

Package: cancerscreening (via r-universe)

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Title Streamline Access to Cancer Screening Data
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Description Retrieve cancer screening data for cervical, breast and colorectal cancers from the Kenya Health Information System <<https://hiskenya.org>> in a consistent way.
License MIT + file LICENSE
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<https://github.com/damurka/cancerscreening>
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cancerscreening-configuration
cancerscreening configuration

Description

Some aspects of cancerscreening behaviour can be controlled via an option.

Usage

```
with_cancerscreening_quiet(code)
```

```
local_cancerscreening_quiet(env = parent.frame())
```

Arguments

code	Code to execute quietly
env	The environment to use for scoping

Value

No return value, called for side effects

No return value, called for side effects

No return value, called for side effects

Messages

The `cancerscreening_quiet` option can be used to suppress messages from `cancerscreening`. By default, `cancerscreening` always messages, i.e. it is *not* quiet.

set `cancerscreening_quiet` to `TRUE` to suppress message, by one of these means, in order of decreasing scope:

- Put `options(cancerscreening_quiet = TRUE)` in the start-up file, such as `.Rprofile`, or in your R script
- Use `local_cancerscreening_quiet()` to silence `cancerscreening` in a specific scope
- Use `with_cancerscreening_quite` to run small bit of code silently

`local_cancerscreening_quiet` and `with_cancerscreening` follow the conventions of the `withr` package (<https://withr.r-lib.org>).

Examples

```
## Not run:
# message: "The credentials have been set."
khis_cred(username = 'username', password = 'password')

# suppress messages for a small amount of code
with_cancerscreening_quiet(
  khis_cred(username = 'username', password = 'password')
)

## End(Not run)

## Not run:
# message: "The credentials have been set."
khis_cred(username = 'username', password = 'password')

# suppress messages for a in a specific scope
local_cancerscreening_quiet()

# no message
khis_cred(username = 'username', password = 'password')

# clear credentials
khis_cred_clear()

## End(Not run)
```

get_analytics_formatted

Retrieves Analytics Table Data from KHIS

Description

`get_analytics_formatted()` fetches data from the KHIS analytics data tables for a given period and data element(s), without performing any aggregation.

Usage

```
get_analytics_formatted(  
  element_ids,  
  start_date,  
  end_date = NULL,  
  level = c("country", "county", "subcounty", "ward", "facility"),  
  organisations = NULL,  
  ...  
)
```

Arguments

<code>element_ids</code>	A vector of data element IDs for which to retrieve data. Required.
<code>start_date</code>	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
<code>end_date</code>	The ending date for data retrieval (default is the current date).
<code>level</code>	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
<code>organisations</code>	A list of organization units ids to be filtered.
<code>...</code>	Other options that can be passed onto KHIS API.

Details

- Retrieves data directly from KHIS analytics tables.
- Supports optional arguments for providing organization lists, data elements, and categories.
- Allows specifying KHIS session objects, retry attempts, and logging verbosity.

Value

A tibble with detailed information, including:

- Geographical identifiers (country, county, subcounty, ward, facility, depending on level)
- Reporting period (month, year, fiscal year)
- Data element names
- Category options
- Reported values

Examples

```
# Clinical Breast Examination data elements
# XEX93uLsAm2 = CBE Abnormal
# cXe64Yk0QMY = CBE Normal
element_id = c('cXe64Yk0QMY', 'XEX93uLsAm2')

# Download data from February 2023 to current date
data <- get_analytics_formatted(element_ids = element_id,
                               start_date = '2023-02-01')

data
```

get_breast_cbe	<i>Retrieves Data for Clinical Breast Examinations (CBE) Conducted</i>
----------------	--

Description

get_breast_cbe() retrieves data for CBE conducted within a specified period from the KHIS API server.

Usage

```
get_breast_cbe(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing data for CBE conducted with the following columns:

- country - Name of the country
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.

- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- age_group - The age group category of the report (25-34, 35-39, 40-55, 56-74, or 75+).
- category - Additional category if available.
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
cbe_data <- get_breast_cbe(start_date = '2023-02-01')
cbe_data
```

get_breast_mammogram *Retrieves Data for Mammograms Conducted*

Description

get_breast_mammogram() retrieves data for mammograms conducted within a specified period from the KHIS API server.

Usage

```
get_breast_mammogram(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing data for mammograms conducted with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report (25-34, 35-39, 40-55, 56-74, or 75+).
- category2 - Additional category if available.
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
mammogram_data <- get_breast_mammogram(start_date = '2023-02-01')
mammogram_data
```

get_breast_ultrasound *Retrieves Data for Breast Ultrasound Conducted*

Description

get_breast_ultrasound() retrieves data for breast ultrasounds conducted within a specified period from the KHIS API server.

Usage

```
get_breast_ultrasound(  
  start_date,  
  end_date = NULL,  
  level = c("country", "county", "subcounty", "ward", "facility"),  
  organisations = NULL,  
  ...  
)
```

Arguments

<code>start_date</code>	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
<code>end_date</code>	The ending date for data retrieval (default is the current date).
<code>level</code>	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
<code>organisations</code>	A list of organization units ids to be filtered.
<code>...</code>	Other options that can be passed onto KHIS API.

Value

A tibble containing data for breast ultrasound conducted with the following columns:

- `country` - Name of the country.
- `county` - Name of the county. Optional if the level is county, subcounty, ward or facility.
- `subcounty` - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- `ward` - Name of the ward. Optional if the level is ward or facility.
- `facility` - Name of the health facility. Optional if the level facility.
- `period` - The month and year of the data.
- `fiscal_year` - The financial year of the report(July-June Cycle).
- `year` - The calendar year of the report.
- `month` - The month name of the report.
- `category` - The age group category of the report (25-34, 35-39, 40-55, 56-74, or 75+).
- `category2` - Additional category if available.
- `element` - The data element.
- `value` - The number reported.

Examples

```
# Download data from February 2023 to current date
ultrasound_data <- get_breast_ultrasound(start_date = '2023-02-01')
ultrasound_data
```

 get_cervical_hiv_screened

Retrieves Cervical Cancer Screening Data on HIV Positive Women

Description

get_cervical_hiv_screened() retrieves cervical cancer screening and positivity data for HIV positive women for a specified period from the KHIS API server.

Usage

```
get_cervical_hiv_screened(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing cervical cancer screening data on HIV positive women with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report (<25, 25-49, 50+).

- category2 - Additional category if available.
- element - The data element (HPV, VIA or Pap Smear).
- source - The source report (MOH 711 or MOH 745).
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
screened <- get_cervical_hiv_screened(start_date = '2023-02-01')
screened
```

get_cervical_positive *Retrieves Cervical Cancer Screening Data with Positive Results*

Description

get_cervical_positive() retrieves cervical cancer screening data with positive results for a specified period from the KHIS API server.

Usage

```
get_cervical_positive(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing cervical cancer screening data with positive results with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.

- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report (<25, 25-49, 50+).
- category2 - Additional category if available.
- element - The data element (HPV, VIA or Pap Smear).
- source - The source report (MOH 711 or MOH 745).
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
positive <- get_cervical_positive(start_date = '2023-02-01')
positive
```

get_cervical_screened *Retrieves Cervical Cancer Screening Data*

Description

get_cervical_screened() retrieves cervical cancer screening data for a specified period from the KHIS API server.

Usage

```
get_cervical_screened(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing cervical cancer screening data with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report (<25, 25-49, 50+).
- category2 - Additional category if available.
- element - The data element (HPV, VIA or Pap Smear).
- source - The source report (MOH 711 or MOH 745).
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
screened <- get_cervical_screened(start_date = '2023-02-01')
screened
```

get_cervical_treated *Retrieves Cervical Cancer Precancerous Treatment Data*

Description

get_cervical_treated() retrieves cervical cancer precancerous treatment data for a specified period from the KHIS API server.

Usage

```
get_cervical_treated(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing cervical cancer precancerous treatment data with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report (<25, 25-49, 50+).
- category2 - Additional category if available.
- element - The data element (HPV, VIA or Pap Smear).
- source - The source report (MOH 711 or MOH 745).
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
treated <- get_cervical_treated(start_date = '2023-02-01')
treated
```

```
get_colorectal_colonoscopy
```

Retrieves Data for Colorectal Screening using Colonoscopy

Description

get_colorectal_colonoscopy() retrieves data for colorectal screening using colonoscopy within a specified period from the KHIS API server.

Usage

```
get_colorectal_colonoscopy(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing data for colorectal screening with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report (45-54, 55-64, or 65-75).
- category2 - Additional category if available.
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
data <- get_colorectal_colonoscopy(start_date = '2023-02-01')
data
```

get_colorectal_fobt *Retrieves Data for Colorectal Screening Using FOBT*

Description

get_colorectal_fobt() retrieves data for colorectal screening using FOBT within a specified period from the KHIS API server.

Usage

```
get_colorectal_fobt(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing data for colorectal screening with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.

- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report (45-54, 55-64, or 65-75).
- category2 - Additional category if available.
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
data <- get_colorectal_fobt(start_date = '2023-02-01')
data
```

```
get_filtered_population
```

Filters the Population

Description

get_filtered_population() filters the population based on age and level and projects the population base on the year provided

Usage

```
get_filtered_population(
  year,
  min_age,
  max_age,
  modifier = 1,
  level = c("country", "county", "subcounty"),
  pop_sex = c("female", "male", "both"),
  rate = 0.022
)
```

Arguments

year	The year to project the population
min_age	The minimum age to include in the filtered data
max_age	The maximum age to include in the filtered data
modifier	A multiplier that affect the population projection. Default 1
level	The desired level of the organization unit hierarchy to retrieve data for: "country", "county" or "subcounty".
pop_sex	The desired population sex: "male", "female" (default), "both"
rate	The population growth

Value

A tibble containing the target population

Examples

```
# Get the female population in 2022 aged 25-49 years
filtered_population <- get_filtered_population(2022, 25, 49, pop_sex = 'female')
filtered_population

# Get 5% male population in 2022 aged 40-75 years
filtered_population <- get_filtered_population(2022, 40, 75, modifier = 0.05, pop_sex = 'male')
filtered_population
```

get_lab_bone_marrow *Retrieves the Bone Marrow Laboratory Data*

Description

get_lab_bone_marrow() retrieves bone marrow lab data for a specified period from the KHIS API server.

Usage

```
get_lab_bone_marrow(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing bone marrow lab data with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.

- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
data <- get_lab_bone_marrow(start_date = '2023-02-01')
data
```

```
get_lab_fluid_cytology
```

Retrieves the Fluid Cytology Data

Description

get_lab_fluid_cytology() retrieves fluid cytology lab data for a specified period from the KHIS API server.

Usage

```
get_lab_fluid_cytology(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing fluid cytology lab data with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
data <- get_lab_fluid_cytology(start_date = '2023-02-01')
data
```

get_lab_fna

Retrieves the Fine-Needle Aspiration Laboratory Data

Description

get_lab_fna() retrieves fine-needle aspiration lab data for a specified period from the KHIS API server.

Usage

```
get_lab_fna(  
  start_date,  
  end_date = NULL,  
  level = c("country", "county", "subcounty", "ward", "facility"),  
  organisations = NULL,  
  ...  
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing fine-needle aspiration lab data with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
data <- get_lab_fna(start_date = '2023-02-01')
data
```

get_lab_smears

Retrieves the Cytology Smears Laboratory Data

Description

get_lab_smears() retrieves cytology smears lab data for a specified period from the KHIS API server.

Usage

```
get_lab_smears(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing cytology smears lab data with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
data <- get_lab_smears(start_date = '2023-02-01')
data
```

```
get_lab_tissue_histology
```

Retrieves the Tissue Histology Laboratory Data

Description

get_lab_tissue_histology() retrieves tissue histology lab data for a specified period from the KHIS API server.

Usage

```
get_lab_tissue_histology(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble containing tissue histology lab data with the following columns:

- country - Name of the country.
- county - Name of the county. Optional if the level is county, subcounty, ward or facility.
- subcounty - Name of the subcounty. Optional if the level is subcounty, ward or facility.
- ward - Name of the ward. Optional if the level is ward or facility.
- facility - Name of the health facility. Optional if the level facility.
- period - The month and year of the data.
- fiscal_year- The financial year of the report(July-June Cycle).
- year - The calendar year of the report.
- month - The month name of the report.
- category - The age group category of the report
- element - The data element.
- value - The number reported.

Examples

```
# Download data from February 2023 to current date
data <- get_lab_tissue_histology(start_date = '2023-02-01')
data
```

```
get_screening_reporting_analytics
  Retrieves Reporting Metrics for Screening Tool
```

Description

get_screening_reporting_analytics() It fetches the reporting metrics for the screening tool (MOH 745).

Usage

```
get_screening_reporting_analytics(
  start_date,
  end_date = NULL,
  level = c("country", "county", "subcounty", "ward", "facility"),
  organisations = NULL,
  ...
)
```

Arguments

start_date	The start date to retrieve data. It is required and in the format YYYY-MM-dd.
end_date	The ending date for data retrieval (default is the current date).
level	The desired data granularity: "country" (the default), "county", "subcounty", "ward", or "facility".
organisations	A list of organization units ids to be filtered.
...	Other options that can be passed onto KHIS API.

Value

A tibble with the reporting metrics.

Examples

```
# Download screening metric from February 2023 to current date
data <- get_screening_reporting_analytics(start_date = '2023-02-01')
data
```

target_population *Screening Target Populations*

Description

These functions subsets the Kenyan population to the desirable screening population.

Usage

```
get_cervical_target_population(
  year,
  level = c("country", "county", "subcounty")
)
```

```
get_breast_cbe_target_population(
  year,
  level = c("country", "county", "subcounty")
)
```

```
get_breast_mammogram_target_population(
  year,
  level = c("country", "county", "subcounty")
)
```

```
get_colorectal_target_population(
  year,
  level = c("country", "county", "subcounty")
)
```

Arguments

year	Year for which to estimate population.
level	The desired level of the organization unit hierarchy to retrieve data for: "country" (default), "county" or "subcounty".

Details

get_cervical_target_population() subsets the target population for cervical cancer screening: females aged between 25 years and 50 years

get_breast_cbe_target_population() subsets the target population for clinical breast examination: females aged between 25 years and 74 years

get_breast_mammogram_target_population() subsets the target population for breast cancer screening through mammography: females aged between 40 years to 74 years

get_colorectal_target_population() subsets the target population for colorectal cancer screening: males and females aged between 45 years to 75 years

These target populations are guided by the [Kenya National Cancer Screening Guidelines 2018](#). The population projection for counties and the national level are calculated based on population growth 2.2% obtained from the [Kenya National Bureau of Statistics](#). The annual targets follows the guidance of screening guidelines and for cervical cancer it is also guided by the WHO publication 'Planning and implementing cervical cancer prevention programs: A manual for managers.'

Value

A tibble containing the target screening population

- county - name of the county. Optional if the level is county or subcounty
- subcounty - name of the county. Optional if the level if subcounty
- target - number to be screened

A tibble containing the target screening population

A tibble containing the target screening population

A tibble containing the target screening population

Examples

```
# Get the country projection for cervical cancer screening for the year 2024
target_population <- get_cervical_target_population(2024)
target_population

# Get the projection for cervical cancer screening for 2022 by county
target_population <- get_cervical_target_population(2022, level = 'county')
target_population

# Get the projection for CBE for 2022 by county
target_population <- get_breast_cbe_target_population(2022, level = 'county')
target_population

# Get the country projection of women to perform mammogram for the year 2024
target_population <- get_breast_mammogram_target_population(2024)
target_population

# Get the country projection colorectal cancer screening for the year 2024
target_population <- get_colorectal_target_population(2024)
target_population
```

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