

# Package ‘hybridEHR’

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**Type** Package

**Title** Synthetic Hybrid Electronic Health Records Dataset Generator  
with COVID/CT Research Views

**Version** 0.1.0

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**Description** Tools to generate synthetic electronic health records including patients, encounters, vitals, labs, medications, procedures, and allergies, with optional COVID-19-focused and computed tomography (CT)-research views, and export them to comma separated values ('CSV'), 'SQLite', and 'Excel' formats for researchers and developers.

**License** MIT + file LICENSE

**Encoding** UTF-8

**Depends** R (>= 4.1.0)

**Imports** dplyr, tidyr, tibble, lubridate, jsonlite, openxlsx, DBI,  
RSQLite, magrittr

**Suggests** knitr, rmarkdown

**RoxygenNote** 7.3.3

**VignetteBuilder** knitr

**NeedsCompilation** no

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**Repository** CRAN

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**export\_hybrid\_ehr\_dataset**

*Export a hybrid EHR dataset to disk*

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**Description**

Export a hybrid EHR dataset to disk

**Usage**

```
export_hybrid_ehr_dataset(dataset, output_dir, verbose = TRUE)
```

**Arguments**

- |            |   |
|------------|---|
| dataset    | A list as returned by <a href="#">generate_hybrid_ehr_dataset()</a> . |
| output_dir | Directory to write files into.  |
| verbose    | Logical; if TRUE, print messages.                                     |

**Value**

The output directory (invisibly).

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**generate\_hybrid\_ehr**

*High-level wrapper to generate and export a hybrid EHR dataset*

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**Description**

High-level wrapper to generate and export a hybrid EHR dataset

**Usage**

```
generate_hybrid_ehr(  
  n_patients = 500,  
  n_sites = 3,  
  covid.Focused = TRUE,  
  include_ct_links = FALSE,  
  output_dir,  
  seed = NULL,  
  verbose = TRUE  
)
```

## Arguments

n_patients	Number of unique patients.
n_sites	Number of sites/hospitals to simulate.
covid.Focused	Logical; if TRUE, use COVID-era encounter and lab patterns.
include_ct_links	Logical; if TRUE, add CT timing variables and a CT severity score in the CT research view.
output_dir	Directory for exported files.
seed	Optional integer used to set the random seed for reproducibility.
verbose	Logical; if TRUE, print progress messages to the console.

## Value

A list with:

**dataset** The in-memory dataset list (as from generate\_hybrid\_ehr\_dataset).

**output\_dir** The output directory path where files were written.

A list containing:

**dataset** Generated dataset object

**output\_dir** Path to exported files

## Examples

```
ehr <- generate_hybrid_ehr_dataset(  
  n_patients = 10,  
  seed = 123,  
  verbose = FALSE  
)  
  
export_hybrid_ehr_dataset(  
  ehr,  
  output_dir = tempdir(),  
  verbose = FALSE  
)
```

**generate\_hybrid\_ehr\_dataset***Generate synthetic hybrid EHR tables*

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**Description**

Generate synthetic hybrid EHR tables

**Usage**

```
generate_hybrid_ehr_dataset(
  n_patients = 500,
  n_sites = 3,
  covid.Focused = TRUE,
  include_ct_links = FALSE,
  seed = NULL,
  verbose = TRUE
)
```

**Arguments**

n_patients	Number of unique patients.
n_sites	Number of sites/hospitals to simulate.
covid.Focused	Logical; if TRUE, use COVID-era encounter and lab patterns.
include_ct_links	Logical; if TRUE, add CT timing variables and a CT severity score in the CT research view.
seed	Optional integer used to set the random seed for reproducibility.
verbose	Logical; if TRUE, print progress messages to the console.

**Value**

A list with elements:

- tables** Named list of core EHR tables (patients, encounters, vitals, labs, medications, procedures, allergies).
- research** Named list with ct\_research\_view (if covid.Focused) and ml\_flat\_view (aggregated ML-ready table).
- metadata** List of high-level generation settings and table metadata.

**Examples**

```
ehr <- generate_hybrid_ehr_dataset(  
  n_patients = 10,  
  n_sites = 2,  
  covid.Focused = TRUE,  
  include_ct_links = FALSE,  
  seed = 123,  
  verbose = FALSE  
)  
  
names(ehr$tables)  
head(ehr$tables$patients)
```

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