

# Package ‘readepi’

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**Title** Read Data from Relational Database Management Systems and Health Information Systems

**Version** 1.0.1

**Description** Import Data from Relational Database Management Systems (RDBMS) and Health Information Systems ('HIS'). The current version of the package supports importing data from RDBMS including 'MS SQL', 'MySQL', 'PostGRESQL', and 'SQLite', as well as from two HIS platforms: 'DHIS2' and 'SORMAS'.

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**URL** <https://epiverse-trace.github.io/readepi/>,  
<https://github.com/epiverse-trace/readepi/>,  
<https://github.com/epiverse-trace/readepi>

**BugReports** <https://github.com/epiverse-trace/readepi/issues>

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---

dhis2_login	<i>Establish connection to a DHIS2 instance</i>
-------------	---

---

### Description

Establish connection to a DHIS2 instance

### Usage

```
dhis2_login(base_url, user_name, password)
```

**Arguments**

base\_url        A character with the base URL of the target DHIS2 instance  
user\_name       A character with the user name  
password        A character with the user's password

**Value**

An httr2\_response object if the connection was successfully established

**Examples**

```
## Not run:  
dhis2_log <- dhis2_login(  
  base_url = "https://play.im.dhis2.org/stable-2-42-1",  
  user_name = "admin",  
  password = "district"  
)  
  
## End(Not run)
```

---

get_api_version	<i>Get DHIS2 API version</i>
-----------------	------------------------------

---

**Description**

Get DHIS2 API version

**Usage**

```
get_api_version(login)
```

**Arguments**

login            A httr2\_response object returned by the dhis2\_login() function

**Value**

A numeric with minor version of the API

**Examples**

```
## Not run:  
# login to the DHIS2 instance  
dhis2_login <- login(  
  type = "dhis2",  
  from = "https://smc.moh.gm/dhis",  
  user_name = "test",  
  password = "Gambia@123"
```

```
)  
  
# get the API version  
api_version <- get_api_version(login = dhis2_login)  
  
## End(Not run)
```

---

get\_data\_elements      *Get all data elements from a specific DHIS2 instance*

---

### **Description**

Get all data elements from a specific DHIS2 instance

### **Usage**

```
get_data_elements(login)
```

### **Arguments**

login                    A httr2\_response object returned by the dhis2\_login() function

### **Value**

A data frame with the following two columns: the data elements IDs and their corresponding names.

### **Examples**

```
## Not run:  
# establish the connection to the system  
dhis2_login <- login(  
  type = "dhis2",  
  from = "https://smc.moh.gm/dhis",  
  user_name = "test",  
  password = "Gambia@123"  
)  
  
# retrieve the data elements  
data_elements <- get_data_elements(login = dhis2_login)  
  
## End(Not run)
```

---

`get_organisation_units`*Get the organization units from a specific DHIS2 instance*

---

**Description**

Retrieves all organisational reporting units and their levels, then builds a hierarchy for each unit by tracing its ancestries from the deepest level up to the root.

**Usage**

```
get_organisation_units(login)
```

**Arguments**

`login` A `httr2_response` object returned by the `dhis2_login()` function

**Details**

1. Fetches all organisation units via the `get_org_units()` function,
2. Fetches all organisational unit levels via the `get_org_unit_levels()` function,
3. Filters for organisational units at the deepest level,
4. Traces the parent hierarchy of each deepest unit up to the root,
5. Constructs a tabular structure where each row is a full lineage.

**Value**

A data frame where each row represents a full hierarchy for the last-level unit by keeping the hierarchical organizational unit's name and ID at each level, using the official level names provided by the DHIS2 instance like "Country Name", "Country ID", etc.

**Examples**

```
## Not run:  
# establish the connection to the system  
dhis2_login <- login(  
  type = "dhis2",  
  from = "https://smc.moh.gm/dhis",  
  user_name = "test",  
  password = "Gambia@123"  
)  
  
# fetch the organisation units  
org_units <- get_organisation_units(login = dhis2_login)  
  
## End(Not run)
```

---

get_programs	<i>get all programs from a given specific DHIS2 instance</i>
--------------	--

---

### Description

The function first fetches all programs from the DHIS2 Aggregate system, then distinguishes the Tracker and Aggregate programs.

### Usage

```
get_programs(login)
```

### Arguments

login            A httr2\_response object returned by the dhis2\_login() function

### Value

A data frame with the following columns: the program ID, the program name, and the program type specifying whether the program is part of the Aggregate or Tracker system.

### Examples

```
## Not run:
# establish the connection to the system
dhis2_login <- login(
  type = "dhis2",
  from = "https://smc.moh.gm/dhis",
  user_name = "test",
  password = "Gambia@123"
)

# fetch the programs
programs <- get_programs(login = dhis2_login)

## End(Not run)
```

---

get_program_org_units	<i>Get organisation units that are associated with a given program</i>
-----------------------	--

---

### Description

Get organisation units that are associated with a given program

### Usage

```
get_program_org_units(login, program, org_units = NULL)
```

**Arguments**

login	A httr2_response object returned by the dhis2_login() function
program	A character with the program ID or name
org_units	A data frame with all organisation units from target DHIS2 instance. This is the output from the get_organisation_units() function

**Value**

A data frame with the organisation units associated with the provided program

**Examples**

```
## Not run:
# login to the DHIS2 instance
dhis2_login <- login(
  type = "dhis2",
  from = "https://smc.moh.gm/dhis",
  user_name = "test",
  password = "Gambia@123"
)

# fetch the organisation units
org_units <- get_organisation_units(login = dhis2_login)

# get the organisation units associated with the following program
'E5IUQuHg3Mg'
target_org_units <- get_program_org_units(
  login = dhis2_login,
  program = "E5IUQuHg3Mg",
  org_units = org_units
)

## End(Not run)
```

---

get\_program\_stages      *Get program stages for one or more DHIS2 programs*

---

**Description**

Retrieves the stages associated with specified DHIS2 program IDs, or all programs if none are specified. If any of the supplied program names or IDs are not found, the function displays a message and proceeds with the valid ones.

**Usage**

```
get_program_stages(login, programs = NULL, program = NULL)
```

**Arguments**

login	A httr2_response object returned by the dhis2_login() function
programs	A data frame with the program IDs and names obtained from the get_programs() function
program	A character with the program ID or name

**Value**

A data frame with the following columns:

1. program\_id: the unique ID of the program
2. program\_name: the displayed name of the program
3. program\_stage\_name: the name of each stage associate with the program
4. program\_stage\_id: the ID of each program stage

**Examples**

```
## Not run:  
# establish the connection to the DHIS2 instance  
dhis2_login <- login(  
  type = "dhis2",  
  from = "https://smc.moh.gm/dhis",  
  user_name = "test",  
  password = "Gambia@123"  
)  
  
# get the list of all program stages from the DHIS2 instance  
all_program_stages <- get_program_stages(  
  login = dhis2_login,  
  program = "E5IUQuHg3Mg",  
  programs = NULL  
)  
  
## End(Not run)
```

---

get\_tracked\_entities *Get tracked entity attributes, their corresponding IDs and event IDs*

---

**Description**

Get tracked entity attributes, their corresponding IDs and event IDs

**Usage**

```
get_tracked_entities(login, api_version, org_unit, program, org_units)
```



**Arguments**

login	A httr2_response object returned by the dhis2_login() function
api_version	A numeric with the API version obtained from the get_api_version() function
org_unit	A character with the organisation unit ID or name
program	A character with the program ID or name
org_units	A data frame with all organisation units from target DHIS2 instance. This is the output from the get_organisation_units() function

**Value**

A data frame with the tracked entity attributes alongside their events and tracked entity IDs

**Examples**

```
## Not run:
# login to the DHIS2 instance
dhis2_login <- login(
  type = "dhis2",
  from = "https://smc.moh.gm/dhis",
  user_name = "test",
  password = "Gambia@123"
)

# set the program and org unit IDs
program <- "E5IUQuHg3Mg"
org_unit <- "GcLhRNAFppR"

# get the api version
api_version <- get_api_version(login = dhis2_login)

# get all the organisation units from the DHIS2 instance
org_units <- get_organisation_units(login = dhis2_login)

# get the tracked entity attributes
tracked_entity_attributes <- get_tracked_entities(
  login = dhis2_login,
  api_version = api_version,
  org_unit = org_unit,
  program = program,
  org_units = org_units
)

## End(Not run)
```

---

 login

*Establish a connection to the HIS of interest.*


---

### Description

The current version of the package supports basic authentication (using the username and password) and personal authentication key (using API key and bearer token).

### Usage

```
login(
  from,
  type,
  user_name = NULL,
  password = NULL,
  driver_name = NULL,
  db_name = NULL,
  port = NULL
)
```

### Arguments

from	The URL to the HIS of interest. For APIs, this must be the base URL (required).
type	The source name (required). The current version of the package covers the following RDBMS and HIS types: "ms sql", "mysql", "postgres", "sqlite", "dhis2", and "sormas".
user_name	The user name (optional).
password	The user's password (optional). When the password is not provided (set to NULL), the user will be prompted to enter the password.
driver_name	The driver name (optional). This is only needed for connecting to RDBMS only.
db_name	The database name (optional). This is only needed for connecting to RDBMS only.
port	The port ID (optional). This is only needed for connecting to RDBMS only.

### Value

A connection object

### Examples

```
# connect to the test MySQL server
## Not run:
login <- login(
  from = "mysql-rfam-public.ebi.ac.uk",
  type = "mysql",
  user_name = "rfamro",
```

```
    password = "",
    driver_name = "",
    db_name = "Rfam",
    port = 4497
  )

## End(Not run)

# connect to a DHIS2 instance
## Not run:
dhi2s_login <- login(
  type = "dhis2",
  from = "https://smc.moh.gm/dhis",
  user_name = "test",
  password = "Gambia@123"
)

## End(Not run)

# connect to SORMAS
## Not run:
sormas_login <- login(
  type = "sormas",
  from = "https://demo.sormas.org/sormas-rest",
  user_name = "SurvSup",
  password = "Lk5R7JXeZSEc"
)

## End(Not run)
```

---

lookup\_table

*Lookup table*

---

### Description

The lookup table is used to establish the correspondence between R operators, provided in an expression, and the DHIS2 or SQL operators.

### Usage

```
lookup_table
```

### Format

A data frame of 11 rows and 3 columns

---

read_dhis2	<i>Import data from DHIS2</i>
------------	-------------------------------

---

**Description**

Import data from DHIS2

**Usage**

```
read_dhis2(login, org_unit, program)
```

**Arguments**

login	A htr2_response object returned by the dhis2_login() function
org_unit	A character with the organisation unit ID or name
program	A character with the program ID or name

**Value**

A data frame that contains both the tracked entity attributes and their event data.

**Examples**

```
## Not run:
# login to the DHIS2 instance
dhis2_login <- login(
  type = "dhis2",
  from = "https://smc.moh.gm/dhis",
  user_name = "test",
  password = "Gambia@123"
)
program = "E5IUQuHg3Mg"
org_unit = "GcLhRNAFppR"
data <- read_dhis2(
  login = dhis2_login,
  org_unit = org_unit,
  program = program
)

# fetch data from the test DHIS2 instance
dhis2_login <- login(
  type = "dhis2",
  from = "https://play.im.dhis2.org/stable-2-42-1",
  user_name = "admin",
  password = "district"
)
org_unit <- "DiszpKrYNg8"
program <- "IpHINAT79UW"
```

```
data <- read_dhis2(  
  login = dhis2_login,  
  org_unit = org_unit,  
  program = program  
)  
  
## End(Not run)
```

---

read\_rdbms

*Import data from relational database management systems (RDBMS).*

---

### Description

The function assumes the user has read access to the database. Importing data from RDBMS requires the installation of the appropriate driver that is compatible with the server version hosting the database. For more details, see the [vignette](#) on how to install the driver.

### Usage

```
read_rdbms(login, query)
```

### Arguments

login	The connection object obtained from the <code>login()</code> function.
query	An SQL query or a list with the following elements: <ol style="list-style-type: none"><li>1. table: a string with the table name</li><li>2. fields: a vector of column names. When specified, only those columns will be returned. Default is NULL.</li><li>3. filter: an expression or a vector of values used to filter the rows from the table of interest. This should be of the same length as the value for the 'select'. Default is NULL.</li></ol>

### Value

A data.frame with the requested data as specified in the query argument.

### Examples

```
## Not run:  
# establish the connection to the database  
rdbms_login <- login(  
  from = "mysql-rfam-public.ebi.ac.uk",  
  type = "mysql",  
  user_name = "rfamro",  
  password = "",  
  driver_name = "",  
  db_name = "Rfam",  
  port = 4497
```

```

)

# import data where query parameters are specified as a list
authors_list <- read_rdbms(
  login = rdbms_login,
  query = list(table = "author", fields = NULL, filter = NULL)
)

# import data where query parameters is within an SQL query
authors_list <- read_rdbms(
  login = rdbms_login,
  query = "select * from author"
)

## End(Not run)

```

---

read\_sormas

*Import data from SORMAS*


---

## Description

The function returns the following columns by default: case\_id, person\_id, sex, date\_of\_birth, case\_origin, country, city, latitude, longitude, case\_status, date\_onset, date\_admission, outcome, date\_outcome, contact\_id, date\_last\_contact, date\_first\_contact, Ct\_values.

## Usage

```
read_sormas(login, disease, since = 0)
```

## Arguments

login	A list with the user's authentication details
disease	A character with the target disease name
since	A Date value in ISO8601 format (YYYY-mm-dd). Default is 0 i.e. to fetch all cases from the beginning of data collection.

## Details

Note that the some values in the date\_of\_birth column of the output object might not have some missing elements such a missing year (NA-12-26), month (2025-NA-01) or date (2025-12-NA), or a combination of two missing elements.

## Value

A data frame with the case data of the specified disease.

**Examples**

```
## Not run:
# establish the connection to the SORMAS system
sormas_login <- login(
  type = "sormas",
  from = "https://demo.sormas.org/sormas-rest",
  user_name = "SurvSup",
  password = "Lk5R7JXeZSEc"
)
# fetch all COVID (coronavirus) cases from the test SORMAS instance
covid_cases <- read_sormas(
  login = sormas_login,
  disease = "coronavirus"
)

## End(Not run)
```

---

request_parameters	<i>Request parameters</i>
--------------------	---------------------------

---

**Description**

The request\_parameters data frame contains the information related to which request parameters should be used when fetching data from a specific version of DHIS2. It currently contains parameters for versions ranging from 2.22 to 2.42.

**Usage**

```
request_parameters
```

**Format**

A data frame of 3 rows and 12 columns

---

show_tables	<i>Display the list of tables in a database</i>
-------------	---

---

**Description**

Display the list of tables in a database

**Usage**

```
show_tables(login)
```

**Arguments**

login            The connection object obtained from the login() function.

**Value**

a character that contains the list of all tables found in the specified database.

**Examples**

```
## Not run:
# connect to the test MySQL server
rdbms_login <- login(
  from      = "mysql-rfam-public.ebi.ac.uk",
  type      = "mysql",
  user_name = "rfamro",
  password  = "",
  driver_name = "",
  db_name   = "Rfam",
  port      = 4497
)

# display the list of available tables from this database
tables <- show_tables(login = rdbms_login)

## End(Not run)
```

---

sormas\_get\_api\_specification

*Download the API specification file from SORMAS*

---

**Description**

Download the API specification file from SORMAS

**Usage**

```
sormas_get_api_specification(path = tempdir(), overwrite = TRUE)
```

**Arguments**

path            A character with the path to the folder where the downloaded data dictionary should be stored. Default is NULL i.e. the data dictionary will be stored in tempdir()

overwrite       A logical used to specify whether to overwrite to overwrite the existing data dictionary or not. Default is TRUE



**Value**

Invisibly returns the path to the folder where the file is stored. When path = NULL, the file will be stored in the R temporary folder as: api\_specification.json

**Examples**

```
# save the SORMAS API specification into the temporary R folder
path_api_specs <- sormas_get_api_specification()
```

---

```
sormas_get_data_dictionary
      Download SORMAS data dictionary
```

---

**Description**

Download SORMAS data dictionary

**Usage**

```
sormas_get_data_dictionary(path = tempdir(), overwrite = TRUE)
```

**Arguments**

path	A character with the path to the folder where the downloaded data dictionary should be stored. Default is NULL i.e. the data dictionary will be stored in tempdir()
overwrite	A logical used to specify whether to overwrite to overwrite the existing data dictionary or not. Default is TRUE

**Value**

A character with path to the folder where the data dictionary is stored. When path = NULL, the file will be stored in the R temporary folder as: dictionary.xlsx

**Examples**

```
# download and save the data dictionary in the default R temporary directory
path_to_dictionary <- sormas_get_data_dictionary()
```

---

sormas\_get\_diseases    *Get the list of disease names from a SORMAS instance*

---

**Description**

Get the list of disease names from a SORMAS instance

**Usage**

```
sormas_get_diseases(login)
```

**Arguments**

login                    A list with the user's authentication details

**Value**

A vector of the list of disease names in a SORMAS instance

**Examples**

```
## Not run:
# establish the connection to the SORMAS system
sormas_login <- login(
  type = "sormas",
  from = "https://demo.sormas.org/sormas-rest",
  user_name = "SurvSup",
  password = "Lk5R7JXeZSEc"
)
disease_names <- sormas_get_diseases(
  login = sormas_login
)

## End(Not run)
```

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