

A semi-automated procedure to clean and enrich electronic monitoring adherence data The CleanADHdata.R script

ESPACOMP Adherence Data Analysis workshop November 17th, 2022

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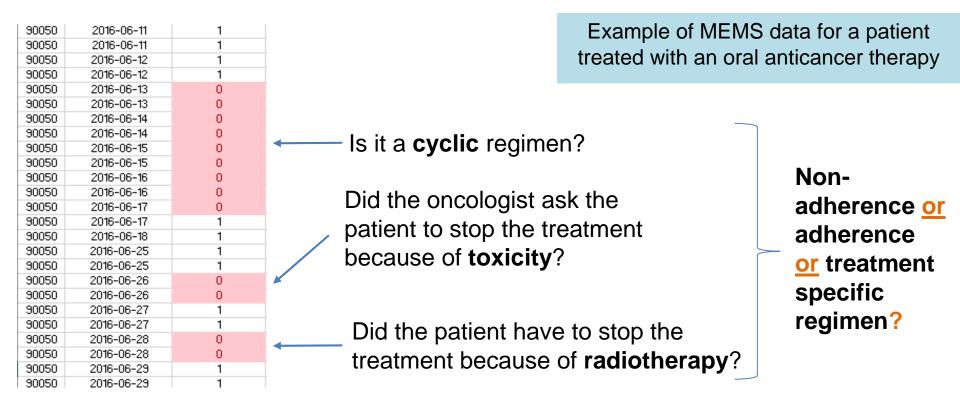






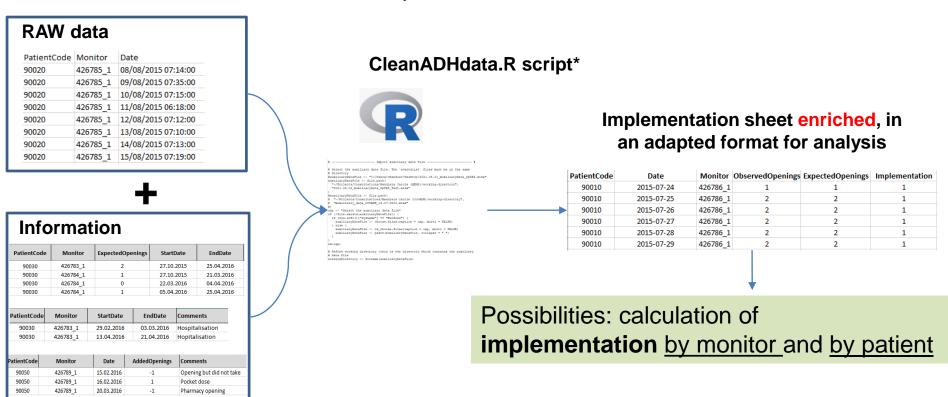
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Context: need to clean electronic monitoring (EM) adherence database

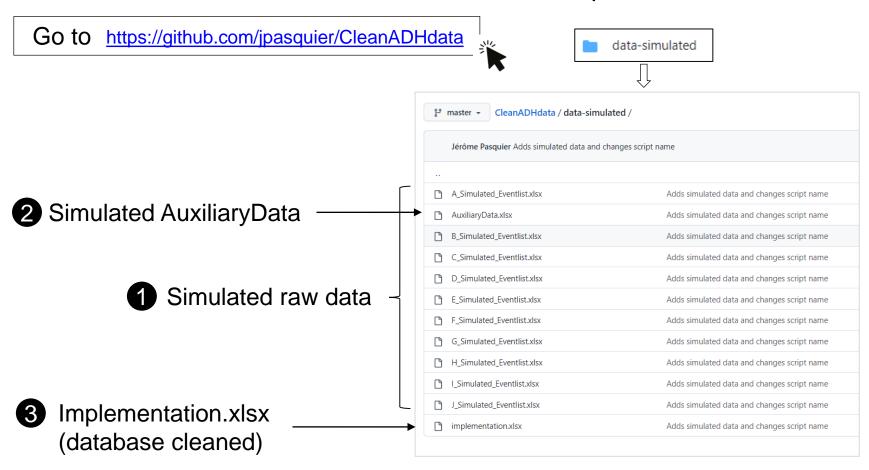


Aim: developing a script in the statistical software R to clean and enrich MEMS data with information provided upon patient's report and medical record

Methods: CleanADHdata.R script cleans MEMS data



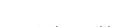
^{*}Implementation at each day=1 if the patient opens the MEMS <u>at least</u> the number of times it is expected, otherwise implementation=0



1. Extraction of the EM raw data

From the MEDAMIGO software AARDEX

From an Excel sheet in the required format

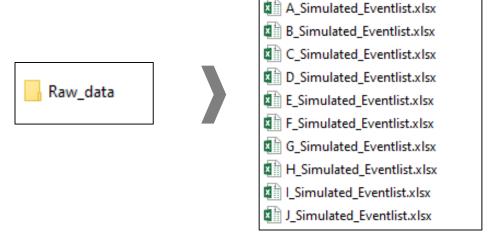


The three columns are to be entitled "PatientCode", "Monitor" and "Date" in the first row

PatientCode	Monitor	Date
Α	125A	2020-10-05 07:50:01 UTC
Α	125A	2020-10-05 14:54:55 UTC
Α	125A	2020-10-06 18:09:57 UTC

2. Gathering the patients' EM raw data into the same file folder

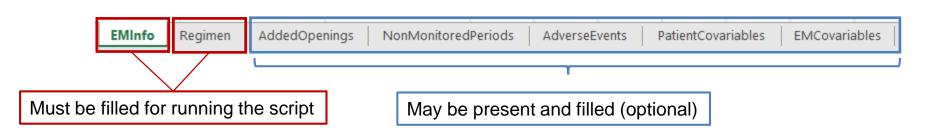
Filenames have to contain the "eventslist" or "eventlist" character string

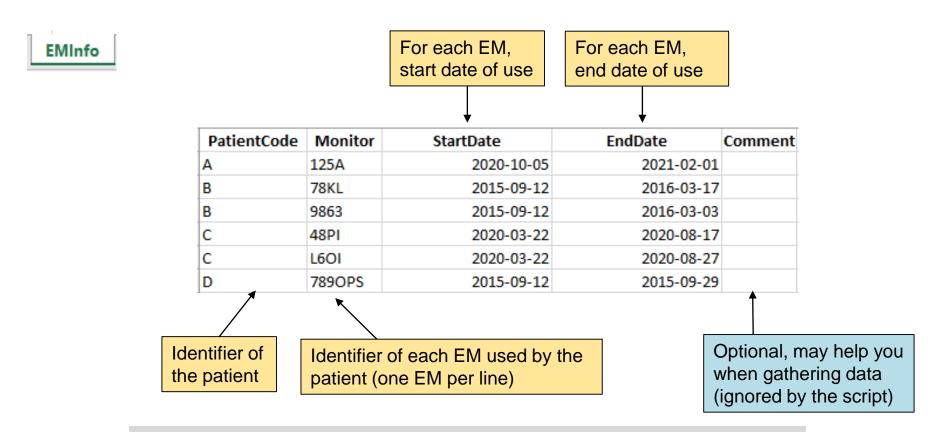


3. Preparing the **Auxiliary data** Excel sheet

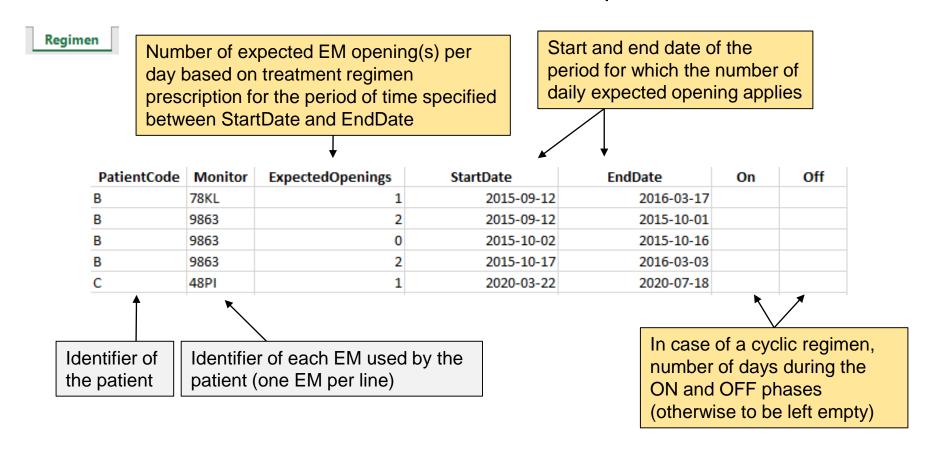
Investigators need to determine in advance which relevant variables to collect (e.g., regimen changes, hospitalization dates, periods of EM non-use)

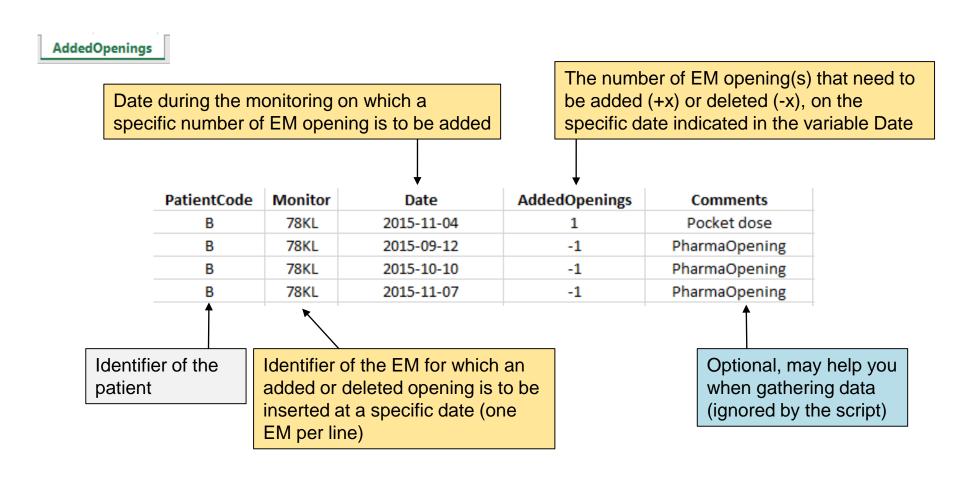
Composition of the Auxiliary Data Excel Sheet: 7 different sheets

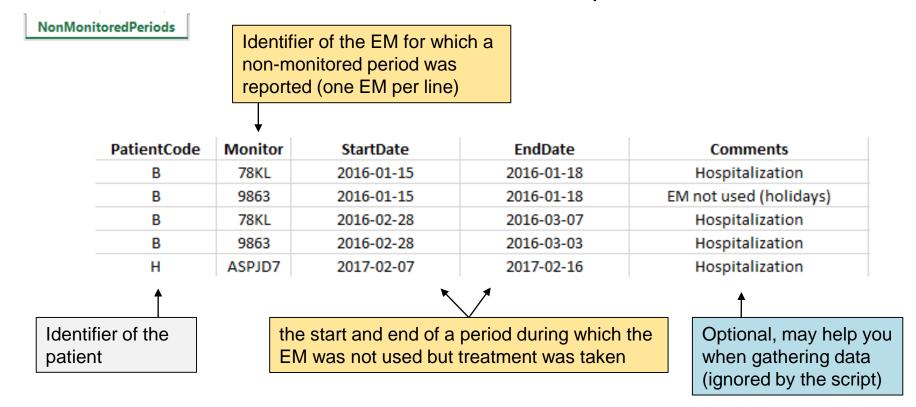




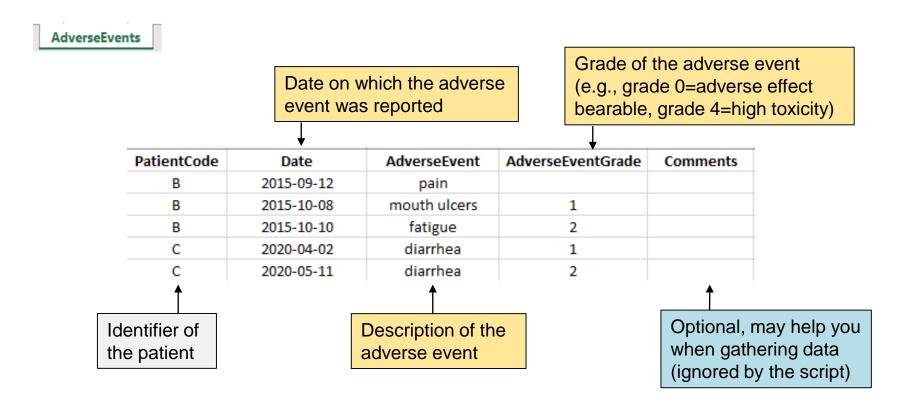
The cleaned database is truncated by the **start and end dates** of each EM use.







The implementation component of adherence is not calculated during nonmonitored periods and has to be considered as missing during the analyses.



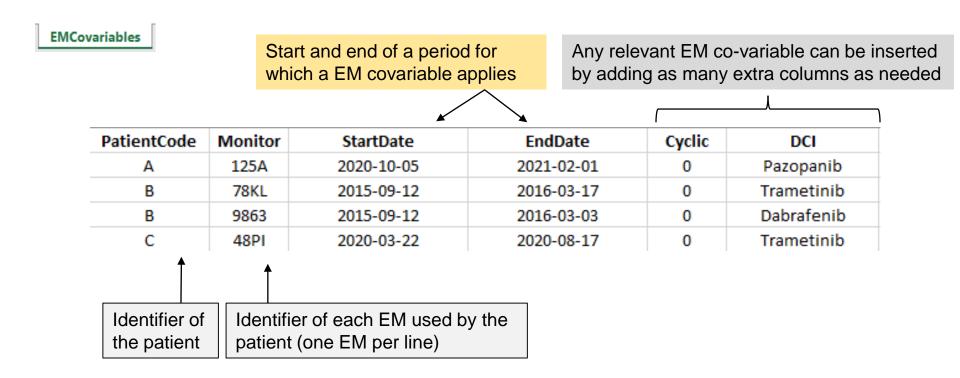
PatientCovariables

Any relevant patient co-variable can be inserted by adding as many extra columns as needed

PatientCode	StartDate	EndDate	Gender	Age	Discontinuation_censure
Α	2020-10-05	2021-02-01	1	65	0
В	2015-09-12	2016-03-17	2	48	0
С	2020-03-22	2020-08-27	2	61	1

Identifier of the patient

start and end of a period for which a patient covariable applies

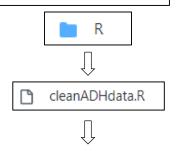


4. Running the CleanADHData.R script (available on https://github.com/jpasquier/CleanADHdata/tree/master/R)

➤ Open R



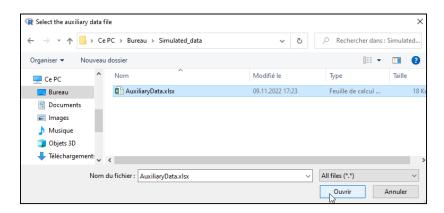
> Run the script

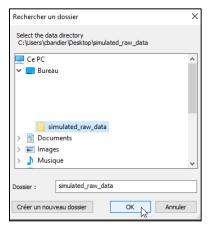


```
6 # TODO:
8 # POSSIBLE OPTIONS:
 9 # * Also read CSV files for raw data (in addition to Excel files))
10 # * Case-insensitive variable names
14 # Install missing packages and load required libraries
15 pkgs <- c("readxl", "writexl")</pre>
npkgs <- pkgs[!(pkgs %in% installed.packages()[, "Package"])]</pre>
17 if(length(npkgs)) {
18 install.packages(npkgs)
19 }
20 for (pkg in pkgs) {
21 library(pkg, character.only = TRUE)
23 if (Sys.info()["sysname"] != "Windows") {
24 if (!("tcltk" %in% installed.packages()[, "Package"])) {
       install.packages("tcltk")
27
     library(tcltk)
28 }
29 rm(pkgs, npkgs, pkg)
```

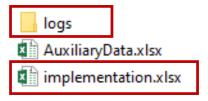
i) A window allows to select the AuxiliaryData Excel file

ii) A second window opens up to select the folder that contains the EM raw data

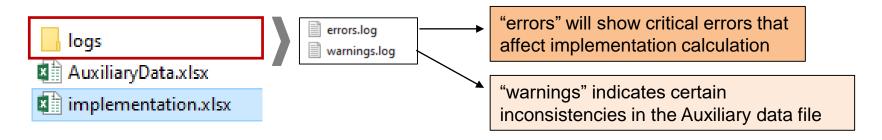




- 5. After the end of processing, the R script generates:
- ➤ the cleaned and enriched EM dataset, called "implementation.xlsx" in the same file folder as the AuxiliaryData Excel sheet
- Logs file

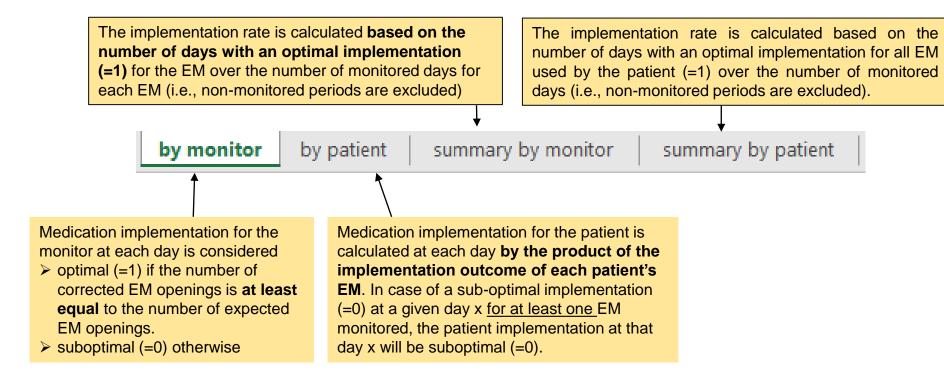


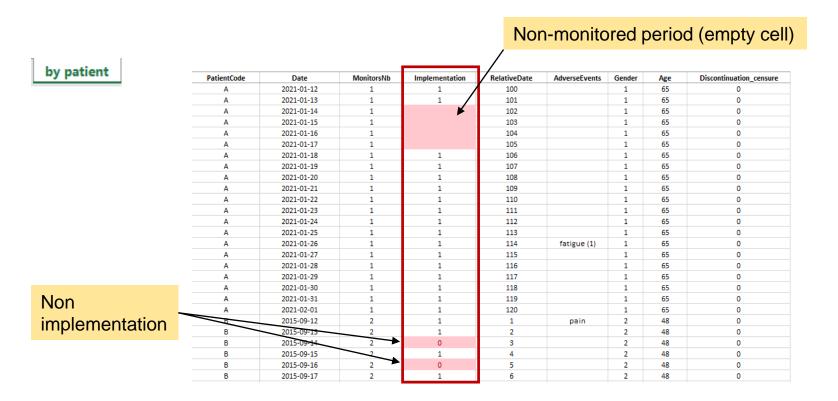
- 6. Checking errors with the log files
- ➤ 2 log files are created, located in the same file folder as the AuxiliaryData Excel sheet



After all errors or warnings are resolved, implementation.xlsx is ready for analysis

7. Implementation.xlsx





Cleaned EM database ready for analysis

summary by monitor

A first overview of the results before analysis

PatientCode	Monitor	Implementation
Α	125A	0.99
С	48PI	0.97
F	7374WOS	0.97
E	74RPZ	0.98
D	789OPS	1.00
В	78KL	0.86
В	9863	0.97
Н	ASPJD7	0.96
G	AU7	0.98
Н	J878D	0.93
С	L6OI	0.99
J	OJ5	0.99
I	S7	0.97
I	SOD7	0.97
J	SZU12	0.98



Patient B implemented less Trametinib (monitor 78KL) compared to Dabrafenib (monitor 9863)

summary by patient

A first overview of the results before analysis

Discussion

- Based on the cleaned and enriched dataset we can evaluate implementation, persistence and adherence analysis (e.g., with Generalized Estimating Equations), including covariates
- The calculation of medication implementation can be adapted directly in the script under the section "implementation calculation" (line 736)

```
# ----- Implementation calculation ----- #
```

- A tutorial will be submitted for publication to guide future users of the script
- Would you use the CleanADHdata.R script?
- Would you have any inputs?