



REGIONAL TRAINING ON CAPACITY DEVELOPMENT OF DATA ANALYTICS AND DISSEMINATION USING "R" SOFTWARE

AMMAN, JORDAN, 3 - 7 DECEMBER, 2023





Outline

- Wrap-up
- Import data to R
- Exploring imported data
- Data processing part I
 - mutate
 - Select
 - Filter
 - Working with variable names
 - clean_names
 - rename
 - recode
 - De-duplication
- Q&A



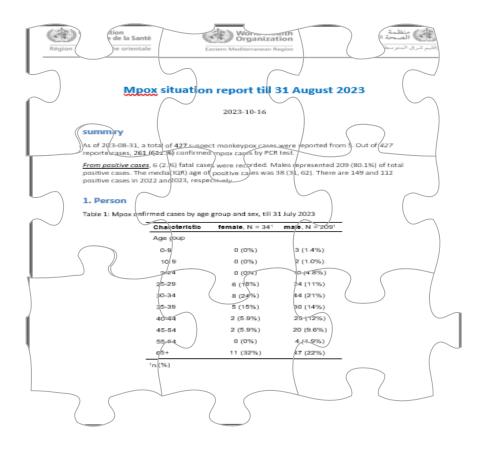


Desired output at end of the training





Desired output at end of the training







Data importation to R





Session 2 Agenda

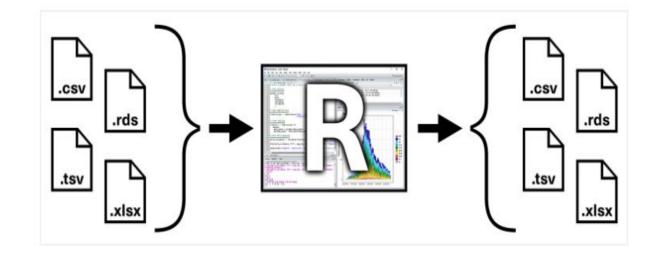
- 9:00 9: 30 (30 min): **Wrap-up**
- 9:30 9:50 (20 min): Presentation "Data importation to R"
- 9:50 10:20 (30 min): **Demonstration**
- 10:20 10:40 (20 min): **Stretching / coffee break**
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Importing data to R

- Different file formats can be imported to R:
 - Spreadsheets (most common): .csv, .xls, .xlsx
 - SPSS dataset: .sav
 - Stata dataset: .dta
 - SAS dataset: .sas7sdat
 - SQL databases
 - JSON files



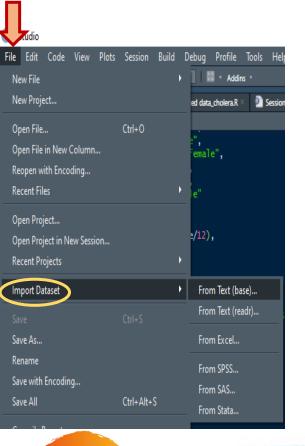




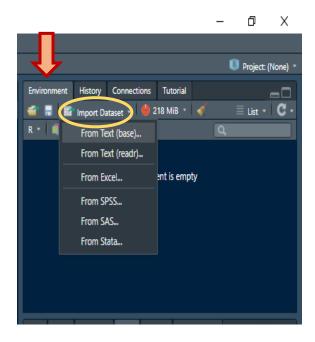
How to import data to R?

R built-in options What is the file format?!!

Installed packages



and action



Where is that file?!!

Many packages are available to support importing different datasets to R. We will use some of:

- Specific for .csv
 - readr::read_csv()
- Many file formats
 - rio::import()
 - data.table::fread





How to import data to R? Cont.





Where is that file?!!

Absolute path

- A specific path for the file that is unique to the user's computer
- Against the concept of reproducible analysis

C:/Users/abdelgawadb/Desktop/Regional R training workshop/Data/Cholera case study/cholera_20231102.csv'

Relative path

- Fixing the file path to the root of R project
- Best for reproducibility
- here::here() is used

import(here("Data", "Cholera case study", "cholera_20231102.csv"))



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Data checking

Initial checking of the imported dataset (data frame):

- Ensure that data is correctly imported
- Explore for any missing, duplicates, inconsistencies

```
> dim(cholera)
[1] 108 53
```

> nrow(cholera) [1] 108 > ncol(cholera) [1] 53

and action

Dimensions of the data frame (rows, columns)

Separately, number of rows/observations Number of columns/variables

```
> str(cholera)
                108 obs. of 53 variables:
'data.frame':
                      "cholera-001" "cholera-003" "cholera-005" "cholera-007" ...
$ recordx id
                      "Place 1" "Place 2" "Place 1" "Place 2" ...
$ hospital
                       "female" "male" "female" "female" ...
$ sex
 $ age
                       "year" "year" "year" "month" ...
 $ Age.unit
                                    "2023-07-07" "2023-07-17" "2023-07-24" ...
 $ adm.date
 $ temp
                      NA NA 36.4 36.9 37.1 36.8 36.2 NA 37 NA ...
 $ cons
 $ cap_fill
$ pulse
$ wt.kg.
$ ht..cm.
                      162 167 93 156 109 147 121 105 163 142 ...
                       NA NA NA NA NA ...
 $ muac
```

Display the structure of the data frame and its variables



Data checking cont.

Checking numeric and date variables

```
summary(cholera$age)
  Min. 1st Qu.
                Median
                           Mean 3rd Qu.
                                            Max.
                          16.59
           7.00
                   9.00
                                  24.00
                                           56.00
> summary(cholera$onset_date)
                                Median
                  1st Qu.
                                                Mean
                                                                                         NA's
                                                          3rd Qu.
                                                                           Max.
"2021-05-06" "2022-02-11" "2022-12-12" "2022-09-12" "2023-07-02" "2023-07-29"
                                                                                          "4"
```

Checking any variable

```
> table(cholera$f_diag, useNA = "always")

confirmed not a case probable suspect <NA>
24 4 74 6 0
```

- Other useful functions:
 - skimr::skim()
 - class()
 - head()/tail()
 - duplicated()
 - is.na()
 - Other simple mathematical functions





Data checking cont.

The concept of tidy data



TIDY DATA is a standard way of mapping the meaning of a dataset to its structure. • •

-HADLEY WICKHAM

In tidy data:

- each variable forms a column
- each observation forms a row
- each cell is a single measurement

each column a variable							
	id	name	color				
	1	floof	gray	K			
	2	max	black	each row an			
	3	cat	orange	bobservation			
	4	donut	gray	2//			
	5	merlin	black	4/			
	6	panda	calico	7			

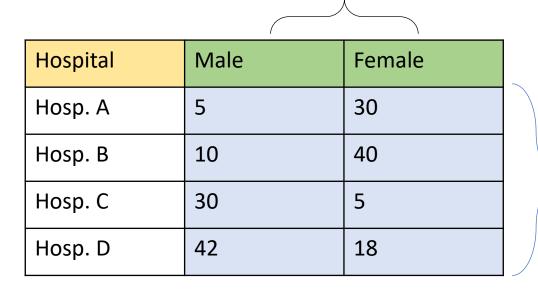
Wickham, H. (2014). Tidy Data. Journal of Statistical Software 59 (10). DOI: 10.18637/jss.v059.i10





Data checking cont.

The concept of tidy data





Hospital	Gender	Value
Hosp. A	Male	5
Hosp. A	Female	30
Hosp. B	Male	10
Hosp. B	Female	40
Hosp. C	Male	30
Hosp. C	Female	5
Hosp. D	Male	42
Hosp. D	Female	18





Demonstration





Exercise: Importing and Exploring Cholera Data

- Open your training R project "Regional_R_training"
- Create a new R script "cholera.R" and save it to the scripts folder
- Structure the script with sections (e.g about the script, load packages,....)
- **Load** the required packages
- **Import** cholera case study "cholera_20231102.csv" dataset located in the "Data" folder, use one of the following:
 - import + here functions
 - fread function

- Make sure the relevant packages are installed/loaded!
- Do not forget to assign the imported data to an object "cholera0"

- **Explore** the imported dataset:
 - What are the number of observations and variables?
 - What is the class() of the imported data?
 - Summary() of the imported data
 - What is the class of "admission date" and "outcome date" variables?





Exercise cont...

- Open your training R project
- Create a new R script "cholera.R" and save to scripts folder
- Import cholera case study "cholera_20231102.csv" dataset
- **Explore** the imported dataset:
 - How many duplicates are in the imported dataset?
 - What is the latest symptom onset date?
 - What is the range and the mean of reported cases' age? [how would you read it?!!]
 - What is the distribution of sex among reported cases? [HINT: table()]
- + Bonus!!
- What are the names of the reported countries? [HINT: unique()]
- Cross-tabulate sex "sex" with final diagnosis "f_diag" [HINT: table()]





Data processing – part l





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Data processing

Data steps to prepare the dataset for analysis:

1. On dataset level:

Proposed function

- Drop/keep columns - - → select()
- De-duplicate ----→ distinct(), duplicated()
- Filter specific observations - → filter()

2. On variable level:

- Variable Name ----- clean_names(), rename()
- Manage inconsistencies --- → recode()
- Change class of variable --- as.Date(), as.numeric(),...
- Create a new variable ---- mutate(), case_when(), if_else()
- Modify same variable ---- mutate(), case_when(), if_else()







Commonly used function

Package::function	Utility		
%>%	Arithmetic operators		
dplyr::filter()	Subset rows		
dplyr::select()	Subset columns		
dplyr::mutate()	Create & transform columns		
<pre>lubridate::mdy(), dmy(), ymd()</pre>	Tell R how to understand names		
janitor::clean_names	Standardize names		
dplyr::rename()	Manual renaming		
dplyr::case_when()	Complex logical re-coding of values		
ifelse()	Simple logical re-coding of values		
dplyr::recode()	Re-code values in a column		
as.Date(), as.numeric()	Convert the class of a column		





Demonstration





Exercise: Data Wrangling in R: Cholera Case Study Processing

- Open your training R project "Regional_R_training"
- Open the R script "cholera.R"
- Add a new section "data processing 1" and do the following data steps: (create a new object "cholera")
 - 1. Standardize the column names [HINT: clean_names()]

✓ It is clear now why column naming is important!

- 2. Rename "f_diag" to a more meaningful name "case_cat"
- 3. Fix inconsistencies in "sex" entries to male & female only [HINT: recode()]
- 4. Create a new column "age_cat" in years:
 - *First:* create new column "age_yr" to standardize all age units to be in years [HINT: mutate() & case_when()]
 - Second: now, create the "age_cat" column [HINT: cut() or epikit::age_categories()]
- 5. Drop all unnecessary columns (e.g. blank ones)
- + Bonus!!
- 6. Create a body mass index "bmi" column as weight(kg)/height(m²)









