

# {cleanepi

0.0.2

Stage of  
development:  
Experimental

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MRC Unit  
The  
Gambia

LONDON  
SCHOOL of  
HYGIENE  
& TROPICAL  
MEDICINE



***clean, curate and standardize  
epidemiological data***

<https://epiverse-trace.github.io/cleanepi>

 [epiverse-trace/cleanepi](https://github.com/epiverse-trace/cleanepi)

# functionalities

- `scan_data()`
- `standardize_date()`
- `check_subject_ids()`
- `calculate_age()`
- `check_date_sequence()`
- `find_duplicates()`
- `remove_duplicates()`
- `convert_to_numeric()`
- `Clean_using_dictionary()`
- `clean_data()`



# scan\_data(data = test\_data)

study_id	event	ccode	country	Date admitted	D.O.B	pcr_date	sex
PS001P2	day 0	2	Gambia	01/12/2020	06/01/1972	Dec 01, 2020	1
PS002P2	day 0	2	Gambia	28/01/2021	02/20/1952	Jan 01, 2021	1
PS004P2-1	day 0	2	Gambia	15/02/2021	06/15/1961	Feb 11, 2021	-99
PS003P2	day 0	2	Gambia	11/02/2021	11/11/1947	Feb 01, 2021	1
P0005P2	day 0	2	Gambia	17/02/2021	09/26/2000	Feb 16, 2021	2
PS006P2	day 0	2	Gambia	17/02/2021	-99	May 02, 2021	2



determine datatypes

- numeric, character and date
- % missing in every column

data_type	study_id	event	ccode	country	Date admitted	D.O.B	pcr_date	sex
missing	0	0	0	0	0	0.0	0	0
numeric	0	0	1	0	0	0.1	0	1
date	0	0	0	0	1	0.9	1	0
character	1	1	0	1	0	0.0	0	0
logical	0	0	0	0	0	0.0	0	0



# standardize\_date()

- convert date columns into %Y-%m-%d format
- convert character columns into Date



Date admitted	D.O.B	Pcr date
01/12/2020	06/01/1972	"Dec 01, 2020"
28/01/2021	02/20/1952	"Jan 01, 2021"
15/02/2021	06/15/1961	"Feb 11, 2021"
11/02/2021	11/11/1947	"Feb 01, 2021"
17/02/2021	09/26/2000	"Feb 16, 2021"
17/02/2021	-99	"May 02, 2021"

Date_admitted	D_O_B	Pcr_date
2020-12-01	01-06-1972	2020-12-01
2021-01-28	1952-02-20	2021-01-01
2021-02-15	1961-06-15	2021-02-11
2021-02-11	1947-11-11	2021-02-01
2021-02-17	2000-09-26	2021-02-16
2021-02-17	NA	2021-05-02

# check\_subject\_ids()

detect and remove incorrect subject IDs



study_id
PS001P2
PS002P2
PS004P2-1
PS003P2
P0005P2
PB500P2

study_id
PS001P2
PS002P2
PS004P2-1
PS003P2
P0005P2
PB500P2

# calculate\_age()

in years / months / weeks / days



study_id	DOB
PS001P2	1-Jun-1972
PS002P2	20-Feb-1952
PS004P2	06/15/1961
PS003P2	11/11/1947
P0005P2	26/09/2000
PS006P2	-99

study_id	DOB	age_months
PS001P2	1972-06-01	615
PS002P2	1952-02-20	859
PS004P2	1961-06-15	747
PS003P2	1947-11-11	910
P0005P2	2000-09-26	276
PS006P2	NA	NA



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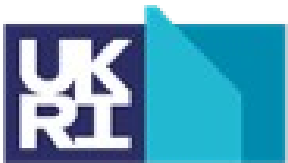
# check\_date\_sequence()

DOB < date\_admitted



Date admitted	D.O.B
06/01/1972	01/12/2020
28/01/2021	02/20/1952
15/02/2021	06/15/1961
11/02/2021	11/11/1947
09/26/2000	17/02/2021
17/02/2021	13/03/2022

DOB	date_admitted
01/12/2020	06/01/1972
02/20/1952	28/01/2021
06/15/1961	15/02/2021
11/11/1947	11/02/2021
17/02/2021	09/26/2000
13/03/2022	17/02/2021



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# find\_duplicates()



group_id	dt_onset	dt_report	sex	outcome
1	2015-05-21	2015-06-03	M	Alive
3	2015-05-31	2015-06-02	M	Dead
2	2015-05-30	2015-06-06	M	Alive
1	2015-05-21	2015-06-03	M	Alive
2	2015-05-30	2015-06-06	M	Alive



# remove\_duplicates()

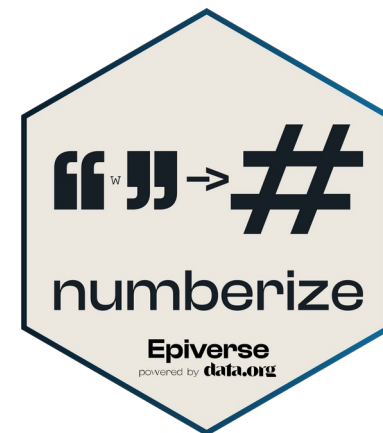


group_id	dt_onset	dt_report	sex	outcome
1	2015-05-21	2015-06-03	M	Alive
3	2015-05-31	2015-06-02	M	Dead
2	2015-05-30	2015-06-06	M	Alive

# convert\_to\_numeric()

```
x <- c(2, "ten", NA, 2, "twenty", 6)
```

```
x <- c(2, 10, NA, 2, 20, 6)
```



# clean\_using\_dictionary()

# THE DATA DICTIONARY

options	values	grp	orders
1	male	sex	1
2	female	sex	2

Add\_to\_dictionary()

options	values	grp	orders
1	male	sex	1
2	female	sex	2
-99	unknown	sex	3

sex
1
1
-99
1
2
2

sex
male
female
unknown
male
female
female



# clean\_data()

{

- scan\_data()
- standardize\_date()
- check\_subject\_ids()
- calculate\_age()
- check\_date\_sequence()
- find\_duplicates()
- remove\_duplicates()
- Convert\_to\_numeric()

}



contributors



Reviewers: Hugo, Pratik, Andree

Colleagues in the Statistics and Bioinformatics, Data management and App development teams