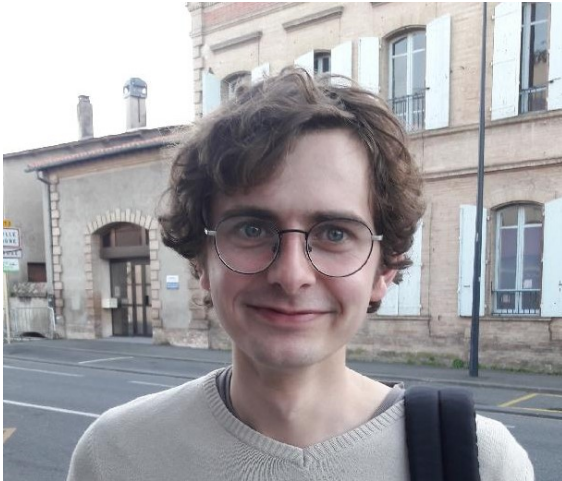


Data-wrangling on R

Rémi Mahmoud

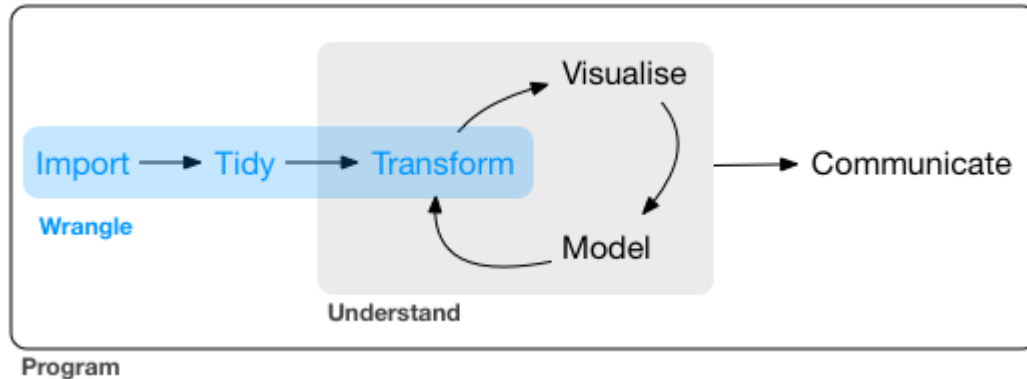
Who is talking ?



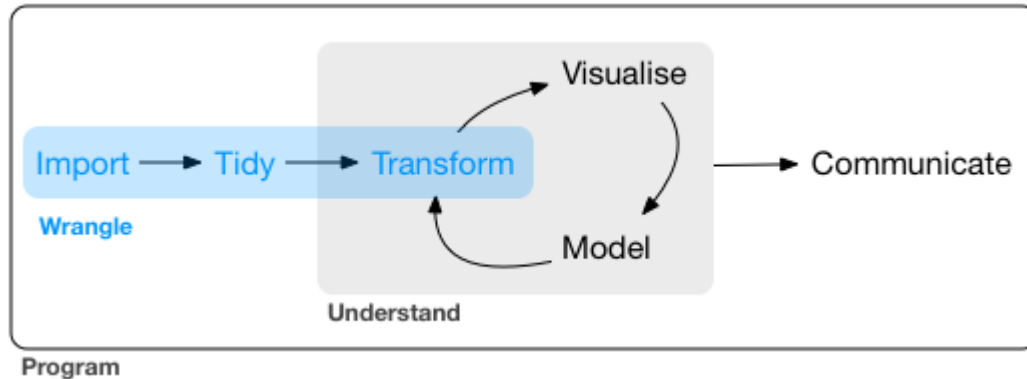
INRAE

What is data-wrangling ?

What is data-wrangling ?



What is data-wrangling ?



- data-wrangling is the set of operations on **raw** data that leads to **non messy** (tidy) data.

What we will talk about today

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- Data importation

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- Manipulate data (filtering, arranging data etc.)

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What we will NOT talk about today

What we will talk about today

- Data importation
- Manipulate data (filtering, arranging data etc.)
- Tidy data

What we will NOT talk about today

- Dealing with missing values / outliers

Framework

All manipulations will be done in the `tidyverse` framework.

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Hence, you should, if not already done, run the following command in R **NOW**

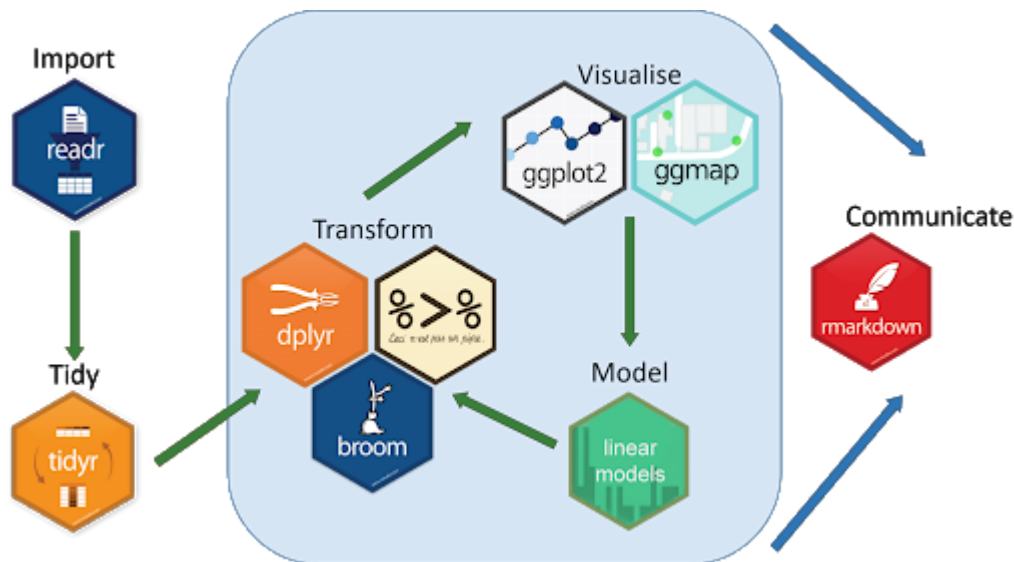
```
install.packages("tidyverse")
```

Tidyverse = Tidy universe

Tidyverse is a set of packages with different purposes, that share the same syntax and that are designed to work in a complementary way

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You can list the packages available in the tidyverse by running the following command:

```
tidyverse::tidyverse_packages()
```

```
## [1] "broom"      "cli"        "crayon"     "dbplyr"     "dplyr"
## [6] "forcats"    "ggplot2"    "haven"      "hms"        "httr"
## [11] "jsonlite"   "lubridate"  "magrittr"   "modelr"     "pillar"
## [16] "purrr"      "readr"      "readxl"     "reprex"     "rlang"
## [21] "rstudioapi" "rvest"      "stringr"    "tibble"     "tidyr"
## [26] "xml2"       "tidyverse"
```


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```

You can see that `ggplot2` that you discovered yesterday belongs to the tidyverse. But there are *many other packages* !

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You can see that `ggplot2` that you discovered yesterday belongs to the tidyverse. But there are *many other packages* !

For instance, the `forcats` package allows to work in a convenient way with factors, `lubridate` with dates etc. .

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```

You can see that `ggplot2` that you discovered yesterday belongs to the tidyverse. But there are *many other packages* !

For instance, the `forcats` package allows to work in a convenient way with factors, `lubridate` with dates etc. .

For now, we will take a closer look to the `readr` and to a lesser extent `readxl` packages. These packages are useful to **import** data.

Import data with readr

The `read_csv` function of `readr` allows to read csv files.

Import data with readr

The `read_csv` function of `readr` allows to read csv files.

```
data_work <- readr::read_csv('data/iris.csv')
```

```
## Parsed with column specification:
## cols(
##   Sepal.Length = col_double(),
##   Sepal.Width = col_double(),
##   Petal.Length = col_double(),
##   Petal.Width = col_double(),
##   Species = col_character()
## )
```

`read_csv` is faster than base R `read.csv` and it parses well different types of columns.

This function has also other arguments that may be useful for you when using it:

This function has also other arguments that may be useful for you when using it:

- `skip` to specify the number of lines to skip before reading the file
- `na` to specify what should be considered as NA (for ex: you could put `na = "Not answered"`)
- `col_names` to specify the names of the columns you want to have in your dataset.
- See `?read_csv` for other arguments.

Tibble vs dataframe

Let us take a look at the data we have imported.

```
head(data_work)
```

```
## # A tibble: 6 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <chr>
## 1         5.1         3.5         1.4         0.2 setosa
## 2         4.9         3         1.4         0.2 setosa
## 3         4.7         3.2         1.3         0.2 setosa
## 4         4.6         3.1         1.5         0.2 setosa
## 5         5         3.6         1.4         0.2 setosa
## 6         5.4         3.9         1.7         0.4 setosa
```


Tibble vs dataframe

Let us take a look at the data we have imported.

```
head(data_work)
```

```
## # A tibble: 6 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <chr>
## 1         5.1         3.5         1.4         0.2 setosa
## 2         4.9         3         1.4         0.2 setosa
## 3         4.7         3.2         1.3         0.2 setosa
## 4         4.6         3.1         1.5         0.2 setosa
## 5         5         3.6         1.4         0.2 setosa
## 6         5.4         3.9         1.7         0.4 setosa
```

Note the particular output of the print:

- *A tibble*
- The type of each column is written under each col_name

The tibble is an alternative to the classical `data.frame` of base R

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As part of the tidyverse, it is mainly used in the tidyverse' packages.

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The difference should not worry you: the main difference with a classical dataframe is the nicer output when printing (run `iris` in R to see the difference).

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As part of the tidyverse, it is mainly used in the tidyverse' packages.

The difference should not worry you: the main difference with a classical dataframe is the nicer output when printing (run `iris` in R to see the difference).

By the way, note that a tibble **is a `data.frame`**

```
is.data.frame(data_work)
```

```
## [1] TRUE
```

An other package of the tidyverse: `readxl` to read `.xlsx` files

The function `read_xlsx` allows you to read `.xlsx` files.

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The function `read_xlsx` allows you to read `.xlsx` files.

Some arguments are useful for you:

- `sheet` : name of the sheet of the file you want to read (if you provide a string), or position of the sheet you want to read (if you provide an integer)
- same arguments as `read_csv` (`na`, `skip` etc.)
- see `?read_xlsx` for details.

Is everything ok until now ?



Exercises

~ 30 minutes

Exercises

~ 30 minutes

*Download the zip Github depository, open
"Import_data_exercises.Rmd" with Rstudio*

Manipulate your data using `dplyr`

`dplyr` is a package of the `tidyverse` designed to manipulate your data easily.

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■ what do we mean by manipulating the data easily ?

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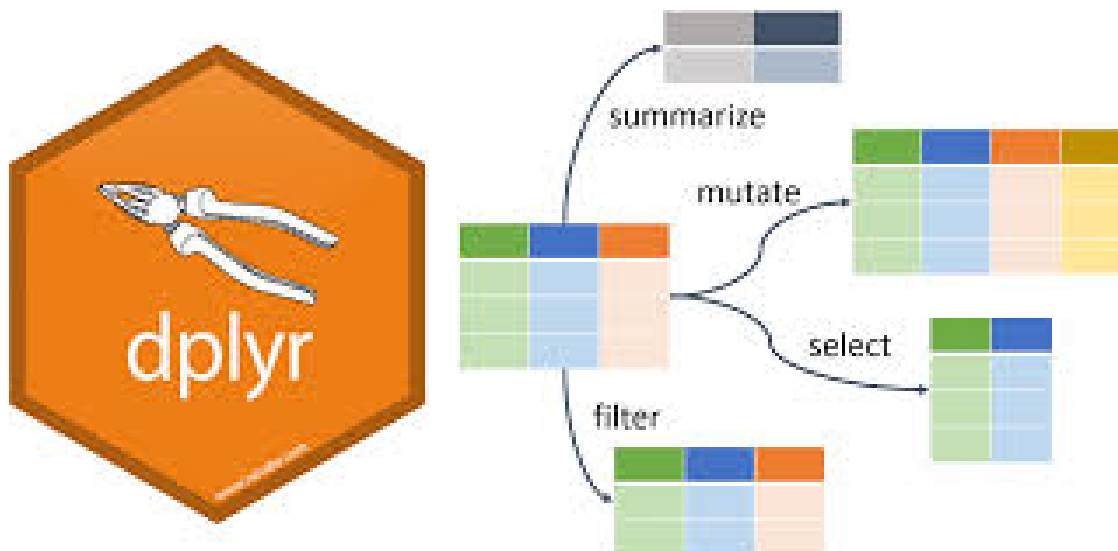
Select columns, filter their rows, create new columns etc.

Manipulate your data using dplyr

dplyr is a package of the tidyverse designed to manipulate your data easily.

what do we mean by manipulating the data easily ?

Select columns, filter their rows, create new columns etc.



Let us consider the dataset `data_work` previously introduced (it is simply the well know `iris` dataset turned into tibble).

```
## # A tibble: 150 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <chr>
## 1         5.1         3.5         1.4         0.2 setosa
## 2         4.9         3         1.4         0.2 setosa
## 3         4.7         3.2         1.3         0.2 setosa
## 4         4.6         3.1         1.5         0.2 setosa
## 5         5         3.6         1.4         0.2 setosa
## 6         5.4         3.9         1.7         0.4 setosa
## # ... with 144 more rows
```

Let us see the manipulations we can do on this dataset.

`dplyr::select`

In a data analysis, we could be interested in:

- *select* some columns, for instance:
 - *select* the 3rd column

dplyr::select

In a data analysis, we could be interested in:

- *select* some columns, for instance:
 - *select* the 3rd column

```
select(data_work, 3)
```

```
## # A tibble: 150 x 1
##   Petal.Length
##         <dbl>
## 1           1.4
## 2           1.4
## 3           1.3
## 4           1.5
## 5           1.4
## 6           1.7
## # ... with 144 more rows
```

`dplyr::select`

In a data analysis, we could be interested in:

- *select* some columns, for instance:
 - *select* column Sepal.Width

dplyr::select

In a data analysis, we could be interested in:

- *select* some columns, for instance:
 - *select* column Sepal.Width

```
select(data_work, Sepal.Width)  
# Note the absence of " around Sepal .Width
```

```
## # A tibble: 150 x 1  
##   Sepal.Width  
##         <dbl>  
## 1         3.5  
## 2          3  
## 3         3.2  
## 4         3.1  
## 5         3.6  
## 6         3.9  
## # ... with 144 more rows
```

`dplyr::select`

In a data analysis, we could be interested in:

- *select* some columns, for instance:
 - *select* all columns except Sepal.width and Sepal.Length)

dpplyr::select

In a data analysis, we could be interested in:

- *select* some columns, for instance:
 - *select* all columns except Sepal.width and Sepal.Length)

```
select(data_work, - c(Sepal.Width, Sepal.Length))
```

```
## # A tibble: 150 x 3
##   Petal.Length Petal.Width Species
##         <dbl>         <dbl> <chr>
## 1         1.4         0.2 setosa
## 2         1.4         0.2 setosa
## 3         1.3         0.2 setosa
## 4         1.5         0.2 setosa
## 5         1.4         0.2 setosa
## 6         1.7         0.4 setosa
## # ... with 144 more rows
```

dpplyr::select

In a data analysis, we could be interested in:

- *select* some columns, for instance:
 - *select* all columns except Sepal.width and Sepal.Length)

```
select(data_work, - c(Sepal.Width, Sepal.Length))
```

```
## # A tibble: 150 x 3
##   Petal.Length Petal.Width Species
##         <dbl>         <dbl> <chr>
## 1         1.4         0.2 setosa
## 2         1.4         0.2 setosa
## 3         1.3         0.2 setosa
## 4         1.5         0.2 setosa
## 5         1.4         0.2 setosa
## 6         1.7         0.4 setosa
## # ... with 144 more rows
```

Note the absence of " around Sepal.Width and Sepal.Length, and the - that means **except**

`dplyr::select:helpers()`

`select()` is provided with many *functions helpers* that you can use to select columns, for instance:

- `select(data_work, contains("pal"))`: all columns of `data_work` with a name containing "pal"

`dplyr::select:helpers()`

`select()` is provided with many *functions helpers* that you can use to select columns, for instance:

- `select(data_work, contains("pal"))`: all columns of `data_work` with a name containing "pal"
- `select(data_work, starts_with("Se"))`: *can you guess it ?*

dp<code>lyr::select:helpers()

`select()` is provided with many *functions helpers* that you can use to select columns, for instance:

- `select(data_work, contains("pal"))`: all columns of `data_work` with a name containing "pal"
- `select(data_work, starts_with("Se"))`: *can you guess it ?*
- `select(data_work, ends_with("th"))`: *can you guess it ?*

`dp_lyr::select:helpers()`

`select()` is provided with many *functions helpers* that you can use to select columns, for instance:

- `select(data_work, contains("pal"))`: all columns of `data_work` with a name containing "pal"
- `select(data_work, starts_with("Se"))`: *can you guess it ?*
- `select(data_work, ends_with("th"))`: *can you guess it ?*
- `select(data_work, matches("*th"))`: *can you guess it ?* (select columns with a name matching a regular expression)

`dplyr::select:helpers()`

`select()` is provided with many *functions helpers* that you can use to select columns, for instance:

- `select(data_work, contains("pal"))`: all columns of `data_work` with a name containing "pal"
- `select(data_work, starts_with("Se"))`: *can you guess it ?*
- `select(data_work, ends_with("th"))`: *can you guess it ?*
- `select(data_work, matches("*th"))`: *can you guess it ?* (select columns with a name matching a regular expression)

That's one of the assets of the `dplyr` syntax: it looks like almost natural language.

`dplyr::filter`

In a data analysis, we could be interested in:

- *filter* rows based on the values of some columns (predicates), for instance:
 - *filter* rows of `data_work` with individuals having their length of Sepal greater than 4

dplyr::filter

In a data analysis, we could be interested in:

- *filter* rows based on the values of some columns (predicates), for instance:
 - *filter* rows of `data_work` with individuals having their length of Sepal greater than 4

```
filter(data_work, Sepal.Length > 4)
```

```
## # A tibble: 150 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##   <dbl>        <dbl>        <dbl>        <dbl> <chr>
## 1         5.1         3.5         1.4         0.2 setosa
## 2         4.9         3         1.4         0.2 setosa
## 3         4.7         3.2         1.3         0.2 setosa
## 4         4.6         3.1         1.5         0.2 setosa
## 5         5         3.6         1.4         0.2 setosa
## 6         5.4         3.9         1.7         0.4 setosa
## # ... with 144 more rows
```

`dplyr::filter`

In a data analysis, we could be interested in:

- *filter* rows based on the values of some columns (predicates), for instance:
 - *filter* rows of `data_work` of species "virginica"

dplyr::filter

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- *filter* rows based on the values of some columns (predicates), for instance:
 - *filter* rows of data_work of species "virginica"

```
filter(data_work, Species == "virginica")
```

```
## # A tibble: 50 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <chr>
## 1         6.3           3.3           6           2.5 virginica
## 2         5.8           2.7           5.1          1.9 virginica
## 3         7.1           3            5.9          2.1 virginica
## 4         6.3           2.9           5.6          1.8 virginica
## 5         6.5           3            5.8          2.2 virginica
## 6         7.6           3            6.6          2.1 virginica
## # ... with 44 more rows
```

`dplyr::filter`

You can put multiple conditions, for instance:

- *filter* rows based on the values of some columns (predicates), for instance:
 - *filter* rows of `data_work` of species "virginica" and with their Width of Petal smaller than 2

dplyr::filter

You can put multiple conditions, for instance:

- *filter* rows based on the values of some columns (predicates), for instance:
 - *filter* rows of `data_work` of species "virginica" and with their Width of Petal smaller than 2

```
filter(data_work, Species == "virginica", Petal.Width < 2)
```

```
## # A tibble: 21 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <chr>
## 1         5.8           2.7           5.1           1.9 virginica
## 2         6.3           2.9           5.6           1.8 virginica
## 3         4.9           2.5           4.5           1.7 virginica
## 4         7.3           2.9           6.3           1.8 virginica
## 5         6.7           2.5           5.8           1.8 virginica
## 6         6.4           2.7           5.3           1.9 virginica
## # ... with 15 more rows
```

| Again, it looks like the natural language !

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That's one of the nicer things in the `dplyr` syntax.

`dplyr::mutate`

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`mutate` is the verb used to create new columns.

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For instance, suppose we want to compute the sum of the lengths of the Sepal and the Petal in our dataset.

dpplyr::mutate

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For instance, suppose we want to compute the sum of the lengths of the Sepal and the Petal in our dataset.

```
mutate(data_work, sum_lengths = Sepal.Length + Petal.Length)
```

```
## # A tibble: 150 x 6
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species sum_lengths
##         <dbl>      <dbl>      <dbl>      <dbl>    <chr>      <dbl>
## 1         5.1         3.5         1.4         0.2  setosa         8.6
## 2         4.9         3         1.4         0.2  setosa         7.9
## 3         4.7         3.2         1.3         0.2  setosa         8.2
## 4         4.6         3.1         1.5         0.2  setosa         8.4
## 5         5         3.6         1.4         0.2  setosa         9.2
## 6         5.4         3.9         1.7         0.4  setosa         9.4
## # ... with 144 more rows
```

dp1yr: other useful functions

dp1yr provides many useful functions. You can guess their purposes just by their name:

dplyr: other useful functions

dplyr provides many useful functions. You can guess their purposes just by their name:

- `arrange`: `arrange(data_work, Species, desc(Petal.Length))`

dplyr: other useful functions

dplyr provides many useful functions. You can guess their purposes just by their name:

- `arrange`: `arrange(data_work, Species, desc(Petal.Length))`

```
## # A tibble: 150 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <chr>
## 1           4.8           3.4           1.9           0.2 setosa
## 2           5.1           3.8           1.9           0.4 setosa
## 3           5.4           3.9           1.7           0.4 setosa
## 4           5.7           3.8           1.7           0.3 setosa
## 5           5.4           3.4           1.7           0.2 setosa
## 6           5.1           3.3           1.7           0.5 setosa
## # ... with 144 more rows
```

dp1yr: other useful functions

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dplyr: other useful functions

dplyr provides many useful functions. You can guess their purposes just by their name:

- `distinct`: `distinct(data_work, Species)`

dplyr: other useful functions

dplyr provides many useful functions. You can guess their purposes just by their name:

- `distinct: distinct(data_work, Species)`

```
## # A tibble: 3 x 1
##   Species
##   <chr>
## 1 setosa
## 2 versicolor
## 3 virginica
```

dp1yr: other useful functions

dp1yr provides many useful functions. You can guess their purposes just by their name:

dplyr: other useful functions

dplyr provides many useful functions. You can guess their purposes just by their name:

- `rename`: `rename(data_work, S.Width = Sepal.Width, S.Length = Sepal.Length)`

dplyr: other useful functions

dplyr provides many useful functions. You can guess their purposes just by their name:

- `rename`: `rename(data_work, S.Width = Sepal.Width, S.Length = Sepal.Length)`

```
## # A tibble: 150 x 5
##   S.Length S.Width Petal.Length Petal.Width Species
##   <dbl>   <dbl>         <dbl>         <dbl> <chr>
## 1     5.1     3.5           1.4           0.2 setosa
## 2     4.9     3             1.4           0.2 setosa
## 3     4.7     3.2           1.3           0.2 setosa
## 4     4.6     3.1           1.5           0.2 setosa
## 5     5       3.6           1.4           0.2 setosa
## 6     5.4     3.9           1.7           0.4 setosa
## # ... with 144 more rows
```


Is everything clear ?





Break Time



~ 15 minutes

Chain commands using %>% (pipe) operator

The %>% (pronounce pipe) provides a convenient way to code, as it allows the code to be written in chain.



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IMPORTANT: the keyboard shortcut for %>% is *ctrl + shift + M*

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The %>% (pronounce pipe) provides a convenient way to code, as it allows the code to be written in chain.



IMPORTANT: the keyboard shortcut for %>% is *ctrl + shift + M*

Try it !

For instance, suppose we want to:

1. *filter* rows of `data_work` of species "virginica" and with their Width of Petal smaller than 2
2. *then* compute the sum of the lengths of the Sepal and the Petal in our dataset.
3. *then* select the columns with their name starting with an *S*
4. *then* arrange the result by length of `Sepal.Length`

We would write

For instance, suppose we want to:

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2. *then* compute the sum of the lengths of the Sepal and the Petal in our dataset.
3. *then* select the columns with their name starting with an *S*
4. *then* arrange the result by length of `Sepal.Length`

We would write

```
arrange(select(mutate(filter(data_work, Species == 'virginica', Petal
```


For instance, suppose we want to:

1. *filter* rows of `data_work` of species "virginica" and with their Width of Petal smaller than 2
2. *then* compute the sum of the lengths of the Sepal and the Petal in our dataset.
3. *then* select the columns with their name starting with an *S*
4. *then* arrange the result by length of `Sepal.Length`

We would write

```
arrange(select(mutate(filter(data_work, Species == 'virginica', Petal
```

Isn't it unreadable ?!

Let's write it using the `%>%` operator:

Let's write it using the %>% operator:

```
data_work %>%  
  filter(Species == 'virginica', Petal.Width < 2) %>%  
  mutate(Sum_lengths = Sepal.Length + Petal.Length) %>%  
  select(starts_with("S")) %>%  
  arrange(Sepal.Length)
```

Let's write it using the %>% operator:

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You see how clearer it looks ?

If I run this: `x %>% sum` it is strictly equivalent to `sum(x)`.

It means: take `x` and pass it through the function `sum`.

Another example to see the power of $\%>\%$. Suppose I want to carry out the following steps:

Another example to see the power of %>%. Suppose I want to carry out the following steps:

1. Take `data_work`
2. Select variables containing "Sepal", and "Petal.Width" and "Species"
3. Filter rows with length of Sepal greater than 5
4. Fit a linear model of Petal.Width vs Sepal.Width + Sepal.Length + Species
5. Print a summary of the model

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4. Fit a linear model of Petal.Width vs Sepal.Width + Sepal.Length + Species
5. Print a summary of the model

```
data_work %>% #Step 1
  select(contains("Sepal") ,
         Petal.Width, Species) %>% # Step 2
  filter(Sepal.Length >5) %>% # Step 3
  lm(Petal.Width ~ Sepal.Width + Sepal.Length + Species,
     data= .) %>% # Step 4: NOTE THE .
  summary # Step 5
```

```
##
## Call:
## lm(formula = Petal.Width ~ Sepal.Width + Sepal.Length + Species,
##     data = .)
##
## Residuals:
```

	Min	1Q	Median	3Q	Max
##	-0.48660	-0.10718	-0.00351	0.12237	0.46503

```
##
## Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
## (Intercept)	-1.14888	0.24851	-4.623	1.01e-05 ***

```
data_work %>% #Step 1
  select(contains("Sepal") ,
         Petal.Width, Species) %>% # Step 2
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In this example, it is also important to notice the `.`. When using the pipe, the `"."` is the object referring to what's before the last `%>%`.

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```

In this example, it is also important to notice the `.`. When using the pipe, the `"."` is the object referring to what's before the last `%>%`.

It is important to specify it when the argument that needs the object before the last `%>%` is not the first argument. That's why we had to specify it in the `lm` function and not in the `select` function.

Group operations

An important features of `dplyr` is its ability to *group* tibbles and compute operations on these *grouped* tibbles.

Group operations

An important features of `dplyr` is its ability to *group* tibbles and compute operations on these *grouped* tibbles.

A *key* function of `dplyr` is `group_by`.

`dplyr::group_by`

dplyr::group_by

```
data_work_by_species <- data_work %>%  
  group_by(Species)  
# Equivalent to data_work_by_species <- group_by(data_work, Species)  
  
data_work_by_species
```

```
## # A tibble: 150 x 5  
## # Groups:   Species [3]  
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species  
##           <dbl>         <dbl>         <dbl>         <dbl> <chr>  
## 1           5.1           3.5           1.4           0.2 setosa  
## 2           4.9           3           1.4           0.2 setosa  
## 3           4.7           3.2           1.3           0.2 setosa  
## 4           4.6           3.1           1.5           0.2 setosa  
## 5           5           3.6           1.4           0.2 setosa  
## 6           5.4           3.9           1.7           0.4 setosa  
## # ... with 144 more rows
```


dplyr::group_by

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data_work_by_species <- data_work %>%  
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# Equivalent to data_work_by_species <- group_by(data_work, Species)  
  
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##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species  
##           <dbl>         <dbl>         <dbl>         <dbl> <chr>  
## 1           5.1           3.5           1.4           0.2 setosa  
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## 5           5           3.6           1.4           0.2 setosa  
## 6           5.4           3.9           1.7           0.4 setosa  
## # ... with 144 more rows
```

Note the # Groups: Species [3]. It means that operations on this dataset will be done for each group.

For example, suppose we want to compute the median of the width of the Sepal for each species.

```
data_work_by_species %>%  
  mutate(median_sepal_width = median(Sepal.Width)) %>%  
  select(starts_with("S"), median_sepal_width)
```

```
## # A tibble: 150 x 4  
## # Groups:   Species [3]  
##   Sepal.Length Sepal.Width Species median_sepal_width  
##           <dbl>         <dbl> <chr>           <dbl>  
## 1           5.1           3.5 setosa           3.4  
## 2           4.9           3   setosa           3.4  
## 3           4.7           3.2 setosa           3.4  
## 4           4.6           3.1 setosa           3.4  
## 5           5             3.6 setosa           3.4  
## 6           5.4           3.9 setosa           3.4  
## # ... with 144 more rows
```

It's nice, but we may also need to summarise the table, just keep a summary of the Species and the median.

```
dplyr::summarise
```

It is easily done by the function `summarise`

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```
data_work_by_species %>%  
  summarise(median_sepal_width = median(Sepal.Width))
```

```
## # A tibble: 3 x 2  
##   Species    median_sepal_width  
##   <chr>          <dbl>  
## 1 setosa          3.4  
## 2 versicolor     2.8  
## 3 virginica       3
```

If you want to take out the grouped structure of your tibble, you just have to use the function `ungroup`

```
data_work_by_species %>% ungroup
```

```
## # A tibble: 150 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <chr>
## 1           5.1           3.5           1.4           0.2 setosa
## 2           4.9           3           1.4           0.2 setosa
## 3           4.7           3.2           1.3           0.2 setosa
## 4           4.6           3.1           1.5           0.2 setosa
## 5           5           3.6           1.4           0.2 setosa
## 6           5.4           3.9           1.7           0.4 setosa
## # ... with 144 more rows
```

Do you have questions ?



Exercises

~ 20 minutes

Exercises

~ 20 minutes

*Open "Manipulate_and_tidy_your_data_exercises.Rmd" with
Rstudio*

Exercises

~ 20 minutes

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Rstudio*

Answer to the questions until section "Tidy your data"

Tidy data

1. Each variable must have its own column
2. Each observation must have its row
3. Each value must have its own cell

country	year	cases	population
Afghanistan	1999	1745	19987071
Afghanistan	2000	2666	20595360
Brazil	1999	30737	172006362
Brazil	2000	80488	174504898
China	1999	214258	1272415272
China	2000	210766	1280425583

variables

country	year	cases	population
Afghanistan	1999	1745	19987071
Afghanistan	2000	2666	20595360
Brazil	1999	30737	172006362
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China	1999	214258	1272415272
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observations

country	year	cases	population
Afghanistan	99	745	19987071
Afghanistan	00	2666	20595360
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Brazil	00	80488	174504898
China	99	214258	1272415272
China	00	210766	1280425583

values

For instance, imagine this dataset, giving the population of different countries in 2002 and 2007:

```
d %>% head
```

```
## # A tibble: 6 x 3
##   country      2002      2007
##   <chr>      <dbl>      <dbl>
## 1 Belgium    10311970  10392226
## 2 France     59925035  61083916
## 3 Germany    82350671  82400996
## 4 Italy      57926999  58147733
## 5 Spain      40152517  40448191
## 6 Switzerland 7361757   7554661
```

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##   country      2002      2007
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- Is this dataset tidy ?

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```
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```

- Is this dataset tidy ?

This dataset is **not** tidy, as the population, which is an observed variable is not in a distinct column (principle 1.). Year is also a variable, so it should have its column too.

Instead, we should have:

```
## # A tibble: 6 x 3
##   country year  population
##   <chr>   <chr>      <dbl>
## 1 Belgium 2002      10311970
## 2 Belgium 2007      10392226
## 3 France  2002      59925035
## 4 France  2007      61083916
## 5 Germany 2002      82350671
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```

Make tidy data

Two key functions, of package `tidyr` are used to tidy the data:

Make tidy data

Two key functions, of package `tidyr` are used to tidy the data:

- `tidyr::pivot_longer` is used to make your dataset *longer* (what a surprise ! :O)
- `tidyr::pivot_wider` is used to make your dataset *wider* (what a surprise ! :O)

In practice:

```
#d is the dataset with the populations of the countries
data_tidy <- d %>% pivot_longer(cols = c("2002", "2007"))

head(data_tidy)
```

```
## # A tibble: 6 x 3
##   country name      value
##   <chr>    <chr>    <dbl>
## 1 Belgium 2002    10311970
## 2 Belgium 2007    10392226
## 3 France  2002    59925035
## 4 France  2007    61083916
## 5 Germany 2002    82350671
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The first argument is the dataset to tidy (which is not necessary to complete because of the %>%).

In practice:

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#d is the dataset with the populations of the countries
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The first argument is the dataset to tidy (which is not necessary to complete because of the %>%).

The second is the name of the columns to gather.

We can also provide another names to the new columns using the arguments `values_to` and `names_to`.

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The columns to gather can also be selected using the select helpers that we've seen previously:

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The columns to gather can also be selected using the select helpers that we've seen previously:

```
#d is the dataset with the populations of the countries
data_tidy <- d %>%
  pivot_longer(cols = -country,
               names_to = "Year", # New name
               #for the columns 2002 and 2007
               values_to = "Population")
#New name for the values

head(data_tidy)
```

```
## # A tibble: 6 x 3
##   country Year   Population
##   <chr>   <chr>         <dbl>
## 1 Belgium 2002     10311970
## 2 Belgium 2007     10392226
## 3 France  2002     59925035
## 4 France  2007     61083916
## 5 Germany 2002     82350671
## 6 Germany 2007     82400996
```

Conversely, `pivot_wider` allows us to come back to the first dataset.

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```
data_tidy %>% pivot_wider(names_from = Year, values_from = Population)
```

```
## # A tibble: 6 x 3
##   country      2002      2007
##   <chr>      <dbl>      <dbl>
## 1 Belgium    10311970  10392226
## 2 France     59925035  61083916
## 3 Germany    82350671  82400996
## 4 Italy      57926999  58147733
## 5 Spain      40152517  40448191
## 6 Switzerland 7361757   7554661
```

Is everything clear ?



Exercises

Exercises

~ 20 minutes

Exercises

~ 20 minutes

Answer to the questions of the section "Tidy your data"

Conclusion

The tidyverse provides many tools to work with data.

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Many topics have not been presented today:

- manipulate factors using `forcats`
- manipulate dates using `lubridate`
- manipulate dates using `stringr`
- join tables using the `*_join` functions of `dplyr`
- apply `mutate`, `filter` or `select` on specific columns using `*_at`, `*_if`, `*_all` suffixes
- ...

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The tidyverse provides many tools to work with data.

Many topics have not been presented today:

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- join tables using the `*_join` functions of `dplyr`
- apply `mutate`, `filter` or `select` on specific columns using `*_at`, `*_if`, `*_all` suffixes
- ...

Feel free to consult this book (available for free online at this adress:

<https://r4ds.had.co.nz/>):

THANKS

THANKS

Any remark, questions ?

THANKS

Any remark, questions ?
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THANKS

Any remark, questions ?

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Lesson contents available at:

https://github.com/RemiMahmoud/data_wrangling