

# Rcourse: Programming in R

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- 1 Back to input files
  - Review on data frame
  - Factors
  
- 2 Programming
  - Conditional execution
  - Loops
  - Executing a command from a script
  - Writing your own functions
  - `lapply()` and `tapply()`
  - How to avoid slow R code

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# Review on data frame

Generic functions:

```
> read.table()  
> write.table()
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Example 1: wghtcls "smoker" lifespan

```
"3" 0 50.3  
3 0 52.8
```

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Example 1: wghtcls "smoker" lifespan

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"3" 0 50.3
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```
3 0 52.8
```

```
> riscfactor <-  
read.table("lifespandata2.txt",header=TRUE)
```

# Review on data frame

Example 2: `wghtcls,smoker,lifespan`

3,0,50.3

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3,0,52.8

```
> riscfactor <- read.csv("lifespandata.csv")
```

```
> riscfactor <-
```

```
read.table("lifespandata.csv",header=TRUE, sep=",",  
fill=TRUE)
```



# Review on data frame

Example 2: wghtcls,smoker,lifespan

3,0,50.3

3,0,52.8

```
> riscfactor <- read.csv("lifespandata.csv")  
> riscfactor <-  
read.table("lifespandata.csv",header=TRUE, sep=",",  
fill=TRUE)
```

Example 3: weight class smoker lifespan

3 0 50.3

3 0 52.8

```
> riscfactor <-  
read.table("lifespandata2.txt",header=TRUE)
```

# Review on data frame

Example 2: wghtcls,smoker,lifespan

3,0,50.3

3,0,52.8

```
> riscfactor <- read.csv("lifespandata.csv")  
> riscfactor <-  
read.table("lifespandata.csv",header=TRUE, sep=",",  
fill=TRUE)
```

Example 3: weight class smoker lifespan

3 0 50.3

3 0 52.8

```
> riscfactor <-  
read.table("lifespandata2.txt",header=TRUE)
```

You have to change the first line of the file because of the space between weight and class.

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# Factors

A variable (numeric or text) can be intended as a factor.

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Example with text:

```
> x <- c("female","male","male","female","female")
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NULL
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> x <- c("female","male","male","female","female")
> levels(x)
NULL
> str(x)
```



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Example with text:

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> x <- c("female","male","male","female","female")
> levels(x)
NULL
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chr [1:5] "female" "male" "male" "female" "female"
```

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A variable (numeric or text) can be intended as a factor.

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> x <- c("female","male","male","female","female")
> levels(x)
NULL
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chr [1:5] "female" "male" "male" "female" "female"
> x <-factor(x)
```

# Factors

A variable (numeric or text) can be intended as a factor.

Example with text:

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> x <- c("female","male","male","female","female")
> levels(x)
NULL
> str(x)
chr [1:5] "female" "male" "male" "female" "female"
> x <-factor(x)
> levels(x)
[1] "female" "male"
> str(x)
Factor w/ 2 levels "female","male":  1 2 2 1 1
```

# Factors

Example with numbers:

```
> y <- rep(c(17,17,18),4); str(y)
num [1:12] 17 17 18 17 17 18 17 17 18 17 ...
```

# Factors

Example with numbers:

```
> y <- rep(c(17,17,18),4); str(y)
num [1:12] 17 17 18 17 17 18 17 17 18 17 ...
> summary(y)
Min. 1st Qu. Median Mean 3rd Qu. Max.
17.00 17.00 17.00 17.33 18.00 18.00
```

# Factors

Example with numbers:

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num [1:12] 17 17 18 17 17 18 17 17 18 17 ...
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17.00 17.00 17.00 17.33 18.00 18.00
> y <- factor(y); str(y)
Factor w/ 2 levels "17","18":  1 1 2 1 1 2 1 1 2 1 ...
> summary(y)
17 18
 8  4
```

# Back to input files

By default `read.table()` sets text variables as factors and not numerical variables.

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This can be changed by specifying the class of the columns.

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riscfactor <-  
read.table("lifespandata.txt", header=TRUE,  
colClasses=c("factor", "numeric", "numeric"))
```



# Back to input files

By default `read.table()` sets text variables as factors and not numerical variables.

This can be changed by specifying the class of the columns.

```
riscfactor <-  
read.table("lifespandata.txt", header=TRUE,  
colClasses=c("factor", "numeric", "numeric"))
```

Or by changing the variables afterwards.

```
riscfactor$wghtcls <- factor(riscfactor$wghtcls)
```

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# Conditional execution

If(), else() and ifelse()

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If(), else() and ifelse()

Syntax:

```
if ( condition ) { commands1 }
```

```
if ( condition ) { commands1 } else { commands2 }
```

```
ifelse ( conditions vector, yes vector, no vector )
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if ( condition ) { commands1 } else { commands2 }
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ifelse ( conditions vector, yes vector, no vector )
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Example:

```
> x <- 4
```

```
> if (x==5) {x <- x+1} else {x <- x*2}
```

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If(), else() and ifelse()

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if ( condition ) { commands1 }
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if ( condition ) { commands1 } else { commands2 }
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ifelse ( conditions vector, yes vector, no vector )
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Example:

```
> x <- 4
```

```
> if (x==5) {x <- x+1} else {x <- x*2}
```

```
> x
```

```
[1] 8
```

# Conditional execution

Other examples:

```
if ( x != 5 & x>3 ) { x <- x+1 ; 17+2 } else { x <- x*2  
; 21+5 }
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if ( x != 5 & x>3 ) { x <- x+1 ; 17+2 } else { x <- x*2  
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[1] 19
```

```
> x
```

```
[1] 9
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if ( x != 5 & x>3 ) { x <- x+1 ; 17+2 } else { x <- x*2  
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```
[1] 19
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```

```
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```
> y <- 1:10
```

```
> ifelse( y<6, y^2, y-1 )
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if ( x != 5 & x>3 ) { x <- x+1 ; 17+2 } else { x <- x*2  
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[1] 19
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```

```
[1] 9
```

```
> y <- 1:10
```

```
> ifelse( y<6, y^2, y-1 )
```

```
[1] 1 4 9 16 25 5 6 7 8 9
```

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# Loops

For(), while() and repeat()

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Syntax:

```
for ( var in set ) { commands }
```

```
while ( condition ) { commands }
```

```
repeat { commands }
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# Loops

For(), while() and repeat()

Syntax:

```
for ( var in set ) { commands }  
while ( condition ) { commands }  
repeat { commands }
```

**break** stops all loops

**next** goes directly to the next iteration of the loop

# Examples

```
> x <- 0 > for ( i in 1:5 ) { if (i==3) { next } ; x <-  
x + i }
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> for ( i in 1:5 ) { if (i==3) { next } ; x <-
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# i=3 is skipped, so x <- 1+2+4+5
> x
[1] 12
```



# Examples

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```

```
# i=3 is skipped, so x <- 1+2+4+5
```

```
> x
```

```
[1] 12
```

```
> y <- 1; j <- 1
```

```
> while ( y < 12 & j < 8 ) { y <- y*2 ; j <- j + 1 }
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```
y is 16 and j is 4
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> x <- 0 > for ( i in 1:5 ) { if (i==3) { next } ; x <-  
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> x
```

```
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> z <- 3
```

```
> repeat { z<- z^2; if ( z>100 ) { break }; print(z)}
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> x <- 0 > for ( i in 1:5 ) { if (i==3) { next } ; x <-  
x + i }
```

# i=3 is skipped, so x <- 1+2+4+5

```
> x
```

```
[1] 12
```

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> y <- 1; j <- 1
```

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> while ( y < 12 & j < 8 ) { y <- y*2 ; j <- j + 1}
```

y is 16 and j is 4

```
> z <- 3
```

```
> repeat { z<- z^2; if ( z>100 ) { break }; print(z)}
```

```
[1] 9
```

```
[1] 81
```

The loop stopped after  $81^2$  so z is 6561.

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# Executing a command from a script

R scripts are stored in `.R` or `.r` files and are executed with the command `source()`

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source(C:/Documents/R/myscript.R)
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You can specify the current working directory using the command `setwd()`



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R scripts are stored in .R or .r files and are executed with the command `source()`

```
source(C:/Documents/R/myscript.R)
```

You can specify the current working directory using the command `setwd()`

```
setwd(C:/Documents/R)
```

```
getwd()
```

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# Basics

Syntax:

```
myfun <- function (arg1, arg2, . . .) { commands }
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## Syntax:

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## Examples:

```
se <- function(x)
{
y<-sqrt(var(x)/length(x))
return(y)
}
```

# Basics

## Syntax:

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myfun <- function (arg1, arg2, . . .) { commands }
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## Examples:

```
se <- function(x)
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se(1:4)
[1] 0.6454972
```

# Basics

Syntax:

```
myfun <- function (arg1, arg2, . . .) { commands }
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Examples:

```
se <- function(x)
{
y<-sqrt(var(x)/length(x))
return(y)
}
se(1:4)
[1] 0.6454972
se("wrong type of argument")
[1] NA
```

Warning message:

```
In var(x) : NAs introduced by coercion
```

# Deal with non correct arguments

```
if (is.numeric(x)!=TRUE)
{
stop("need numeric data")
}
```

# Deal with non correct arguments

```
if (is.numeric(x)!=TRUE)
{
  stop("need numeric data")
}

  se("wrong type of argument")
Error in se("wrong type of argument") : need numeric
data
```



# Deal with the missing data

```
x<-x[is.na(x)==FALSE]
```

# Deal with the missing data

```
x<-x[is.na(x)==FALSE]
```

```
se(c(1:4,NA))
```

```
[1] 0.6454972
```

# Giving default values to the arguments

```
se <- function(x,na.rm=FALSE)
{ if (is.numeric(x)!=TRUE) {stop("need numeric data")}
  if (na.rm==TRUE){x<-x[is.na(x)==FALSE]}
  y<-sqrt(var(x)/length(x))}
```

# Giving default values to the arguments

```
se <- function(x,na.rm=FALSE)
{ if (is.numeric(x)!=TRUE) {stop("need numeric data")}
  if (na.rm==TRUE){x<-x[is.na(x)==FALSE]}
  y<-sqrt(var(x)/length(x))}
```

You can omit to write the name of the argument:

```
se(c(NA,1:4), TRUE))
[1] 0.6454972
```

# Giving default values to the arguments

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se <- function(x,na.rm=FALSE)
{ if (is.numeric(x)!=TRUE) {stop("need numeric data")}
  if (na.rm==TRUE){x<-x[is.na(x)==FALSE]}
  y<-sqrt(var(x)/length(x))}
```

You can omit to write the name of the argument:

```
se(c(NA,1:4), TRUE))
[1] 0.6454972
```

Or give na.rm before the vector.

But not both (omitting name and changing the order of arguments).

# The "...” argument

The "...” argument allows you to pass arguments from one function to another.

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Example:

```
se.sq <- function(x, ...)  
{  
  y<-se(x, ...)  
  return(y^2)  
}
```

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Example:

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se.sq <- function(x, ...)  
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```

```
se.sq(1:4)  
[1] 0.4166667
```



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Example:

```
se.sq <- function(x, ...)  
{  
  y<-se(x, ...)  
  return(y^2)  
}
```

```
se.sq(1:4)  
[1] 0.4166667  
se.sq(c(1:4, NA))  
[1] NA
```

# The "...” argument

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Example:

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se.sq <- function(x, ...)  
{  
  y<-se(x, ...)  
  return(y^2)  
}
```

```
se.sq(1:4)  
[1] 0.4166667  
se.sq(c(1:4, NA))  
[1] NA  
se.sq(c(1:4, NA))  
[1] 0.4166667
```

# Returning several values

To do so use a vector or a list.

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```
ci.norm <- function(x,conf=0.95)
{
q <- qnorm(1-(1-conf)/2)
return(
list(lower=mean(x)-q*se(x),upper=mean(x)+q*se(x)))
}
```

# Returning several values

To do so use a vector or a list.

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ci.norm <- function(x,conf=0.95)
{
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return(
list(lower=mean(x)-q*se(x),upper=mean(x)+q*se(x)))
}
```

```
ci.norm(rnorm(100))
$lower [1] -0.1499551
$upper [1] 0.2754680
```

```
ci.norm(rnorm(100,conf=0.99))
$lower [1] -0.1673693
$upper [1] 0.2443276
```

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# lapply() and tapply()

You use `apply()` and its derivatives to apply the same function to each element of an object.

```
v <- 1:4
```

```
sapply(v,factorial)
```

```
# returns a vector, lapply() would return a list
```

```
[1] 1 2 6 24
```

# lapply() and tapply()

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# returns a vector, lapply() would return a list
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[1] 1 2 6 24
```

`tapply()` is used for data frames.



# apply() and tapply()

You use `apply()` and its derivatives to apply the same function to each element of an object.

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v <- 1:4  
sapply(v,factorial)  
# returns a vector, lapply() would return a list  
[1] 1 2 6 24
```

`tapply()` is used for data frames.

Example: data frame containing lifespan for people from 3 classes of weight. You want the mean lifespan for each class.

```
tapply(lifespan,weightcls,mean)  
1 2 3  
69 61 53
```

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# How to avoid slow R code

- R has to interpret your commands each time you run a script and it takes time to determine the type of your variables.
- So avoid using loops and calling functions again and again if possible
- Think in whole objects such as vectors or lists and apply operations to the whole object instead of looping through all elements.

# How to avoid slow R code

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- So avoid using loops and calling functions again and again if possible
- Think in whole objects such as vectors or lists and apply operations to the whole object instead of looping through all elements.

```
x <- 0  
for(i in 1:1000000) {x <- x+i}
```

```
sum(as.numeric(1:1000000))
```

Try on your computer!