

3 Mars 2021 DU Bioinformatique intégrative Module 3: « R et statistiques »





en explorant des données omiques

Teachers: Anne Badel, Claire Vandiedonck

Helpers: Antoine Bridier-Nahmias, Clémence Réda, Bruno Toupance, Jacques van Helden

Plan de la session 1: R base

- 1. Start-R: introduction au module 3
- 2. Vérification et consolidation des pré-recquis:
 - a. Session R
 - b. Vecteurs
 - c. Matrices

Objectifs d'apprentissage de la session 3

3. Dataframes

- a. Créer un dataframe
- b. Extraire des données d'un dataframe
- c. Manipuler un dataframe: filtrer, créer des sous-dataframes, fusionner des dataframes
- => Travail personnel ce vendredi:
 - a. consolidation des bases de R:
 - vecteurs, matrices, dataframes
 - statistiques decsriptives et figures de base selon le type de données
 - b. facteurs: découverte par un tutorial
 - c. listes: diapo et tutorial

1. Start-R

introduction au module 3 « R - stats »

Equipe pédagogique

Responsables:



Claire Vandiedonck (MCF UP)



Jacques van Helden (Pr AMU, directeur IFB)

Instructeurs:



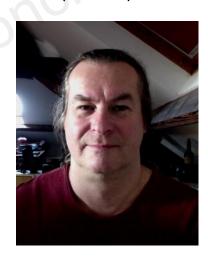
Anne Badel



Magali Berland (IR INRA MétaGénoPolis),



Antoine Bridier-Nahmias (MCF UP)



Olivier Sand (IR CNRS, IFB-core)



Bruno Toupance (MCF UP),



Clémence Réda (DCEM UP),



Yves Clément (MCF UP),



Olivier Taboureau (Pr UP)

Planning du module 3 R-Stats

Jour	Horaire	Description	Instructeurs	Helpers
3 mars	9h30 - 12h30	 R base Manipuler les différents types d'objets dans R Statistiques descriptives Figures de base 	Anne Badel Claire Vandiedonck	Antoine-Bridier-Nahmias Bruno Toupance Clémence Réda Jacques van Helden
4 mars	13h30 - 16h30	Renforcement de R: - Contrôles de flux: exécutions conditionnelles, boucles - Paquets, écrire ses propres fonctions - Customiser ses figures avec R base - Introduction à dplyr/tidyverse/ggplot	Magali Berland Claire Vandiedonck	Antoine-Bridier-Nahmias Yves Clément Bruno Toupance Jacques van Helden
9 mars	14h30 - 17h30	Statistiques pour les données à haut débit RStudio et R markdown	Antoine Bridier-Nahmias Claire Vandiedonck	Anne Badel Clémence Réda Olivier Sand Jacques van Helden
11 mars	9h00 - 12h00	Régression linéaire Corrélation Exploration de données multidimensionnels (ACP/MDS)	Magali Berland Jacques van Helden	Anne Badel Clémence Reda Olivier Taboureau Claire Vandiedonck
29 mars	10h00 - 13h00	Classification supervisée et apprentissage	Olivier Sand Jacques van Helden	Anne Badel Olivier Taboureau Bruno Toupance Claire Vandiedonck
29 mars	14h30 - 17h30	Classification non supervisée (clustering) Analyse d'enrichissment	Anne Badel Olivier Sand Jacques van Helden	Yves Clément Olivier Taboureau Bruno Toupance Claire Vandiedonck

Site web du cours

https://du-bii.github.io/module-3-Stat-R/

module-3-Stat-R



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View the Project on GitHub DU-Bii/module-3-Stat-R

This project is maintained by DU-Bii

Hosted on GitHub Pages — Theme by orderedlist

DUBii - module 3 - Analyse statistique avec R

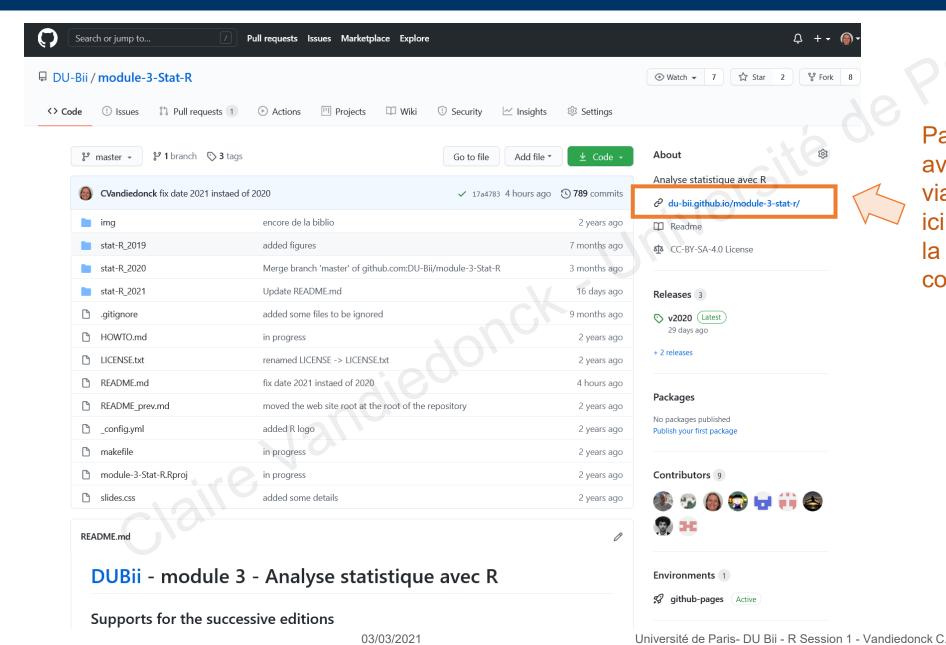
Supports for the successive editions

Edition	Site
2021	stat-R_2021/
2020	stat-R_2020/
2019	stat-R_2019/

Links

Doc	Description	URL
Git pages	Web site of the course (to see the supports)	https://du-bii.github.io /module-3-Stat-R/
Git repo	Repository enabling to download or clone the teaching material on your computer	https://github.com/DU- Bii/module-3-Stat-R

Retour vers le site web depuis le dépôt Github du DU



Pas de panique si vous avez atterri sur cette page via google, cliquez juste ici et vous reviendrez sur la page web correspondante!

Modalités de contrôle des connaissance

Assiduité (25%)

Travail personnel sous forme d'atelier (15%)

1. rendu individuel (7.5%)pour le **19/03** -

2. évaluation par les pairs (7.5%) pour le **02/04** -

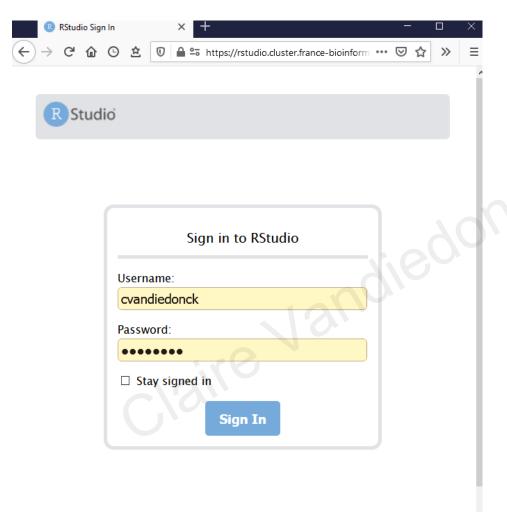
Mini-projet (50%) pour le **10/05** -

- 29/03 matin QCM sur moodle (10%)

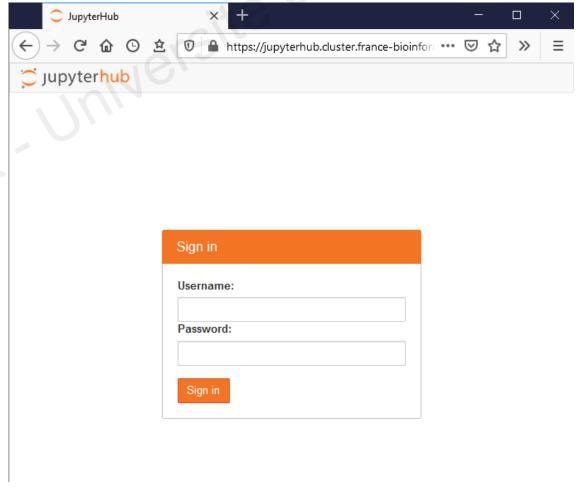
Ressources de l'IFB pour R:

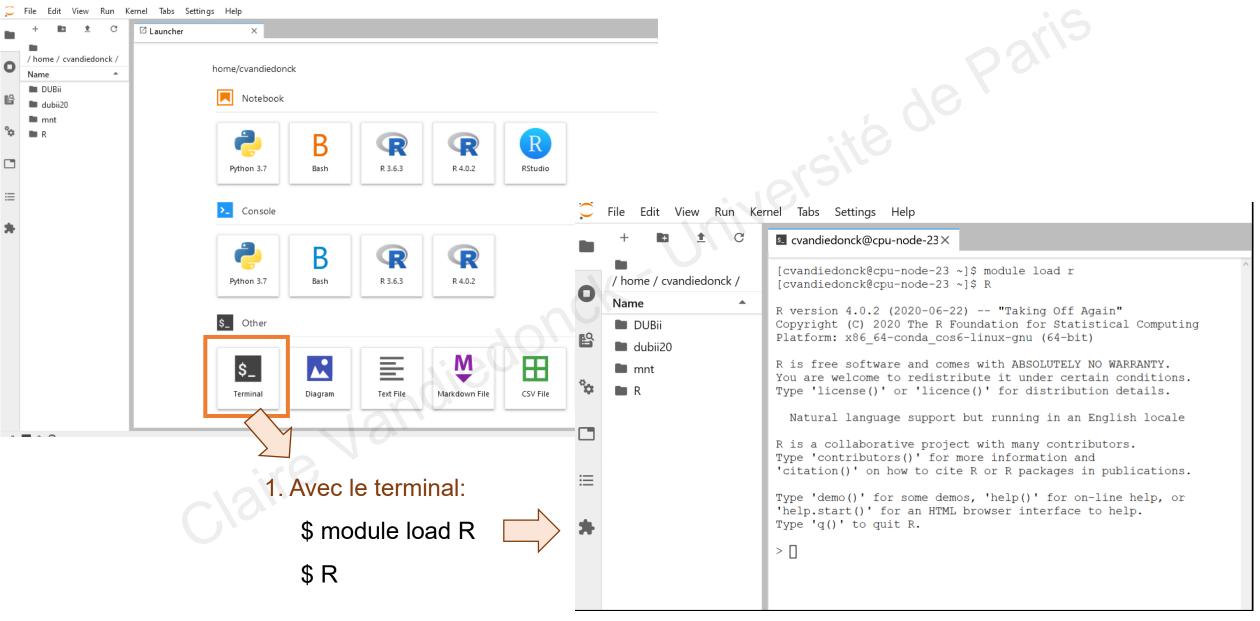
Deux plateformes avec vos mêmes identifiants!

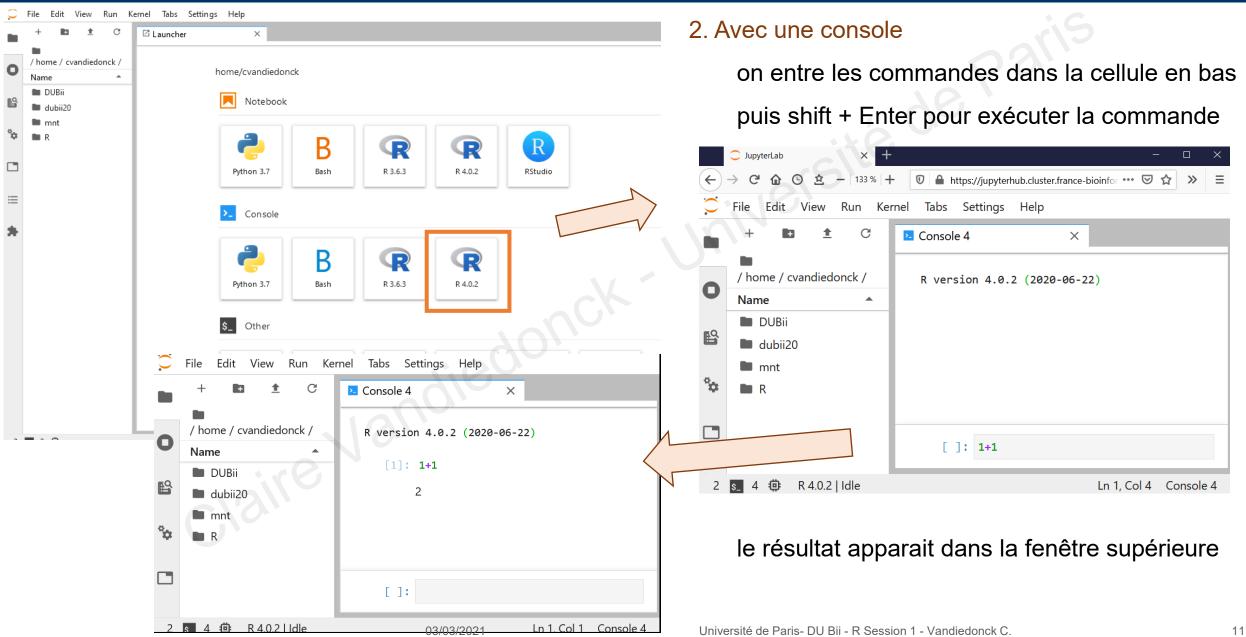
https://rstudio.cluster.france-bioinformatique.fr/



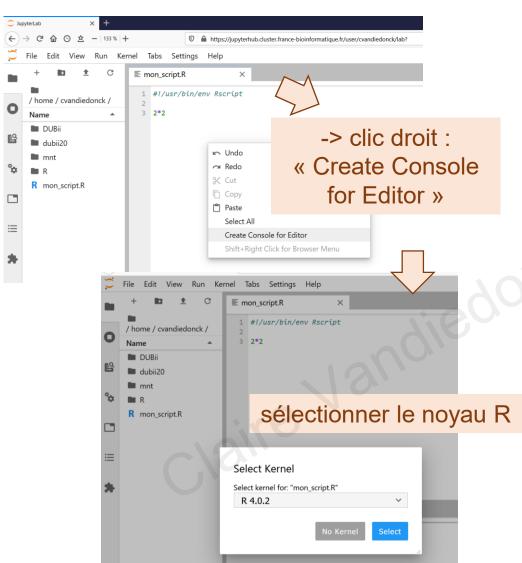
https://jupyterhub.cluster.france-bioinformatique.fr/







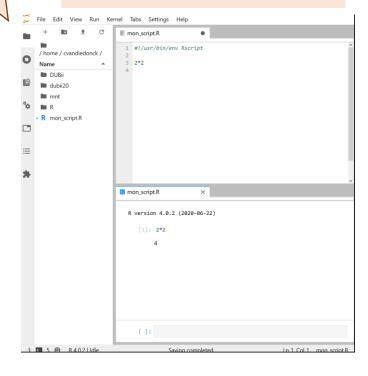
Ouvrez un script

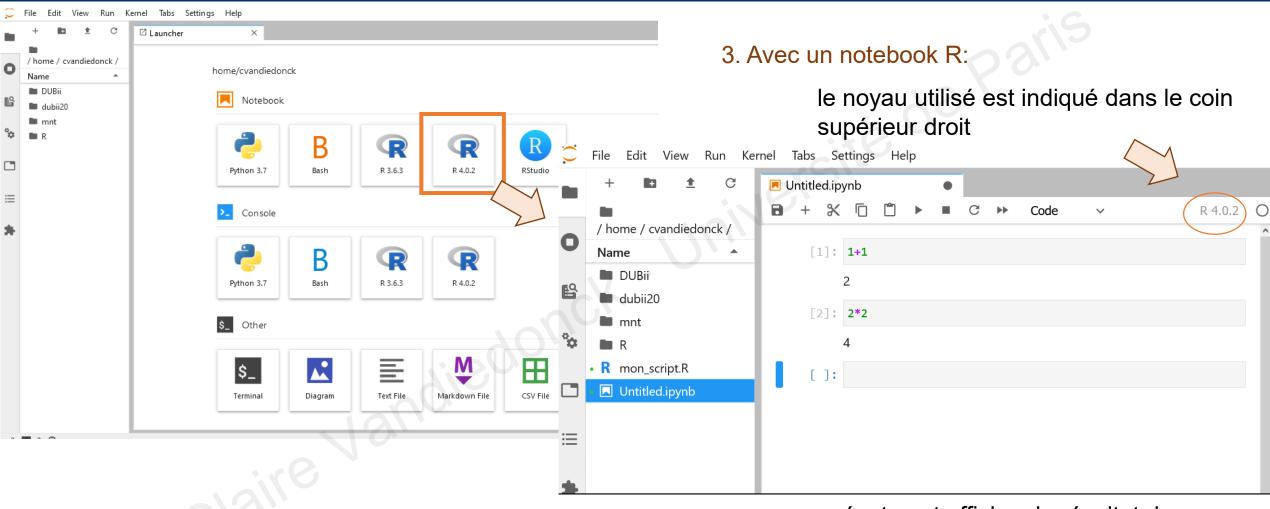


2. Avec une console et un script!

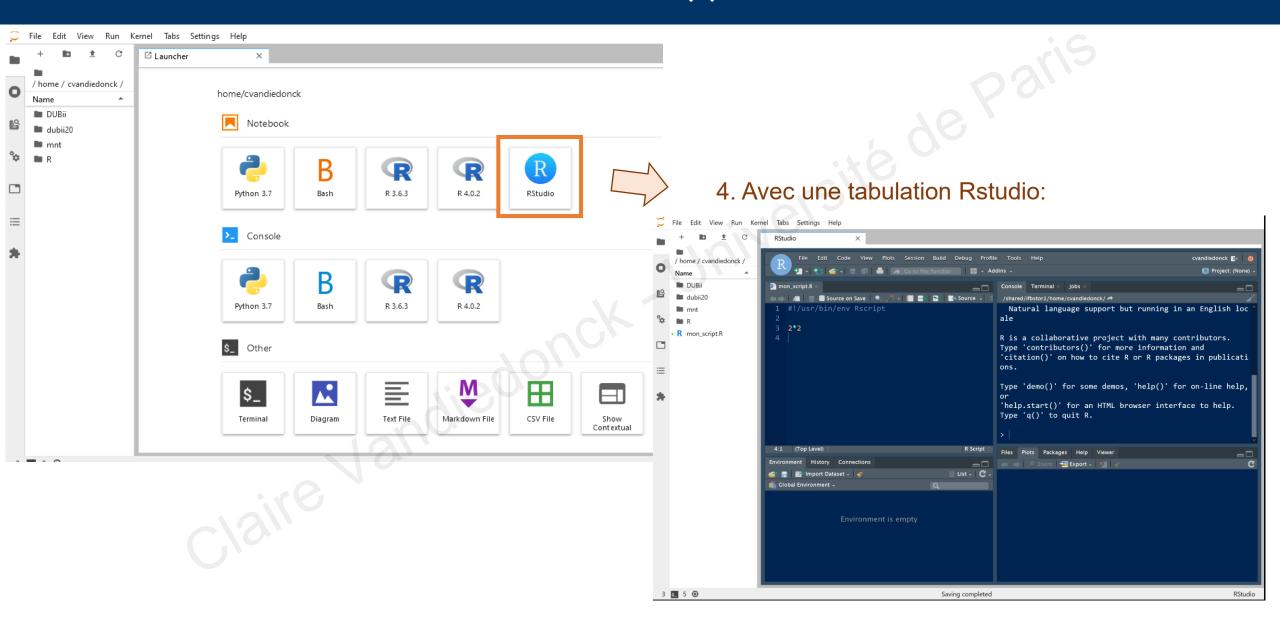


exécuter ensuite les commandes du script dans la console avec shift+Entrée





exécutez et affichez le résultat de vos cellules une par une directement dans le notebook!



Sondage wooclap -> R et vous?





2. Vérification et consolidation des pré-requis:

Session R Vecteurs Matrices

Rappel sur une Session R

sessionInfo()	R version, core packages, version of additional loaded packages
getwd()	get the working directory of the R session on my computer
setwd()	set the working directory of the R session on my computer
ls()	list objects and functions present in my R session
list.files() or dir()	list files and folders in the given directory (by default the working directory) on my computer
save()	to save some objects/functions of my R session on the computer
save.image()	to save all objects/functions of my R session on the computer

data/"data" no quotes if the object is in the R Session, quotes if calling a file which is not in the R Session

Caution when naming R objects/variables: no accent, no special characters (like "-"), cannot start with a number... cf. https://google.github.io/styleguide/Rguide.xml

Rappel sur les types d'objets

Main data structures

object	Heterogeneous = several types may coexist		
vector	no		
matrix	no		
dataframe	yes		
list	yes		

Summary on vectors

```
one-dimension
Format
                     homogeneous: only one type of character, numeric, logical, factor...
Datatype
                     -> ceorcion if heterogeneous
                               - check with class() or mode()
                               - checking type with is.num(), is.charachter(), ...
                               - conversion with as.num(), as.charachter(), ...
                    c(),:,seq(), rep(),sample(),rnorm(),...
Creation
Adding new items
Size
                     length()
Slicing
                    my_vector[i]
                     my_vector[i] <- "toto"
Filling
Naming
                     names()
```

Let's play together!

Exercice sur les vecteurs

Pour chaque question posée, entrez vos solutions de code (ou likez celle de vos collègues):

- 1. Créer un vecteur avec: 5 4 3 2 1 et 0. J'attends au moins 3 façons possibles.
- 2. Créer un vecteur avec: 1_impair, 2_pair, 3_impair, 4_pair, 5_impair, 6_pair
- 3. Ecrire dans R « Il y a 26 lettres dans l'alphabet » en codant la valeur 26
- 4. Comptez le nombre de caractères de la phrase en Q3
- 5. Ecrire la même phrase en lettres capitales.
- 6. Créez un **vecteur x** contenant 100 valeurs tirées aléatoirement à partir des valeurs -10 à 10 et de 10 valeurs manquantes.
- 7. Quelle commande vous permet de compter le nombre de données manquantes dans x?
- 8. Calculez la moyenne sur les 80 1ères valeurs

Summary on matrices

Format two-dimensions

Datatype class() to check it is a matrix

homogeneous: only one type of character, numeric, logical, factor

-> ceorcion if heterogeneous -> check with mode()

Creation matrix(), cbind(), rbind()

Adding new items cbind(), rbind()

Size length() -> nb of items

Dim dim(), str()

Slicing my_matrix [i,j]

Filling my_matrix[i,j] <- "toto"

Naming colnames(), rownames()

Let's play together!

Exercice sur les matrices

entrez vos solutions de code ou réponses (ou likez celle de vos collègues):

- 1. Créer un vecteur myVector avec 12 entiers de 1 à 12
- 2. Regarder la taille puis la dimension de myVector
- 3. Imposer des dim 3 et 4 au vecteur **myVector**
- 4. Quelle est à présent la classe de myVector
- 5. Ajouter une colonne avec un vecteur avec les chaînes de caractères: « one », « two » et « three »
- 6. Qu'est-il advenu aux valeurs numériques?
- 7. Que faire? Un dataframe
- 8. Renommer les colonnes de A à F avec une constante

3. dataframes:

Support pédagogique

Des diapos ci après pour mémo/archivage

Un tutoriel avec un jupyter notebook reprenant les exemples des diapos

« Rsession1_tuto_dataframes.ipynb » sur github

ou dans /shared/projects/dubii2021/trainers/module3/tutorials

Un practical dans un jupyter notebook avec des données omiques

« Rsession1_practicals_dataframes.ipynb » sur github

ou dans /shared/projects/dubii2021/trainers/module3/practicals

Dataframe

Dataframe = two-dimensional object that can be heterogeneous,

Create a dataframe with function data.frame()

Dataframe created with existing vectors

Create a dataframe with function data.frame()

```
> myDataf <- data.frame(weight, size, bmi)
```

```
> myDataf # it looks pretty much like the matrix myData2 weight size bmi
```

```
Fabien 60 1.75 19.59184
Pierre 72 1.80 22.22222
Sandrine 57 1.65 20.93664
Claire 90 1.90 24.93075
Bruno 95 1.74 31.37799
Delphine 72 1.91 19.73630
```

> class(myDataf)# but this is well a dataframe and not a matrix

03/03/2021

[1] "data.frame"

> str(myDataf) # this one is a homogeneous df with numeric vectors

'data.frame': 6 obs. of 3 variables:

\$ weight: num 60 72 57 90 95 72

\$ size: num 1.75 1.8 1.65 1.9 1.74 1.91

\$ bmi : num 19.6 22.2 20.9 24.9 31.4 ...

> dim(myDataf)

[1] 6 3

Important:

If vectors are character chains, use stringsAsFactors= FALSE to avoid their conversion into factors

in R>4, the default is now FALSE !!!

Creating an empty dataframe

```
creating an empty dataframe?
 > d <- data.frame()
 > d
 data frame with 0 columns and 0 rows
 > dim(d)
 [1] 0 0
```

BUT USELESS: impossible to fill!

Better way: converting a matrix in a dataframe with function as.data.frame()

```
> class(myData2)
                                    > d <- as.data.frame(matrix(NA,2,3))</pre>
[1] "matrix"
                                    > d
> class(as.data.frame(myData2))
                                     V1 V2 V3 # by default, col names are V1, V2, etc...
[1] "data.frame"
                                    1 NA NA NA # while if you are using the function
                                    2 NA NA NA # data.frame() and not as.dataframe(),
                                                 #col names are called X1, X2, etc...
                                    > dim(d)
                                    [1] 2 3
                                    > str(d)
                                    'data.frame': 2 obs. of 3 variables:
                                     $ V1: logi NA NA
                                     $ V2: logi NA NA
```

\$ V3: logi, NA NA

You may also use as.data.frame on a matrix generated by binding rows or columns

```
> d2 <- as.data.frame(cbind(1:2, 10:11)
> str(d2)
'data.frame': 2 obs. of 2 variables:
$ V1: int 12
$ V2: int 10 11
```

Row/Column names of dataframes

Either use same fonctions as for matrices rownames() and colnames()

Or better use the ones dedicated to dataframes: row.names() and names()

```
> row.names(d)
[1] "1" "2"
> names(d)
[1] "V1" "V2" "V3"
```

Important: each row name must be unique!

Note: data.frames are a special case of a list of variables of the same number of rows with unique row names

Extracting vectors from dataframes

Getting the vector corresponding to a column from a dataframe:

♦ either by specifying its index

```
> myDataf[,2]
[1] 1.75 1.80 1.65 1.90 1.74 1.91
```

♦ Or by giving its name within the " " inside the squared brackets

```
> myDataf[,"size"]
[1] 1.75 1.80 1.65 1.90 1.74 1.91
```

♦ Or by giving its name after the character « \$ »

```
> myDataf$size
[1] 1.75 1.80 1.65 1.90 1.74 1.91
```

Extracting rows from dataframes

Getting a « dataframe » corresponding to a row from a dataframe:

\$\times\$ either by specifying its index

```
> myDataf
          weight size
                          bmi
                                sex
Fabien
            60 1.75 19.59184
                                Man
Pierre
         72 1.80 22.22222
                                Man
Sandrine
        57 1.65 20.93664
                                Woman
Claire 90 1.90 24.93075
                                Woman
Bruno 95 1.74 31.37799
                                Man
Delphine
        72 1.91 19.73630
                                Woman
> myDataf[2,]
          weight size bmi sex
Pierre
          72 1.8 22.22222 Man
```

- If you extract only one row, you do not get a vector but a dataframe.
- => To convert into a vector, use unlist()

```
> temp <- unlist(myDataf["Pierre",])
> class(temp)
[1] "character"
```

- ♥ Or by giving its name within the " " inside the squared brackets

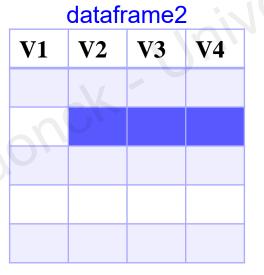
Let's summarize and give it a try

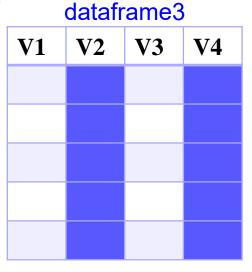
How do we create a dataframe?

Which are the three methods to slice datrames?

Which command should I use to extract the blue cells of the 3 dataframes below?

dataframe1						
V1	V2	V 3	V 4			
			~Q;			
		19				







Now, how to extract the even columns if I have 500 000 columns?

Adding new vectors to a dataframe

Either enter one vector at a time as a new variable

my_dataframe\$new_variable <- my_variable

Or **several vectors or subsets of dataframes** at once

♥ Using data.frame()

mynew dataframe <- data.frame(data.frame1, data.frame2)</pre>

This method will keep the data types of each data.frame

Using cbind() BUT to avoid ->prefer using data.frame()

mynew_dataframe <- cbind(data.frame1, data.frame2)

BE CAREFULL: this method will keep the data types only if the data.frames 1 and 2 have several variables.

If they have only one, these variable will be converted as a vector and cbind() will convert character strings as factors.

```
> d2$new <- 1:2
V1 V2 new
1 1 10 1
2 2 11 2</pre>
```

Reading a text file into R creates a dataframe

Read a text file using read.table():

```
temperatures <- read.table("temperatures.txt", sep="\t", header=T, stringsAsFactors=F)</p>
>temperatures
Month Mean Temp
                                  specify the field
   January
             2.0
                                   separator of the
   February 2.6
                                       text file
   March
          7.9
         11.2
   April
                                                        TRUE if
   May 15.3
   June 22.2
                                                       header in
                                                      the text file
   July 22.9
   August 22.5
                                                                                 FALSE to avoid
   September
                17.3
                                                                                 factorisation of
10 October
             11.7
                                                                                character vectors
11 November
                5.2
12 December
                2.8
> str(temperatures) # the R object is a dataframe !!!!
'data.frame': 12 obs. of 2 variables:
$ Month: chr "January" "February" "March" "April" ...
$ Mean Temp: num 2 2.6 7.9 11.2 15.3 22.2 22.9 22.5 17.3 11.7 ...
```

Reading a text file into R: to factorize or not to factorize?

Warning: use stringsAsFactors=F (now by default in R>4 but TRUE in previous versions) otherwise vectors of character values converted into factors -> see below, the Months were factorized!!!!

```
> temperatures <- read.table("Temperatures.txt", sep="\t", header=T, stringsAsFactors=T)
> str(temperatures)
'data.frame': 12 obs. of 2 variables:
$ Month : Factor w/ 12 levels "April", "August", ..: 5 4 8 1 9 7 6 2 12 11 ...
$ Mean_Temp: num 2 2.6 7.9 11.2 15.3 22.2 22.9 22.5 17.3 11.7 ...
> levels(temperatures$Month) # the levels of the factor are in alphabetic order
[1] "April" "August" "December" "February" "January" "July"
[7] "June"
            "March"
                        "May"
                                 "November" "October" "September"
                                                                 TRUE is by
                                                                default in R<4
```

Factors in R

See tutorial_Factors_in_R.html (on GitHub)

Much care on:

- levels order
- coercion

=> ∈ travail personnel de vendredi

Reading a text file into R

Caution if:

« readxl »

- fewer names than columns in the header
- fewer columns than names in the header -> add argument fill=T to overcome the issue
- some rows with fewer columns -> add argument fill = T to overcome the issue
- using row.names=1 -> this cannot be used when several rows have the same name

or read.xlsx() or read_excel() to read worksheet from an exel file with library « xlsx » or

```
Check the data.frame is as expected using:

str()

head(): displays the first 6 rows

tail(): displays the last 6 rows

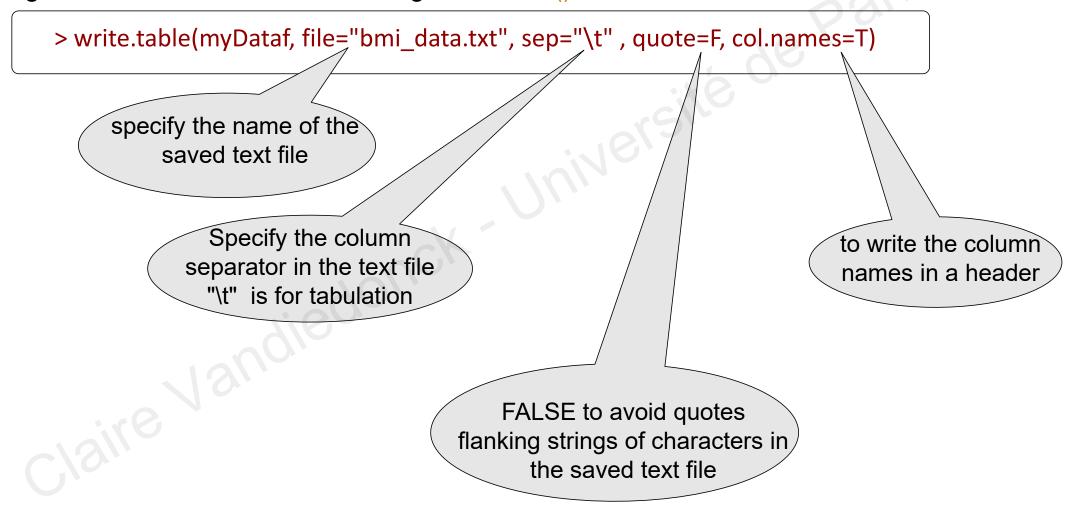
and by displaying some rows in the middle of the file using their index

-> a general habit with any programmation language

Other functions: read.csv(), scan()
```

Saving a dataframe as a text file in the working directory

Saving a dataframe into a text file using write.table()



Filtering dataframes on criteria with which()

It generates a new dataframe

use which() that returns the index of what is TRUE in the condition

```
> which ( myDataf$sex == "Woman")
[1] 3 4 6
> myDataf [ which ( myDataf$sex == "Woman") , ]
         weight size
                          bmi
                                sex
             57 1.65 20.93664 Woman
Sandrine
Claire 90 1.90 24.93075 Woman
Delphine 72 1.91 19.73630 Woman
> str(myDataf [ which ( myDataf$sex == "Woman") , ])
'data.frame': 3 obs. of 4 variables:
 $ weight: num 57 90 72
 $ size : num 1.65 1.9 1.91
  bmi : num 20.9 24.9 19.7
 $ sex
         : chr "Woman" "Woman" "Woman"
```

6^{**}Important:

you may enter this
without including
« which »
BUT this would not deal
with NA values
=> safer to use which()

Or what does not match using "!=" for "different" or "!" for "not" before the test

```
> which ( myDataf$sex != "Man")
[1] 3 4 6
> which (! myDataf$sex == "Man")
[1] 3 4 6
```

What happens if I am not using which()?

CAUTION! If you have NA values, not using which() will also return them!

Let's make a copy of our dataframe and replace the gender of Claire by a missing value

```
> myDataf2 <- myDataf
> myDataf2["Claire", "sex"] <- NA
> myDataf [ myDataf$sex == "Woman", ]
              weight size bmi
                                 sex
Sandrine
              57 1.65 20.93664 Woman
Claire
             90 1.90 24.93075 NA
Delphine 72 1.91 19.73630 Woman
> myDataf2 [ myDataf2$sex == "Woman", ]
              weight size
                                 bmi
                                        sex
Sandrine
              57 1.65 20.93664 Woman
NA
              NA
                    NA
                             NA
                                 \langle NA \rangle
              72 1.91 19.73630 Woman
Delphine
> myDataf2 [ which(myDataf2$sex == "Woman"), ]
              weight size bmi sex
Sandrine
              57 1.65 20.93664 Woman
Delphine
              72 1.91 19.73630 Woman
```



Filtering dataframes on patterns with grep()

\$\times\$ use grep() that returns the index of what matches (even partially)

```
> grep("Wom", myDataf$sex)
[1] 3 4 6
> grep("Woman", myDataf$sex)
[1] 3 4 6
> myDataf [grep("Woman", myDataf$sex), ]
           weight size bmi
                                sex
Sandrine
             57 1.65 20.93664 Woman
Claire 90 1.90 24.93075 Woman
Delphine 72 1.91 19.73630 Woman
> grep("a", row.names(myDataf)) # returns indexes of rows with an "a" in its name
[1] 1 3 4
> myDataf [grep("a", row.names(myDataf)),]
             weight size
                               bmi
                                      sex
Fabien
          60 1.75 19.59184 Man
Sandrine 57 1.65 20.93664 Woman
Claire
             90 1.90 24.93075 Woman
```

Filtering/Subsetting dataframes on criteria with subset()

Subsetting the rows on the columns:

\$\to\$ use subset(): the easiest and most efficient way!

```
> WomenDataf <- subset(myDataf, sex== "Woman")</pre>
```

> WomenDataf

```
      weight size
      bmi sex

      Sandrine
      57 1.65 20.93664
      Woman

      Claire
      90 1.90 24.93075
      Woman

      Delphine
      72 1.91 19.73630
      Woman
```

Filtering dataframes on several criteria

```
logical: & = and, | = or, ! = not
comparison: ==, != (different), >, < , >=, <=
is an element of a vector using : %in%</pre>
```

Either with which()

Or more easily with subset()

```
> subset( myDataf, sex == "Woman" & weight < 80 & bmi > 20)

weight size bmi sex

Sandrine 57 1.65 20.93664 Woman
```

Merging dataframes

Merge two dataframes with a key

- myDataf\$index <- 1:6</p>
- > myDataf

in this example I create a new column for the key but we can also use an existing variable

```
weight size
                                  index
                       bmi
                            sex
Fabien
            60 1.75 19.59184
                              Man
            72 1.80 22.22222
Pierre
                              Man
            57 1.65 20.93664 Woman
Sandrine
Claire
            90 1.90 24.93075 Woman
            95 1.74 31.37799
Bruno
                              Man
            72 1.91 19.73630 Woman
Delphine
```

- > OtherData <- data.frame(c(1:5, 7),rep(c("right-handed","left-handed"),3))
- > names(OtherData) <- c("ID","handedness")</pre>
- > OtherData
 - ID handedness
- 1 1 right-handed
- 2 2 left-handed
- 3 3 right-handed
- 4 4 left-handed
- 5 5 right-handed
- 6 7 left-handed

Merging dataframes

Merge two dataframes with a key

- > myDataf\$index <- 1:6
- > myDataf



> myDataf\$index <- 1:6 > myDataf weight size bmi sex index Fabien 60 1.75 19.59184 Man 1 Pierre 72 1.80 22.22222 Man 2 Sandrine 57 1.65 20.93664 Woman 3 Claire 90 1.90 24.93075 Woman 4 Bruno 95 1.74 31.37799 Man 5	ge two datafra	ames with a key				
> myDataf weight size bmi sex index Fabien 60 1.75 19.59184 Man 1 Pierre 72 1.80 22.22222 Man 2 Sandrine 57 1.65 20.93664 Woman 3 Claire 90 1.90 24.93075 Woman 4	> myDataf\$i	ndex <- 1:6			_	
Fabien 60 1.75 19.59184 Man 1 Pierre 72 1.80 22.22222 Man 2 Sandrine 57 1.65 20.93664 Woman 3 Claire 90 1.90 24.93075 Woman 4	•				•	
Pierre 72 1.80 22.22222 Man 2 Sandrine 57 1.65 20.93664 Woman 3 Claire 90 1.90 24.93075 Woman 4		weight size	bmi	sex	index	46,
Sandrine 57 1.65 20.93664 Woman 3 Claire 90 1.90 24.93075 Woman 4	Fabien	60 1.75	19.59184	Man	1	200
Claire 90 1.90 24.93075 Woman 4	Pierre	72 1.80	22.2222	Man	2	
	Sandrine	57 1.65	20.93664	Woman	3	
Bruno 95 1.74 31.37799 Man 5	Claire	90 1.90	24.93075	Woman	4	
	Bruno	95 1.74	31.37799	Man	5	
Delphine 72 1.91 19.73630 Woman 6	Delphine	72 1.91	19.73630	Woman	6	

- > OtherData <- data.frame(c(1:5, 7),rep(c("right-handed","left-handed"),3))
- > names(OtherData) <- c("ID","handedness")</pre>
- > OtherData



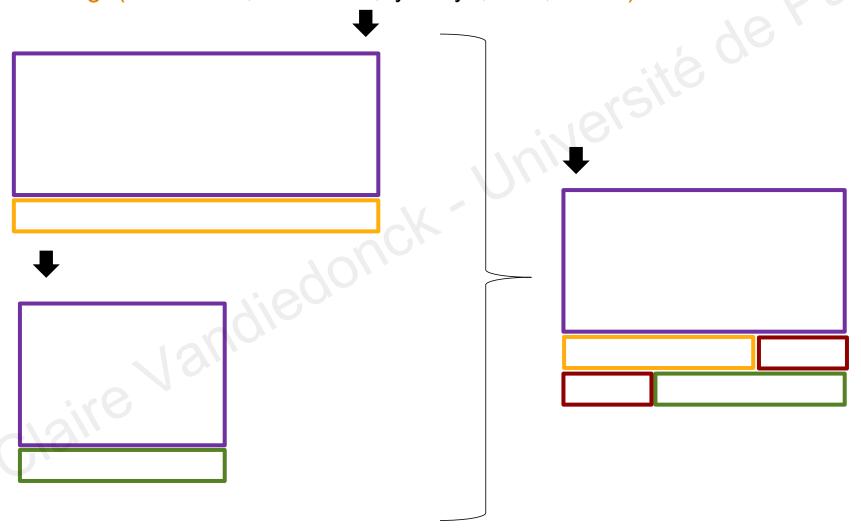
handedness

- 1 right-handed
- 2 left-handed
- 3 right-handed
- left-handed
- 5 5 right-handed
- left-handed 6

Merging dataframes

Merge two dataframes with a key

with merge(dataframe1,dataframe2,by="key" , all=T, sort=F)



Merged dataframe

Merge two dataframes with a key

```
> myMergedDataf <- merge(myDataf, OtherData, by.x="index", by.y="ID", all.x=T, all.y=T, sort=F)
```

> myMergedDataf



	index	weight	: size	e bm:	i sex	k handedness
1	1	60	1.75	19.59184	Man	right-handed
2	2	72	1.80	22.2222	Man	left-handed
3	3	57	1.65	20.93664	Woman	right-handed
4	4	90	1.90	24.93075	Woman	left-handed
5	5	95	1.74	31.37799	Man	right-handed
6	6	72	1.91	19.73630	Woman	<na></na>
7	7	NA	NA	NA	<na></na>	left-handed

- ✓ unless the merge is done on the row.names(), the row.names of initial data.frames are lost -> the new data.frame has its own row names
- ✓ if two columns had the same name, a « .x » or a « .y » is added to the first/second.

Summary on dataframes

Format two-dimensions

Datatype class() to check it is a dataframe

hteterogeneous: type of columns/variable can differ

Creation data.frame() with existing vectors, as.data.frame() when

converting a matrix

Adding new items my_dataf\$new_var to add a new column, rbind() to add a new row

AVOID using cbind() to add a columns -> use data.frame() instead

Dim dim(), str(), nrows(), ncol()

Slicing my_dataf[i,j] by indexes, my_dataf["my_row","my_var"] by name,

my_dataf\$my_var for a given columns

Filling my_dataf[i,j] <- "toto", etc...using slicing methods

Naming names(), row.names()