex, and character) and list storage. It is a powerful way to store data and its usage in the computations extremely simplifies the programming. The examples will demonstrate the power of vector calculations. Temperature from Fahrenheit is easily converted to Celsius with following statement.

***tempAsCels<-5\*(dataFrm\$Temp-32)/9.0***

```

> tempAsCels <- 5 * (dataFrm$Temp - 32)/9
> cat("is.vector(tempAsCels)")
is.vector(tempAsCels)
> print(is.vector(tempAsCels))
[1] TRUE
> cat("View(tempAsCels(1:5))")
View(tempAsCels(1:5))
> print(tempAsCels[1:5])
[1] 19.44444 22.22222 23.33333 16.66667 13.33333

```

It is also easy to dissect and examine data stored in vector format. The following simple snippet will demonstrate how to retrieve the month data from the dataFrm object. We have determined the months which days are 80 degree Fahrenheit. It also shows number of 80 degree days for our data.

```

dataFrm$Month[dataFrm$Temp==80]
length(dataFrm$Month[dataFrm$Temp==80])
[1] 6 6 7 8 9
[1] 5

```

The **which**, function is available from the R base package, is used to determine the indices of the vector when the condition meets up the criteria. The following snippets show which element of the temperature vector was exactly 80 degrees Fahrenheit.

```

which(dataFrm$Temp==80)

```

[1] 45 59 76 106 130

## 6.3 Matrix

The matrix is available in R to store two dimensional **array** in a matrix format. The discussion of matrix without **array** will not be complete to understand data structures in **R** programming. The array can store vector data in one, two or dimensional format. The vector data is stored without array bound information, but **array** and **matrix** needs complete information about dimensions.

The **array** needs the information about data types and dimensions of the array. The **array** stores one to many dimensional data. In the following example, `row_names` stored in a one dimensional **array**.

```
row_Names<-array("",dim=length(unique(dataFrm$Month)))
```

Structure of **array** can be found using **str** from **util** package in R. The **str** function will give data types and dimension of the **array**. The following demonstrates how to use **str** function with **array**.

```
strr(row_Names)
```

Another useful function to check whether data stored in array or not. If the data stored in an array, `is.array(row_Names)` will return *TRUE*.

The **matrix** is a special case of **array** which is a two dimensional data structure. The **matrix** need data types, row and column dimensions. The `tempAsMatrix` is defined as **matrix** in the following example.

```
tempAsMatrix<-matrix("",nrow=length(unique(dataFrm$Month)),ncol=30)
```

```
#form a Matrix from dataFrm data
#the purpose of this function to ceate Temp matrix
formTempMatrix<-function(dataFrm)
{
  monthAsNum<-unique(dataFrm$Month)
  row_Names<-array("",dim=length(unique(dataFrm$Month)))
  tempAsMatrix<-matrix("",nrow=length(unique(dataFrm$Month)),ncol=30)

  for(mIndx in 1:length(monthAsNum)){
    row_Names[mIndx]<-
switch(as.character(monthAsNum[mIndx]), "5"="May", "6"="June", "7"="July", "8"="Aug", "
9"="Sep")
  }
  rownames(tempAsMatrix)<-row_Names

  length(unique(dataFrm$Month))
  rowIndex<-unique(dataFrm$Month)
  for (index in 1:length(rowIndex)){
```

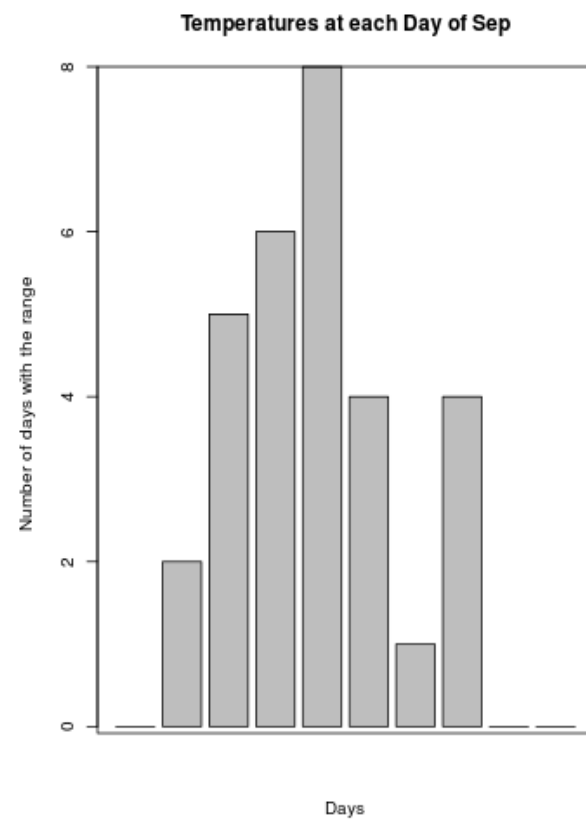
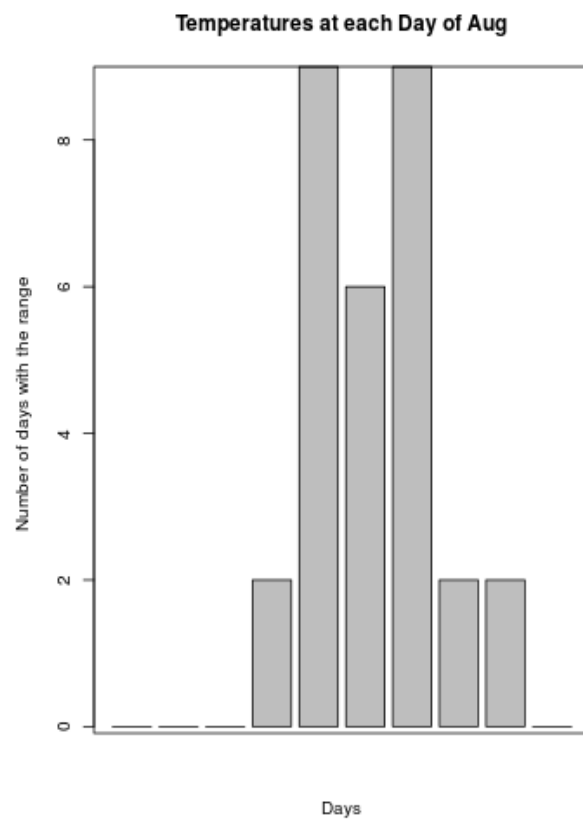
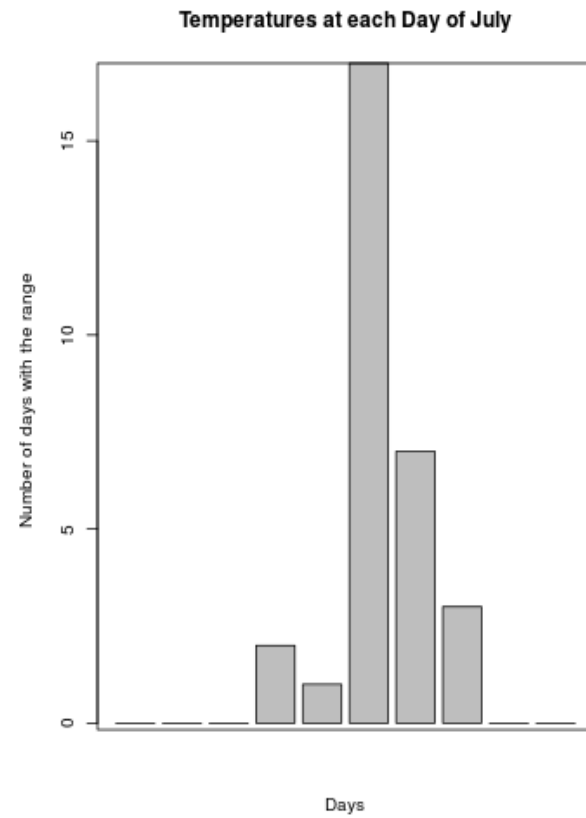
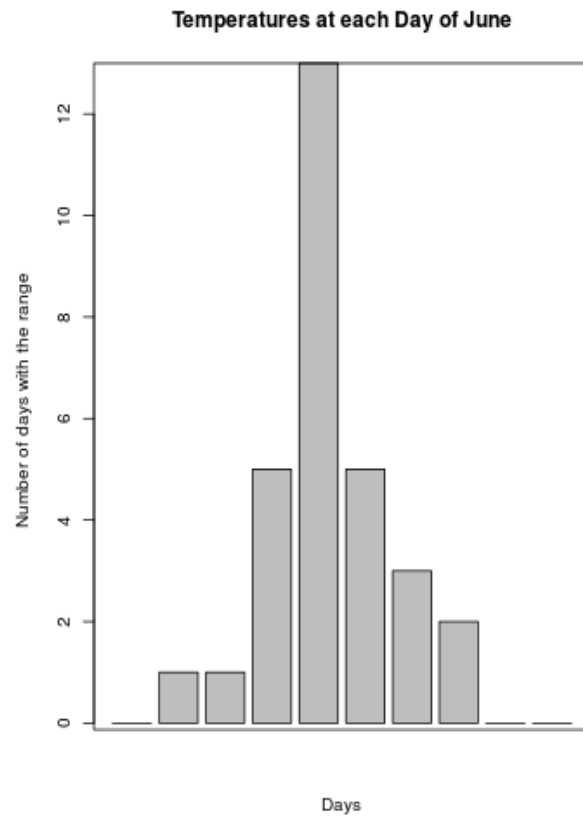


Figure 7 : Example Bar Plot

```

    x<-dataFrm$Temp[dataFrm$Month==rowIndex[index]]
    tempAsMatrix[index,]<-x[1:30]
  }

  return(tempAsMatrix)
}

```

The **str** function for matrix type will return data types, and bounds of the each dimensions. The base package has a function to determine whether data structure is matrix or not. In the following example,

```
is.matrix(tempAsMatrix)
```

the return value will be TRUE.

## 6.4 List

The list is very powerful data structure, heterogeneous data can be stored in a list. The list can contain list, data frame, array and matrix in a structure. The *dataFrm*, *row\_Names* and *tempAsMatrix* data structures are all stored in a **list** called *aList*. The data structure can be retrieved using the *list\$name* structure, such as *aList\$dataFrm*.

```

#demonstate the list
useList<-function(dataFrm,row_Names,tempAsMatrix){
  aList<-list(dataFrm=dataFrm,rowNames=row_Names,matrix=tempAsMatrix)
  str(aList)
  return(aList)
}

str(aList)
List of 3
 $ dataFrm :'data.frame': 153 obs. of  6 variables:
  ..$ Ozone   : int [1:153] 41 36 12 18 999999 28 23 19 8 999999 ...
  ..$ Solar.R: int [1:153] 190 118 149 313 999999 999999 299 99 19 194 ...
  ..$ Wind    : num [1:153] 7.4 8 12.6 11.5 14.3 14.9 8.6 13.8 20.1 8.6 ...
  ..$ Temp    : int [1:153] 67 72 74 62 56 66 65 59 61 69 ...
  ..$ Month   : int [1:153] 5 5 5 5 5 5 5 5 5 5 ...
  ..$ Day     : int [1:153] 1 2 3 4 5 6 7 8 9 10 ...
 $ rowNames: chr [1:5(1d)] "" "" "" "" ...
 $ matrix   : chr [1:5, 1:30] "" "" "" "" ...

```

As seen above, the list data structure is very convenient way to store multiple data structures into a data structure.

## 7 Simple Statistical Analysis

This section is simple introduction to statistical analysis. There will be brief explanation about creating normally distributed variables used statistical analysis, creating simple summary mean, standard deviation. The simple statistical functions are available stats, base, time functional packages. The functional examples will show how to



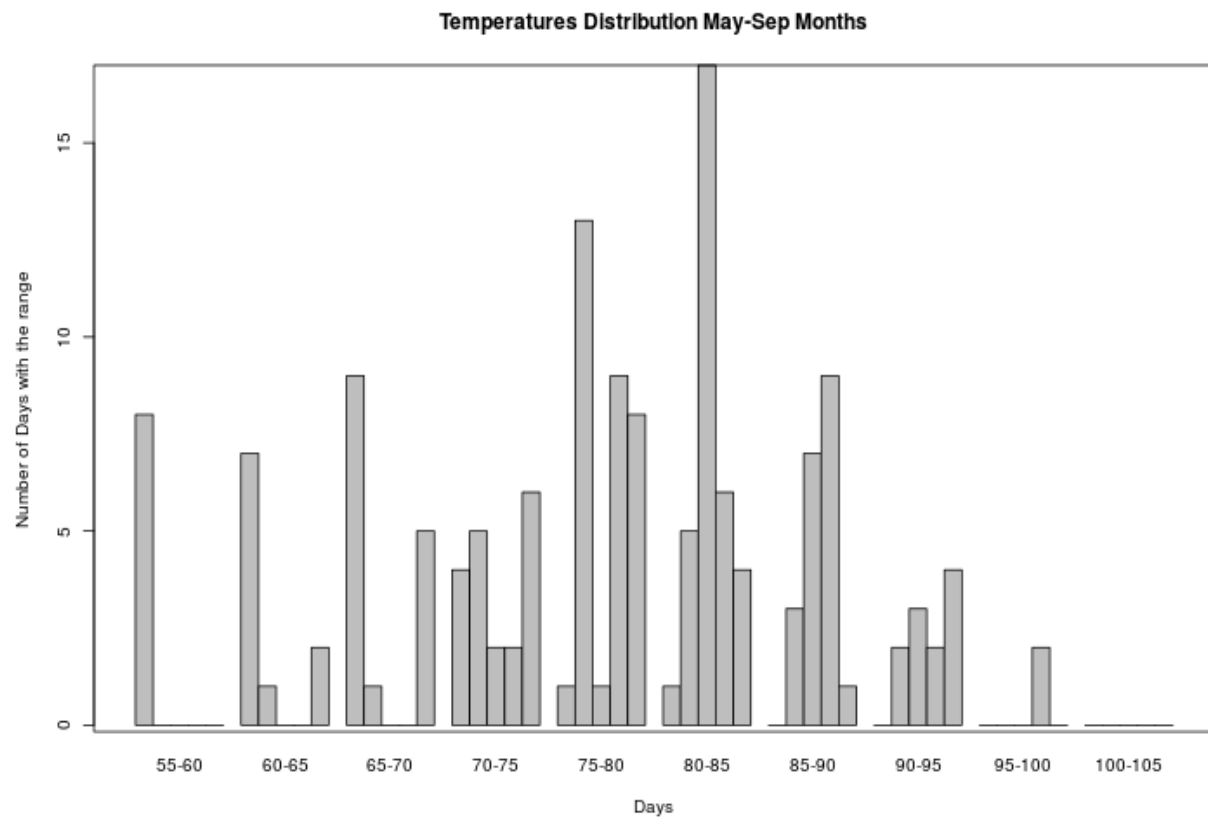


Figure 8 : Example Bar Plot with combined data

use those simple statistics using already defined data structures. Introduction how to use simple statistics against their data will give foundation to extend the readers ability to do further complex statistical analysis. This will also help utilizing the other packages in available by R community.

```
#demonstrate simple distribution and some statistics
simpleStats<-function(n,mean,sd){
x<-rnorm(n,mean=mean,sd)
#mean of variable
mean=mean(x)
ave=ave(x)
max=max(x)
min=min(x)
median=median(x)
sd=sd(x)
sum=sum(x)
summary=summary(x)
return(list(mean=mean,ave=ave,max=max,min=min,median=mean,sd=sd,summary=summary,su
m=sum))
}
```

In the example, normally distributed random variable created using **rnorm** function from **stats** package. Using some of the available functions such as **mean**, **summary** and **sd** from **base** and **stats** packages. The **mean** from **base** package gives the arithmetic mean of the values. The **min** and **max** returns minimum and maximum values for the input values. The **sd** is standard deviation of the input values. The summary function from base package returns **min** and **max** values , the first and third quartiles, **median** and **mean** of input values.

```
xStats<-simpleStats(100,273.16,15)
str(xStats)
List of 8
 $ mean      : num 275
 $ ave       : num [1:100] 275 275 275 275 275 275 ...
 $ max       : num 318
 $ min       : num 241
 $ median    : num 275
 $ sd        : num 14.9
 $ summary:Classes 'summaryDefault', 'table'  Named num [1:6] 241 266 273 275
283 ...
 .. ..- attr(*, "names")= chr [1:6] "Min." "1st Qu." "Median" "Mean" ...
 $ sum       : num 27463
```

Normally distributed random numbers is generated by using **rnorm** function from stats package. The **runif** generates uniformly distributed random numbers. The random number generators ran four times in the example, and selected the fifth element of the storage vectors (**xTempNorm** and **xTempUni**). Since the script is not using fixed seed number for the random number generators, the

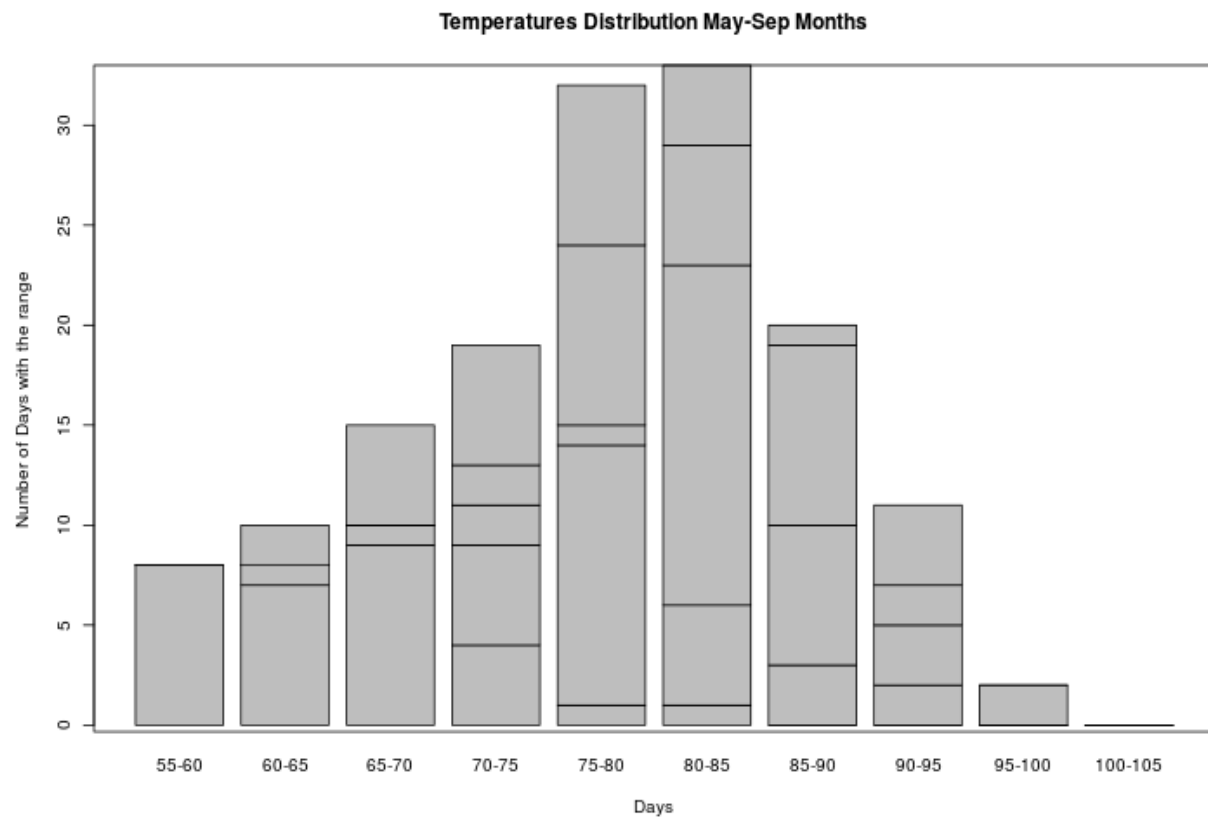


Figure 9 : Example Bar Plot with combined data

storage vectors will consist of different elements as we shown here with the fifth elements of the vectors.

```
#Create randomly number generated vectors
#simple statistics
randomNumGen<-function(seed=FALSE){
print("Normally distributed Random Number")
  for(indx in 0:3){
    print(paste("Iteration ",indx,sep= ""))
    if(seed) set.seed(6543)
    xTempNorm<-rnorm(100,273.16,15)

    print(paste("xTempNorm[5]=",xTempNorm[5]))
    cat('summary(xTempNorm)\n')
    print(summary(xTempNorm))

  }

  print("Uniformly distributed Random Number")
  for(indx in 0:3){
    if(seed)set.seed(6543)
    xTempUni<-runif(100,240,320)
    print(paste("Iteration ",indx,sep= ""))
    print(paste("xTempUni[5]=",xTempUni[5]))
    cat('summary(xTempUni)\n')
    #

    print(summary(xTempUni))

  }

return(list(xTempNorm=xTempNorm,xTempUni=xTempUni))
}
[1] "Normally distributed Random Number"
[1] "Iteration 0"
[1] "xTempNorm[5]= 268.63228295104"
summary(xTempNorm)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 237.9  265.1   274.8   274.4   282.5   302.0
[1] "Iteration 1"
[1] "xTempNorm[5]= 273.256935108386"
summary(xTempNorm)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 234.0  264.8   272.5   272.4   282.3   307.5
[1] "Iteration 2"
[1] "xTempNorm[5]= 248.233597307623"
summary(xTempNorm)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.4  264.2   274.5   274.8   285.5   307.9
[1] "Iteration 3"
[1] "xTempNorm[5]= 278.064499119773"
summary(xTempNorm)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 229.7  263.6   272.6   273.3   283.4   315.9
[1] "Uniformly distributed Random Number"
[1] "Iteration 0"
[1] "xTempUni[5]= 245.648153163493"
```

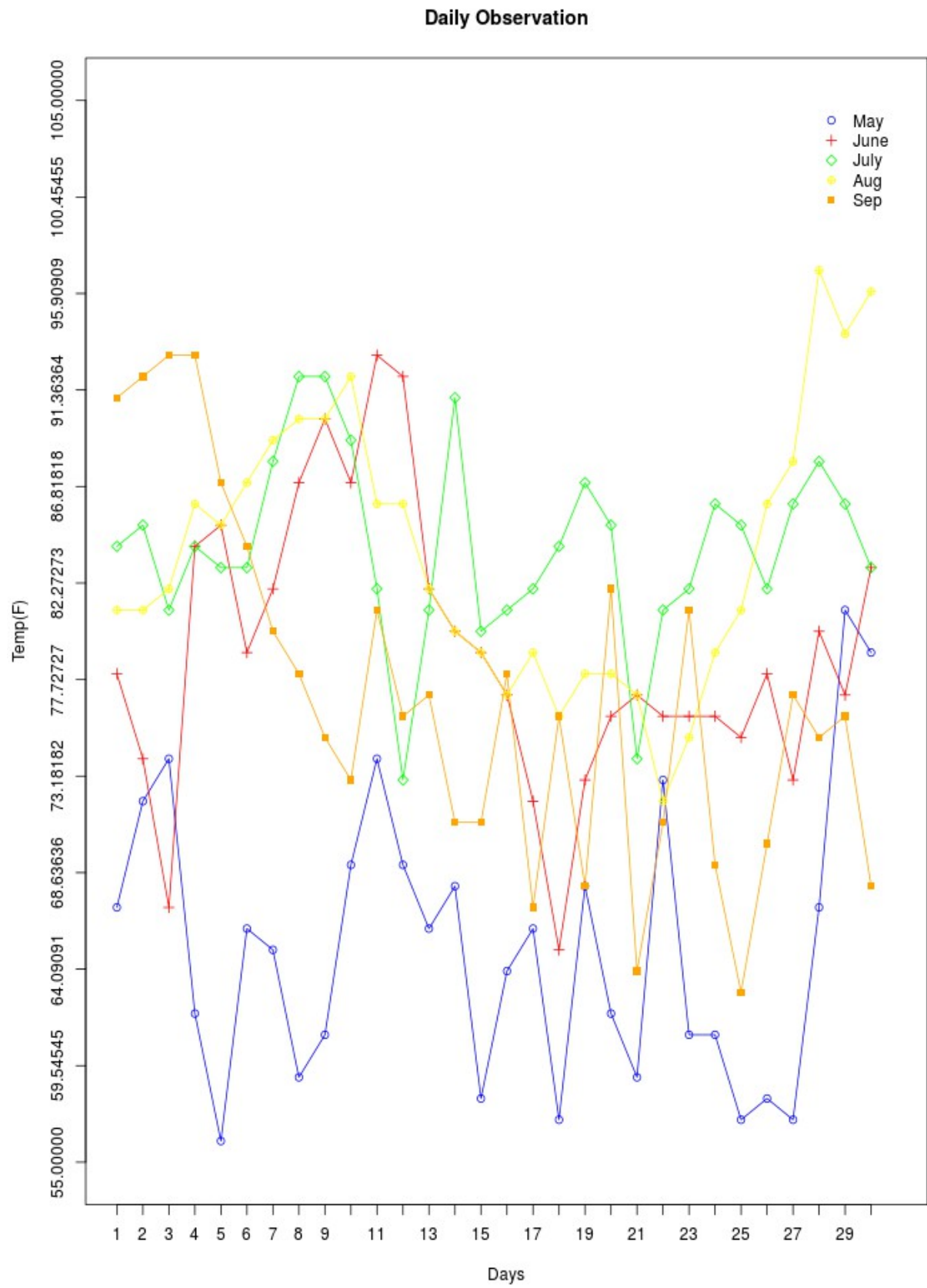


Figure 10 : Observation

```
summary(xTempUni)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.4  256.4  274.5   276.9  298.4   319.5
[1] "Iteration 1"
[1] "xTempUni[5]= 253.614315409213"
summary(xTempUni)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.6  255.2  271.0   276.9  299.8   319.9
[1] "Iteration 2"
[1] "xTempUni[5]= 281.440007463098"
summary(xTempUni)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.3  260.5  279.3   279.9  298.1   319.0
[1] "Iteration 3"
[1] "xTempUni[5]= 265.973723661155"
summary(xTempUni)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.4  256.3  277.3   277.5  300.3   319.9
```

The same example will be repeated for fixed seed numbers for the random number generation. When we set the seed the numbers the produced random number vectors using **rnorm** and **runif** will always contains same elements. The fifth elements for the **xTempNorm** and **xTempUni** vectors is always same for the given seed number 6543. The **summary** of the storage **xTempNorm** and **xTempUni** vector will yield same results because of the same seed number.

```
[1] "Normally distributed Random Number"
[1] "Iteration 0"
[1] "xTempNorm[5]= 288.495445822198"
summary(xTempNorm)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 221.5  259.9  273.3   270.8  281.0   304.0
[1] "Iteration 1"
[1] "xTempNorm[5]= 288.495445822198"
summary(xTempNorm)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 221.5  259.9  273.3   270.8  281.0   304.0
[1] "Iteration 2"
[1] "xTempNorm[5]= 288.495445822198"
summary(xTempNorm)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 221.5  259.9  273.3   270.8  281.0   304.0
[1] "Iteration 3"
[1] "xTempNorm[5]= 288.495445822198"
summary(xTempNorm)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 221.5  259.9  273.3   270.8  281.0   304.0
[1] "Uniformly distributed Random Number"
[1] "Iteration 0"
```

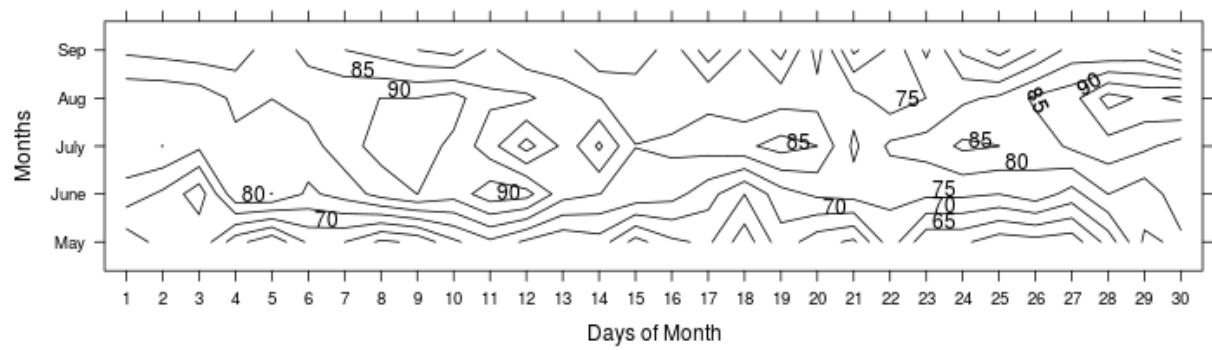


Figure 11 : Daily Temperature Values Plotted as Contours

```
[1] "xTempUni[5]= 272.028888594359"
summary(xTempUni)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.0  262.8  281.6   280.5  297.3   318.4
[1] "Iteration 1"
[1] "xTempUni[5]= 272.028888594359"
summary(xTempUni)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.0  262.8  281.6   280.5  297.3   318.4
[1] "Iteration 2"
[1] "xTempUni[5]= 272.028888594359"
summary(xTempUni)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.0  262.8  281.6   280.5  297.3   318.4
[1] "Iteration 3"
[1] "xTempUni[5]= 272.028888594359"
summary(xTempUni)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 240.0  262.8  281.6   280.5  297.3   318.4
```

The `xRandGenT` list is used to sample two vector datasets. The sample method from base R package used to select 15 random elements from vector datasets. Once the samples assigned the correlations values computed by using different methods available from **stats** package. The covariance between **xTempNorm** and **xTempUni** vector datasets computed using `cov` function from **stats** package. The standard deviations calculated using `sd` function from the **stats** package.

```
corTest<-function(xVals=xVals,yVals=yVals){
x<-sample(xVals,size=15)
y<-sample(yVals,size=15)
#cor(x,y,use="everything",method=c("pearson","kendall","spearman"))
cor.test(x,y,method="pearson")
print(cor.test(x,y,method="pearson"))
sdXsdY=sd(x)*sd(y)
pearsonCor=cov(x,y,method="pearson")/sdXsdY
print(paste("Pearson=",pearsonCor))

cor.test(x,y,method="spearman")
print(cor.test(x,y,method="spearman"))

cor.test(x,y,method="kendall")
print(cor.test(x,y,method="kendall"))

}
corTest(xRandGenT$xTempNorm,xRandGenT$xTempUni)
```

Pearson's product-moment correlation

```
data:  x and y
t = -0.7144, df = 13, p-value = 0.4876
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
```



	May	June	July	Aug	Sep
1	67	78	84	81	91
2	72	74	85	81	92
3	74	67	81	82	93
4	62	84	84	86	93
5	56	85	83	85	87
6	66	79	83	87	84
7	65	82	88	89	80
8	59	87	92	90	78
9	61	90	92	90	75
10	69	87	89	92	73
11	74	93	82	86	81
12	69	92	73	86	76
13	66	82	81	82	77
14	68	80	91	80	71
15	58	79	80	79	71
16	64	77	81	77	78
17	66	72	82	79	67
18	57	65	84	76	76
19	68	73	87	78	68
20	62	76	85	78	82
21	59	77	74	77	64
22	73	76	81	72	71
23	61	76	82	75	81
24	61	76	86	79	69
25	57	75	85	81	63
26	58	78	82	86	70
27	57	73	86	88	77
28	67	80	88	97	75
29	81	77	86	94	76
30	79	83	83	96	68

-0.6426395 0.3530540  
sample estimates:

```

      cor
-0.1943601

[1] "Pearson= -0.194360080385567"

Spearman's rank correlation rho

data:  x and y
S = 766, p-value = 0.1779
alternative hypothesis: true rho is not equal to 0
sample estimates:
      rho
-0.3678571

```

```

Kendall's rank correlation tau

data:  x and y
T = 40, p-value = 0.2395
alternative hypothesis: true tau is not equal to 0
sample estimates:
      tau
-0.2380952

```

## 8 Graphic Outputs

This section is designed to be how to represent data output in a graphical formats.

The visual presentation of data will help audience form an opinion much faster in human's brain. This might be very well represented by the common known phrase “A picture is worth a thousand words” (<http://www.phrases.org.uk/meanings/a-picture-is-worth-a-thousand-words.html>). This is mind, we will show examples of histogram, bar plot, plot, matplot is readily available R **graphics** pckage.

```

#The purpose of this test to see effect of number of Normal distribution
#Normal distribution is back bone statistical theory
#to run this
#testNormal(numSet=100,mean=c(5,15,50),sd=c(10,5,3,1))
#num number of random observations
normalDist<-function(numSet=numSet,meanSet=meanSet,sdSet=sdSet){

  #opar<-par(ask=interactive())
  opar<-par(font.main=1,ps=9)
  layout(matrix(c(1,2,3,4),2,2,byrow=TRUE))
  for(numIndx in 1:length(numSet)){
    num<-numSet[numIndx]
    for(pIndx in 1:length(meanSet)){
      p=meanSet[pIndx]
      for(i in 1:length(sdSet)){

        x<-rnorm(num,mean=p,sd=sdSet[i])

```

```

iqr<-(quantile(x)[4]-quantile(x)[2])
psd<-iqr/1.35

if(sd(x)<psd){

```

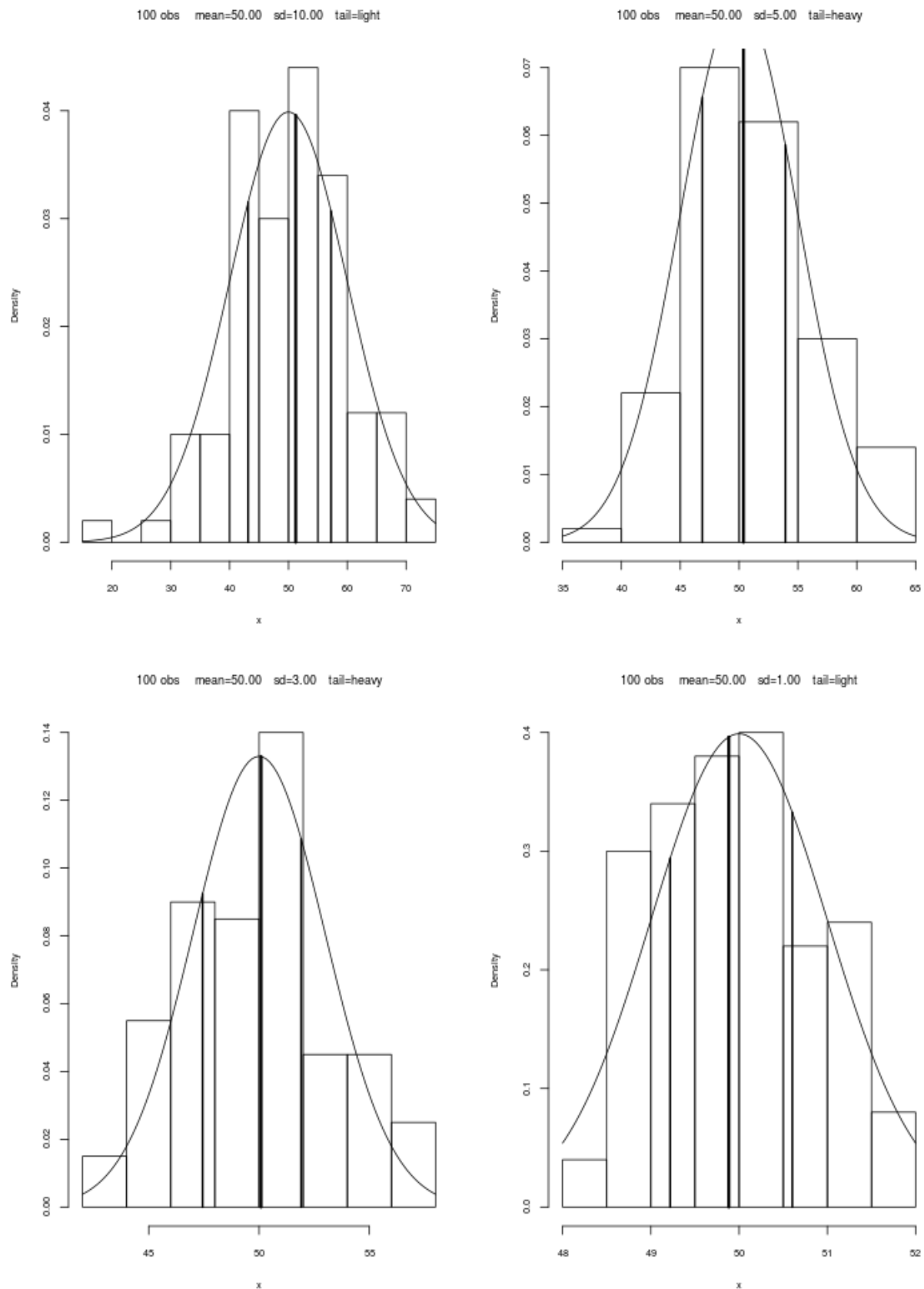


Figure 6 : Example Histogram

```

    tail="light"
  }else{
    tail="heavy"
  }

```

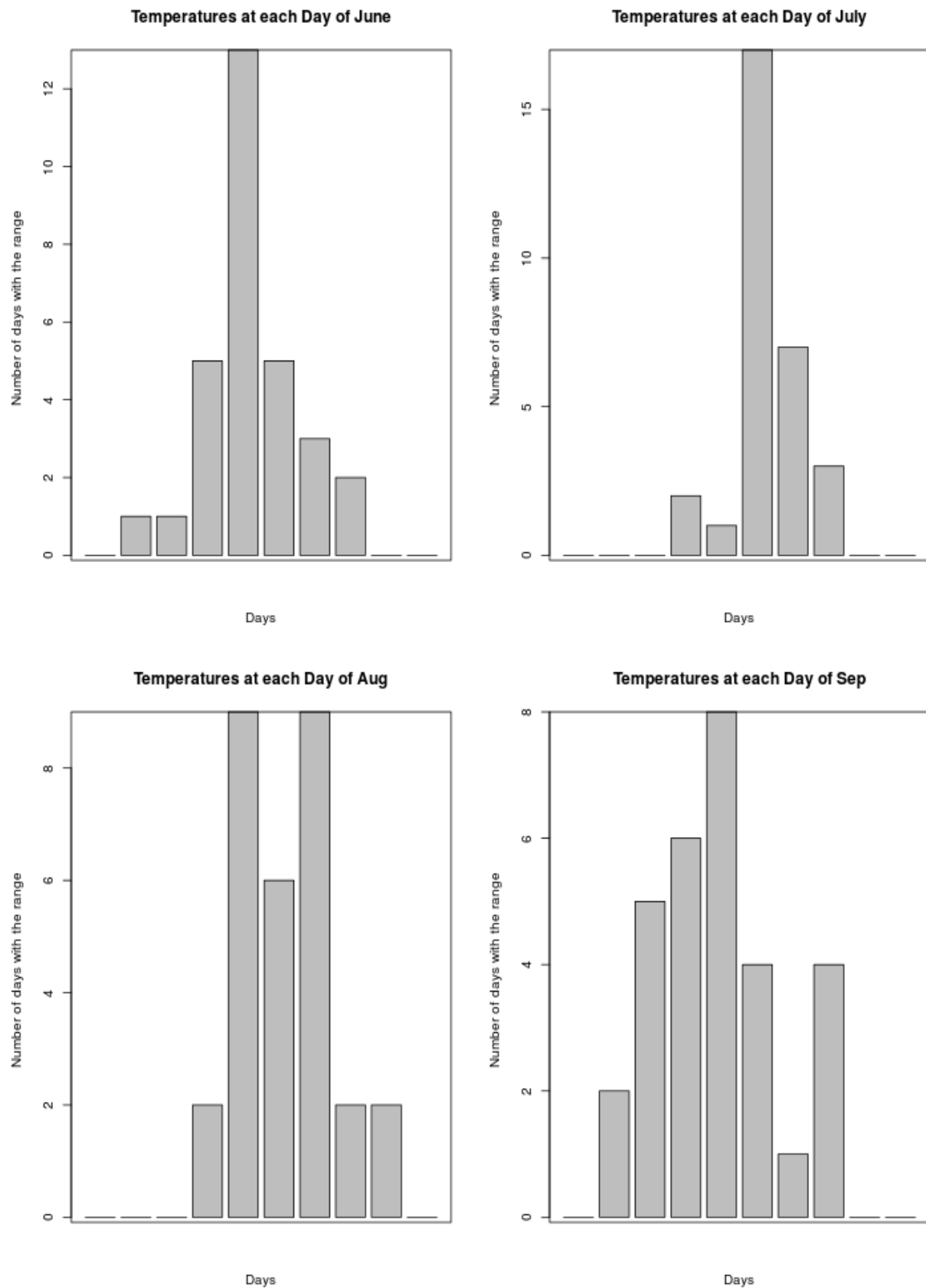


Figure 7 : Example Bar Plot