

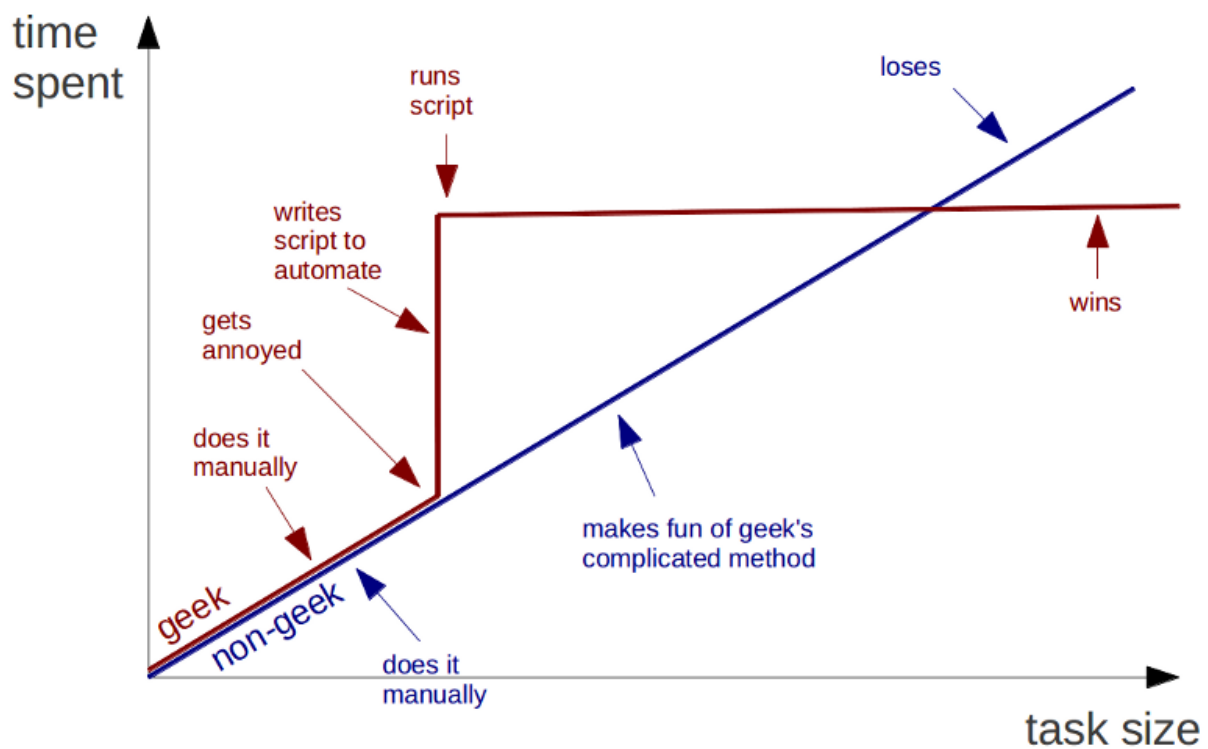
# Repeat “things” in R - loops, functions, apply

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## Introduction

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### Geeks and repetitive tasks



## Why?

- Fewer mistakes
- Your future self

## Tools

- Write a loop

```
for (x in 1:length(input)){
  boxplot(input[,x]~category)
}
```

- Write a function

```
correction<-function(input,correction){
  input-correction
}
```

=====

- Use the apply family

```
apply(input,2,mean)
```

## My examples

- Read Data
- Treat Data
- Create Graphs

## My examples

- Read Data
- **Treat Data**
- **Create Graphs**

Dataset: Doubs river chemistry, provided in ADE4 library

	dfs	alt	slo	flo	pH	har	pho	nit	amm	oxy	bdo
1	3	934	6.176	84	79	45	1	20	0	122	27
2	22	932	3.434	100	80	40	2	20	10	103	19
3	102	914	3.638	180	83	52	5	22	5	105	35
4	185	854	3.497	253	80	72	10	21	0	110	13
5	215	849	3.178	264	81	84	38	52	20	80	62
6	324	846	3.497	286	79	60	20	15	0	102	53

## Functions

- Break long script into small, formalized operations

Why? - Readability - Avoid errors - Facilitate code recycling

# Functions

Define a function

```
function.name<-function(arg1, arg2, arg3=5,...){  
  newVal<-arg1+arg2  
  newVal/arg3  
}
```

Source from file or execute in console

```
function.name(arg1,arg2)
```

## Functions - Example

Define a function

```
np_ratio<-function(arg1, arg2, arg3,...){  
  newVal<-arg1+arg2  
  newVal/arg3  
}
```

```
ratio_doubs<-n_pratio(doubs$env$nit,  
                      doubs$env$amm,  
                      doubs$env$pho)
```

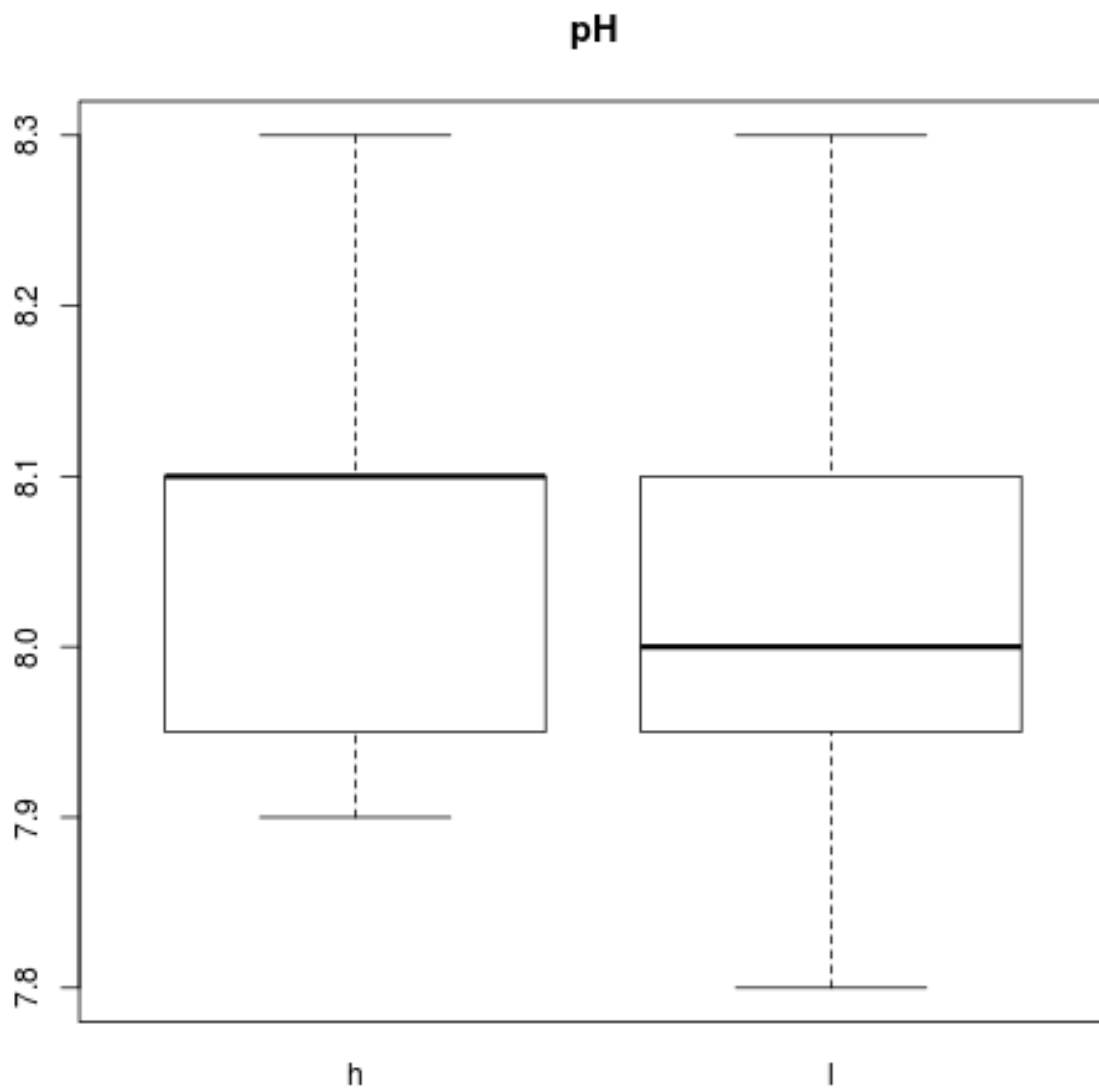
## Functions - Example2

id: function Define a function

```
bp_me<-function(x,cat,t,  
               cnames=levels(as.factor(cat))){  
  #make a boxplot  
  boxplot(x~cat,xlab="",ylab="",names.arg=NA,  
          axes=F,outline=F)  
  #do some fine tuning  
  par(mgp=c(3,1,0))  
  axis(2)  
  #par(mgp=c(3,0,0))  
  par(las=1)  
  axis(side=1,at=c(1:length(cnames)),  
       labels=cnames,lty=0)  
  #par(mgp=c(3,1,0))  
  par(las=2)  
  box()  
  title(main=t)  
}
```

## Functions - Example2

```
categs<-ifelse(  
  doubs$env$salt>=median(doubs$env$salt),"h","l")  
bp_me(doubs$env$pH/10,categs,"pH")
```



## Functions - General remarks

- Choose clear names, make them easy to understand
- Break long functions into small steps
- Avoid using global variables

- use ... in the function definition to allow flexible input

## Loops

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- Avoid!
- Very versatile
- Slow, more difficult to recycle and read

## Loops

```
par(mfrow=c(2,2),las=2,oma = c(1.2,2, 1, 0.5),mar=c(2,4,2,1))
input<-doubs$env[,c("har","pho","nit","amm")]
input[,2:4]<-input[,2:4]/100
```

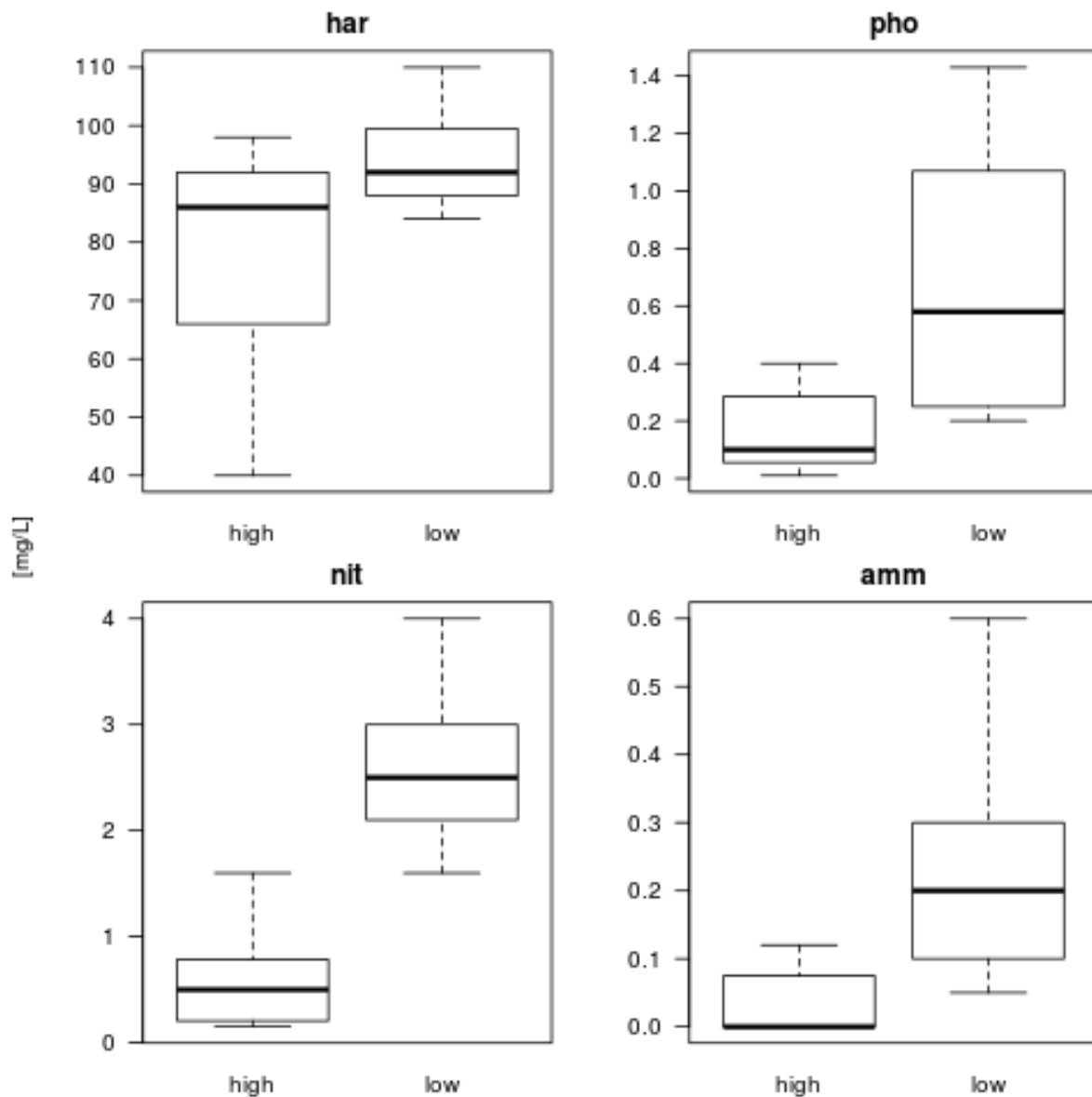
## Loops

title:false

```
for (x in 1:length(input)){
  boxplot(input[,x]~categs,xlab="",ylab="",
          names.arg=NA,axes=F,outline=F)
  axis(2)
  par(las=1)
  axis(side=1,at=c(1:length(
    levels(as.factor(categs)))),
    labels=c("high","low"),lty=0)
  box()
  title(colnames(input)[x])
}
par(las=0)
mtext("[mg/L]", side=2,outer = TRUE, cex = 0.8,)
```

## Loops

title:false



## The apply family

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- Apply a function to a series of objects
- Input and output can vary
- Used in the dplyr package to construct some useful tools

## apply()

```
apply(x,MARGIN,fun(x))
```

```
a<-apply(doubs$env,2,mean)
b<-apply (doubs$fish,1,sum)
head(as.data.frame(b))
```

```
      b
1  3
2 12
3 16
4 21
5 34
6 21
```

```
=====
```

```
head(as.data.frame(b))
```

```
      b
1  3
2 12
3 16
4 21
5 34
6 21
```

## lapply()

```
lapply(X,fun(x))
```

## lapply()

```
title:false
```

```
lapply(as.list(doubs$env),mean)
```

```
$dfs
[1] 1879.033
```

```
$alt
[1] 481.5
```

```
$slo
[1] 2.757733
```

```
$flo  
[1] 2220.1
```

```
$pH  
[1] 80.5
```

```
$har  
[1] 86.1
```

```
$pho  
[1] 55.76667
```

```
$nit  
[1] 165.4
```

```
$amm  
[1] 20.93333
```

```
$oxy  
[1] 93.9
```

```
$bdo  
[1] 51.16667
```

## lapply()

title:false

```
a<-lapply(doubs,function(x){apply(x,2,mean)})  
summary(a)
```

	Length	Class	Mode
env	11	-none-	numeric
fish	27	-none-	numeric
xy	2	-none-	numeric
species	4	-none-	numeric

## mapply()

id: application

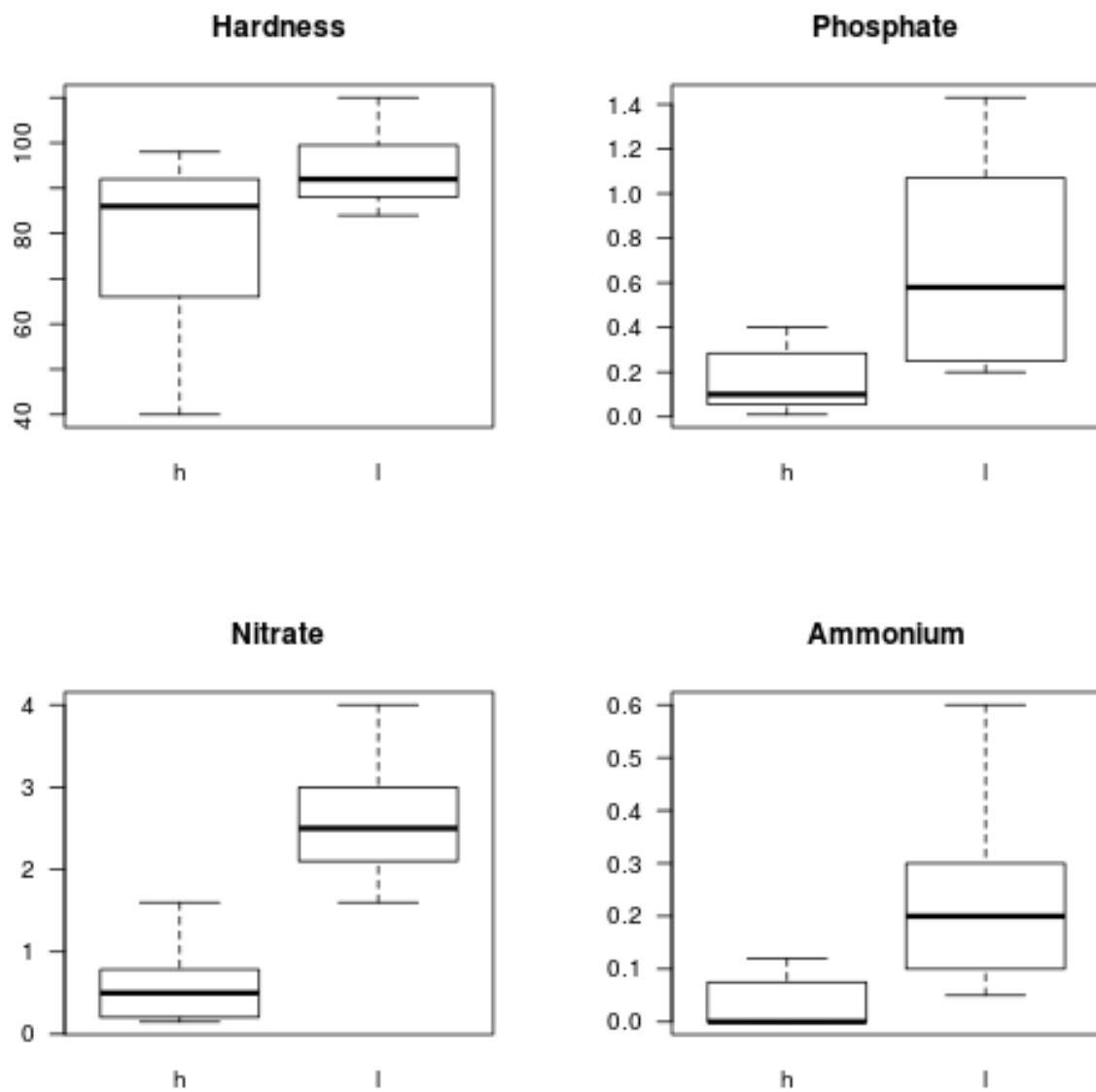
```
mapply(FUN, arg1, arg2,..., MoreArgs = NULL)
```

bp\_me definition

```
titles<-c("Hardness","Phosphate","Nitrate","Ammonium")  
par(mfrow=c(2,2))  
mapply(bp_me,x=input,t=titles,MoreArgs=list(cat=categs))  
summary(a)
```



mapply()



\$har  
NULL

\$pho  
NULL

\$nit  
NULL

\$amm  
NULL

	Length	Class	Mode
env	11	-none-	numeric
fish	27	-none-	numeric
xy	2	-none-	numeric
species	4	-none-	numeric

## Ressources

Two blog posts that I like

<http://nicercode.github.io/guides/repeating-things/> <http://nicercode.github.io/guides/functions/>

And for the advanced:

<http://adv-r.had.co.nz/Functions.html>

And to do nice R presentations wit R Studio:

<http://support.rstudio.com/hc/en-us/articles/200486468> [http://rmarkdown.rstudio.com/authoring\\_pandoc\\_markdown.html](http://rmarkdown.rstudio.com/authoring_pandoc_markdown.html)