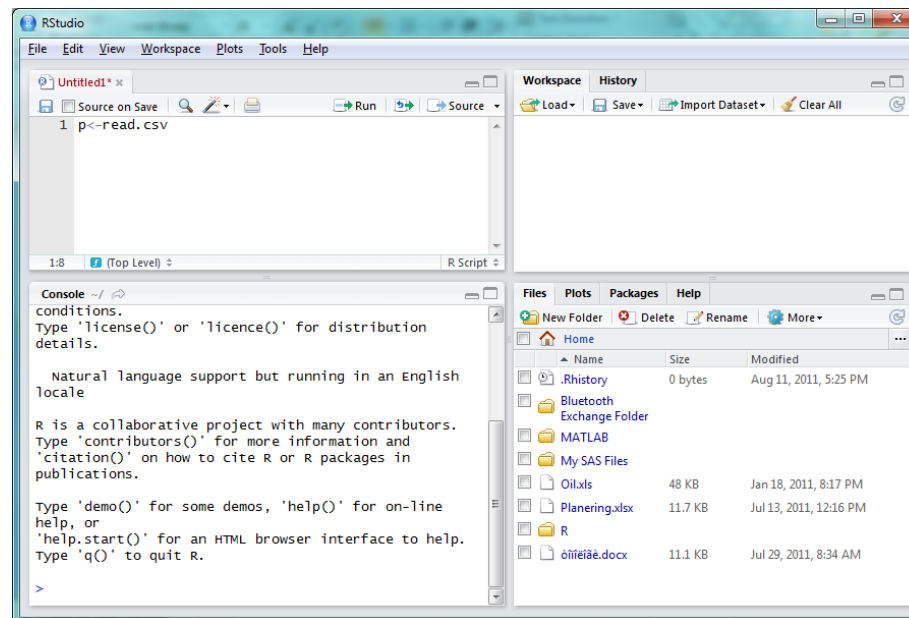


Introduction to R

Lecture 1c

R

- R is a **programming language** of a higher-level
- Constantly increasing amount of packages (new research)
- Free of charge
- Website: <http://www.r-project.org/>
- Code Editor: <http://rstudio.org/>



Software: use RStudio

- Install R: <http://www.r-project.org/>
- Install RStudio: <http://rstudio.org/>

The screenshot shows the RStudio environment with four main panels: Source, Workspace, Console, and Plots. Annotations with arrows point to each panel:

- Program**: Points to the Source editor showing R code for loading data and creating a plot.
- Workspace**: Points to the Environment pane showing the loaded 'diamonds' dataset and the 'p' plot object.
- Execution console**: Points to the Console pane showing the output of the R commands, including summary statistics for the 'diamonds' dataset.
- Plots**: Points to the Plots pane showing a scatter plot titled 'Diamond Pricing' with 'Carat' on the x-axis and 'Price' on the y-axis, colored by clarity.

Source Editor Code:

```
1 library(ggplot2)
2 source("plots/formatPlot.R")
3
4 view(diamonds)
5 summary(diamonds)
6
7 summary(diamonds$price)
8 aveSize <- round(mean(diamonds$carat), 4)
9 clarity <- levels(diamonds$clarity)
10
11 p <- qplot(carat, price,
12            data=diamonds, color=clarity,
13            xlab="carat", ylab="Price",
14            main="Diamond Pricing")
15
```

Console Output:

```
> summary(diamonds$price)
   Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
   326     950    2401    3933    5324   18820
> aveSize <- round(mean(diamonds$carat), 4)
> clarity <- levels(diamonds$clarity)
> p <- qplot(carat, price,
+            data=diamonds, color=clarity,
+            xlab="carat", ylab="Price",
+            main="Diamond Pricing")
> format.plot(p, size=24)
>
```

Workspace Data:

Variable	Value
aveSize	0.7979
clarity	character[8]
p	ggplot[8]

Plots: A scatter plot titled 'Diamond Pricing' showing 'Price' (y-axis, 0 to 15,000) versus 'Carat' (x-axis, 0.0 to 3.5). The points are colored by clarity, with a legend on the right showing categories: I1, SI2, SI1, VS2, VS1, VVS2, VVS1, and IF.

Basics in RStudio

Important to know:

- Create a new file and save it (File menu)
- Running one line or entire code (Edit menu)
- Running one line in console
- Workspace (Observe, Save, Clear)
- Setting current directory (Tools)
- Installing new package (Packages tabs)

Call help

- Specific function
 - `help(function)`
- Help browser
 - `help.start()`
- Search for something in help
 - `help.search("expression")`
- Quick reminder of function arguments:
 - `args(function)`
- Examples of how to use function:
 - `example(function)`
- If some method is not installed on the computer:
 - `RSiteSearch("expression")`

Introduction

- R is case-sensitive (A and a)
- Each command on a new line
- Comment:

```
#R is a very cool language!
```

Initialize/set the variable

Use-> or <- or =

```
a<-3
```

```
a=3
```

```
3->b
```

Vectors

- Create a vector

```
x<-c(1,3)
```

- See the result

```
x
```

```
print(x)
```

```
> x<-c(1,3)
> x
[1] 1 3
> print(x)
[1] 1 3
>
```

- Create an empty vector

```
Y=numeric(10)
```

```
Y
```

```
> Y=numeric(10)
> Y
[1] 0 0 0 0 0 0 0 0 0 0
.
```


Sequence

- Either ' or seq()

R Console

```
> f<-3:5
> f
[1] 3 4 5
> g<-seq(from=3, to=7, by=0.5)
> g
[1] 3.0 3.5 4.0 4.5 5.0 5.5 6.0 6.5 7.0
> |
```


Operation with vectors

- indexing
- Element-wise: $+-*/^{\wedge}$
- log exp sin cos
- length –number of elements
- sum - sum of all elements
- max min sort order
- which.min which.max

Logicals:

TRUE or FALSE:

A=TRUE;

$== > >= < <= != \& \text{ (and)} \mid \text{ (or)}$

```
> a=1:5
> b=c(1,4,-1,3,0)
> a+b
[1] 2 6 2 7 5
> a*b
[1] 1 8 -3 12 0
> b+4
[1] 5 8 3 7 4
> length(a)
[1] 5
> sum(a^2)
[1] 55
> max(b)
[1] 4
> which.max(b)
[1] 2
> order(b)
[1] 3 5 1 4 2
> sort(b)
[1] -1 0 1 3 4
> b[1]
[1] 1
> b[2:4]
[1] 4 -1 3
> b[-2]
[1] 1 -1 3 0
```

Matrices

Use `matrix()`

```
a<-matrix(values,nrow=m,ncol=n)
```

Values should be listed columnwise

`nrow=` and `ncol=` can be skipped

```
R Console
> a<-matrix(c(2,1,1,-1),nrow=2,ncol=2)
> a
      [,1] [,2]
[1,]    2    1
[2,]    1   -1
> |
```

- Create empty matrix

```
> m=matrix(0,nrow=2,ncol=3)
> m
      [,1] [,2] [,3]
[1,]    0    0    0
[2,]    0    0    0
> |
```

Matrix operations

```
> x<-c(1,2)
> a<-matrix(c(2,1,1,-1),2,2)
> b<-matrix(c(1,0,1,1),2,2)
> y=a%*%x
> y
      [,1]
[1,]     4
[2,]    -1
> c=a%*%b
> c
      [,1] [,2]
[1,]     2     3
[2,]     1     0
> |
```

Usual vector operations can also be applied:

```
> m4=matrix(c(1,2,0,1), nrow=2)
> m5=matrix(c(2,2,5,1), nrow=2)
> m4
      [,1] [,2]
[1,]     1     0
[2,]     2     1
> m5
      [,1] [,2]
[1,]     2     5
[2,]     2     1
> m4*m5
      [,1] [,2]
[1,]     2     0
[2,]     4     1
```

Matrix operations

- Matrix operators/functions:

- transpose $b = t(a)$

$$b = a^T$$

- Inverse $b = a^{-1}$

$$b = \text{solve}(a)$$

- Solve $d = a^{-1}b$

$$d = \text{solve}(a, b)$$

```
> a
      [,1] [,2]
[1,]     2     1
[2,]     1    -1
> t(a)
      [,1] [,2]
[1,]     2     1
[2,]     1    -1
> solve(a)
      [,1] [,2]
[1,] 0.3333333 0.3333333
[2,] 0.3333333 -0.6666667
> |
```


Indexing for matrices

- Positive index

`x[1, 6]` `x[2:10,]`

- Negative index

`x[2, -(1:5)]` row 2 and all columns except 1:5

- Entire column or row

`y=x[2,]` entire row 2

- Extraction

`x[x>5]`

```
> b
[1] 1 4 -1 3 0
> d=b[b>0]
> d
[1] 1 4 3
```

Replication

- Replication for vectors
 - rep(what, times)
- Replication for matrices
 - matrix()

```
> v1=rep(3,5)
> v1
[1] 3 3 3 3 3
> v2=rep(c(3,4),2)
> v2
[1] 3 4 3 4
> m1=matrix(1,nrow=2,ncol=2)
> m1
      [,1] [,2]
[1,]    1    1
[2,]    1    1
> m2=matrix(v2,nrow=4,ncol=2)
> m2
      [,1] [,2]
[1,]    3    3
[2,]    4    4
[3,]    3    3
[4,]    4    4
> m3=matrix(v2,nrow=2,ncol=4, byrow=T)
> m3
      [,1] [,2] [,3] [,4]
[1,]    3    4    3    4
[2,]    3    4    3    4
```

Matrix operations

- Dimension
 - `dim(mat)`, `nrow(mat)`, `ncol(mat)`
- Row/column statistics
 - `colMeans`, `rowMeans`, `colSums`, `rowSums`

```
> m4=matrix(c(1,3,5,2,4,6),nrow=3)
> m4
```

```
      [,1] [,2]
[1,]     1     2
[2,]     3     4
[3,]     5     6
```

```
> colMeans(m4)
```

```
[1] 3 4
```

```
> rowSums(m4)
```

```
[1] 3 7 11
```

- Apply a function over vector/matrix
 - `sapply`, `apply`, `lapply`
 - Normally used when function works only element-wise

```
> m2
      [,1] [,2]
[1,]     3     3
[2,]     4     4
[3,]     3     3
[4,]     4     4
> ns=dim(m2)
> ns[2]
[1] 2
```

```
> sapply(v2,log)
[1] 1.098612 1.386294 1.098612 1.386294
> log(v2)
[1] 1.098612 1.386294 1.098612 1.386294
```

Vector/matrix operations

- Create confusion matrix (classification)

- `table(X,Y)`

- Extract diagonal

- `Diag(X)`

```
> X=c(1,3,1,1,2,3,1,2,2,2,1,1,3)
> Xfit=c(2,3,2,1,2,3,1,2,2,2,1,1,1)
> t1=table(Xfit,X)
> t1
      X
Xfit  1 2 3
    1 4 0 1
    2 2 4 0
    3 0 0 2
> t1[1,1]
[1] 4
> diag(t1)
1 2 3
4 4 2
>
```


Factors

- Text values

```
> f1=c("Man", "Woman")
> f1
[1] "Man"    "Woman"
> f2=c("Man", "Woman", "Man")
> table(f2)
f2
  Man Woman
    2     1
> f3=factor(c(1,0,1,1,0), levels=c(0,1), labels=c("Man", "Woman"))
> f3
[1] Woman Man   Woman Woman Man
Levels: Man Woman
>
```

Lists

- List is a collection of objects

```
> d<-15;
> a<-matrix(c(1,2,3,4),2,2);
> a
      [,1] [,2]
[1,]    1    3
[2,]    2    4
> b<-list(first=d, second=a, x="mary")
> b
$first
[1] 15

$second
      [,1] [,2]
[1,]    1    3
[2,]    2    4

$x
[1] "mary"
```

```
> b$x
[1] "mary"
> b[[3]]
[1] "mary"
```

Data frame

Vectors and matrices of the row length can be collected into a data frame

- Used to store the data of different types into a single table

Use `data.frame(object 1, object 2, ... , object k)`

```
> x<-c(1,3)
> y<-c("M", "F")
> z<-data.frame(x,y)
> z
  x y
1 1 M
2 3 F
```

Data frame

- Any column in the data frame can be retrieved by

dataframe\$object

```
> z$x
[1] 1 3
> z[[1]]
[1] 1 3
> z$y
[1] M F
Levels: F M
```

- Any row in the data frame can be extracted by using matrix notation, for ex: **z[1,]**

Read data from CSV file

1. Save as "comma-separated file"(csv)
2. Change current directory, Session→ Set Working Directory or setwd()
3. Use

```
Dataframe=read.csv2(file_name)
```

```
Dataframe=read.csv(file_name)
```

Read from Excel files: packages xlsx, readxl

Conversion between types

- Data frame to matrix
- Matrix to data frame
- Numeric to factor
- Factor to numeric
- List to vector
- Vector to list

```
> df1=data.frame(X=c(1,2,3), Y=c(1,0,1))
> df1
  X Y
1 1 1
2 2 0
3 3 1
> m5=as.matrix(df1)
> m5
      X Y
[1,] 1 1
[2,] 2 0
[3,] 3 1
> df2=as.data.frame(m5)
> df2$X
[1] 1 2 3
```

```
> v6=c(1,4,2,2,1)
> f4=as.factor(v6)
> f4
[1] 1 4 2 2 1
Levels: 1 2 4
> f5=c("1", "0", "1", "1", "1")
> f5
[1] "1" "0" "1" "1" "1"
> as.list(f5)
[[1]]
[1] "1"

[[2]]
[1] "0"

[[3]]
[1] "1"

[[4]]
[1] "1"

[[5]]
[1] "1"

> l1=list(a=1, b=3)
> l1
$a
[1] 1

$b
[1] 3

> as.numeric(l1)
[1] 1 3
```

Loops

```
for (name in expr1 )  
{  
...  
}
```

```
while (condition)  
{  
...  
}
```

```
> for (i in 1:5) {  
+ y=seq(i,8)  
+ print(y) }  
[1] 1 2 3 4 5 6 7 8  
[1] 2 3 4 5 6 7 8  
[1] 3 4 5 6 7 8  
[1] 4 5 6 7 8  
[1] 5 6 7 8  
> |
```

Conditioning and loops

```
If (x==3) {
```

```
...
```

```
...
```

```
} else {
```

```
...
```

```
}
```

```
> m4=matrix(c(1,2,0,1), nrow=2)
> m4
      [,1] [,2]
[1,]    1    0
[2,]    2    1
> n=dim(m4)[1]
> I=numeric(n)
> for (i in 1:n){
+   if(max(m4[i,]>1)) I[i]=1
+ }
> I
[1] 0 1
```


Random number generation

- Random are not random
 - Use `set.seed(12345)` to get identical results
- A plenty of random number generators
 - `Rnorm`
 - `Runif`
 - ...
- Use `d` for density `p` for CDF `q` for quantiles and `r` for simulation:
(ex: `rnorm` `pnorm` `dnorm` `qnorm`)

Distribution	R name	additional arguments
beta	<code>beta</code>	<code>shape1, shape2, ncp</code>
binomial	<code>binom</code>	<code>size, prob</code>
Cauchy	<code>cauchy</code>	<code>location, scale</code>
chi-squared	<code>chisq</code>	<code>df, ncp</code>
exponential	<code>exp</code>	<code>rate</code>
F	<code>f</code>	<code>df1, df2, ncp</code>
gamma	<code>gamma</code>	<code>shape, scale</code>
geometric	<code>geom</code>	<code>prob</code>
hypergeometric	<code>hyper</code>	<code>m, n, k</code>
log-normal	<code>lnorm</code>	<code>meanlog, sdlog</code>
logistic	<code>logis</code>	<code>location, scale</code>
negative binomial	<code>nbinom</code>	<code>size, prob</code>
normal	<code>norm</code>	<code>mean, sd</code>
Poisson	<code>pois</code>	<code>lambda</code>
Student's t	<code>t</code>	<code>df, ncp</code>
uniform	<code>unif</code>	<code>min, max</code>
Weibull	<code>weibull</code>	<code>shape, scale</code>
Wilcoxon	<code>wilcox</code>	<code>m, n</code>

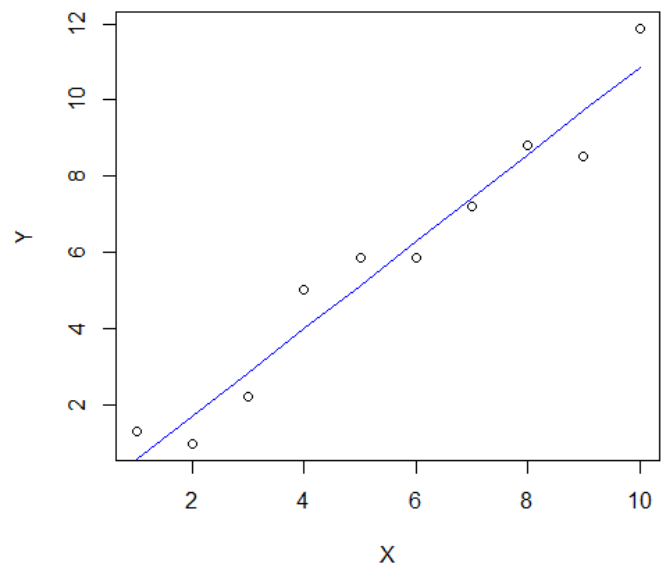
Using a function

- Use `?name_of_function` to see function parameters
 - For ex. `?lm`
- There are some obligatory parameters and optional parameters
- The optional parameters can be specified in different order

```
X=1:10
Y=1:10+rnorm(10)
W=c(rep(1,5), rep(2,5))
mydata=data.frame(X,Y)

result=lm(Y~X, weights=W,data=mydata)
?predict.lm
Fit=predict(result)

plot(X,Y)
points(X,Fit, type="l", col="blue")
```



Writing your own functions

- Function writing must always end with writing the value which should be returned!
- You may also use **"return(value)"** to show what value the function should return

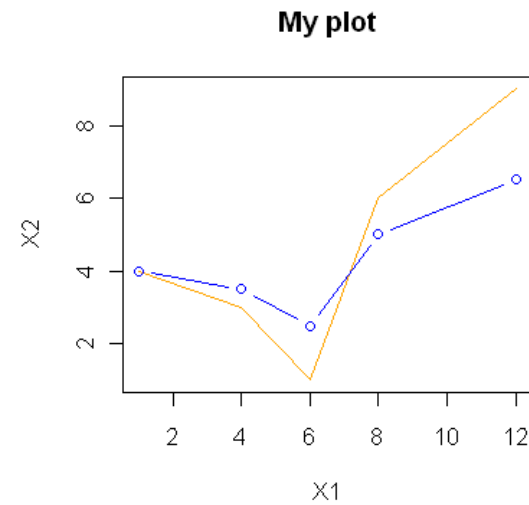
```
> myfun <-function(x=20, y, z)
+ {
+   if(z)
+     result=x+y
+   else
+   {
+     result=log(x)*y
+   }
+   result
+ }
> r<-myfun(1,2, TRUE)
> r
[1] 3
> r<-myfun(z=FALSE, y=0)
> r
[1] 0
> |
```

Graphical procedures

Some common procedures:

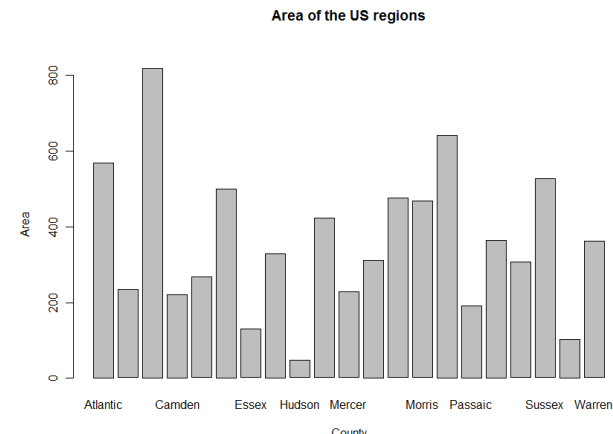
- `plot(x,..)` plots time series
- `plot(x,y)` scatter plot
- `plot(x,y)` followed by `points(x,y)` plots several scatterplots in one coordinate system
- `hist(x,..)` plots a histogram
- `persp(x,y,z,...)` creates surface plots
- `cloud(formula,data..)` creates 3D scatter plot

```
x<-c(1,4,7 8, 12);  
y<-c(4,3,1,6,9);  
  
plot(x,y, type="l", col="orange",  
      main="My plot", xlab="X1", ylab="X2");  
points(x, y/2+2, type="b", col="blue");
```



Graphical parameters

- Adjust color of a graphical object by specifying
 - col=
- Other typical parameters for graphical functions
 - main="text" Title "text"
 - sub="text" Footnote "text"
 - xlab="text" X-axis label
 - ylab="text" Y-axis label

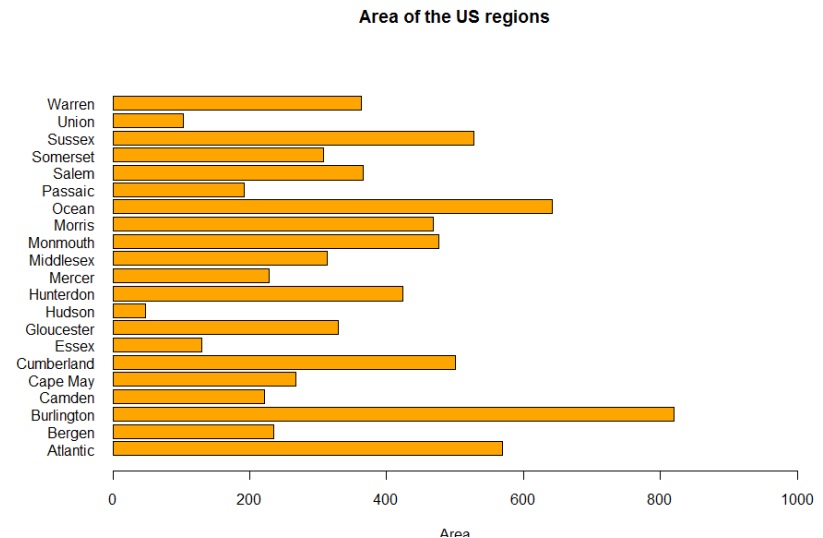


```
mydata<-read.csv2("Counties.csv");  
barplot(mydata$Area, names.arg=mydata$County, main="Area of the US regions",  
xlab="County", ylab="Area");
```

Graphical parameters

- Some parameters need to be specified either in the plotting function or inside `par(...)`
 - `Pch=number` – symbol that is plotted
 - `Lty=number` – linetype
 - `Las=0 1 eller 2` Direction of axis values
 - `mai=c(bottom, left, top, right)` – margins (inch)
 - `adj=`between 0 and 1, horizontal justification

```
barplot(mydata$Area,  
names.arg=mydata$County, horiz=TRUE, las=1,  
xlim=c(0,1000), col="orange", main="Area of  
the US regions", xlab="Area");
```



Some more examples

- Dividing training/test

```
data=data.frame(X=c(1,1,2,2,3), Y=c("M","F","M","M","F"))
n=dim(data)[1]
set.seed(12345)
id=sample(1:n, floor(n*0.5))
train=data[id,]
test=data[-id,]
```

```
> train
  X Y
4 2 M
5 3 F
> test
  X Y
1 1 M
2 1 F
3 2 M
```

- Computing misclassification rate

```
missclass=function(X,X1){
  n=length(X)
  return(1-sum(diag(table(X,X1)))/n)
}
```

```
> X=c(1,3,1,1,2,3,1,2,2,2,1,1,3)
> Xfit=c(2,3,2,1,2,3,1,2,2,2,1,1,1)
> missclass(X,Xfit)
[1] 0.2307692
```

More about R

- **Important:** read manuals of **dplyr** and **tidyr** packages (data manipulation)
- R language manual <https://cran.r-project.org/doc/manuals/r-release/R-intro.pdf>