

Package ‘pathling’

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Type Package

Title A Library for using 'Pathling'

Version 6.4.2

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Description R API for 'Pathling', a tool for querying and transforming electronic health record data that is represented using the 'Fast Healthcare Interoperability Resources' (FHIR) standard - see <<https://pathling.csiro.au/docs>>.

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URL <https://pathling.csiro.au/>

BugReports <https://github.com/aehec/pathling/issues>

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R topics documented:

conditions	3
ds_aggregate	3
ds_extract	4
ds_read	5
ds_write_delta	6
ds_write_ndjson	7
ds_write_parquet	8
ds_write_tables	9
Equivalence	10
ImportMode	11
LOINC_URI	11
MimeType	12
pathling_connect	12
pathling_disconnect	15
pathling_disconnect_all	15
pathling_encode	16
pathling_encode_bundle	17
pathling_examples	18
pathling_example_resource	18
pathling_install_spark	19
pathling_is_spark_installed	20
pathling_read_bundles	20
pathling_read_datasets	21
pathling_read_delta	22
pathling_read_ndjson	23
pathling_read_parquet	24
pathling_read_tables	24
pathling_spark	25
pathling_spark_info	26
pathling_version	26
PropertyType	27
SNOMED_URI	27
StorageType	28
to_array	28
tx_designation	29
tx_display	30
tx_member_of	31
tx_property_of	31
tx_subsumed_by	32
tx_subsumes	33
tx_to_coding	34
tx_to_ecl_value_set	35
tx_to_loinc_coding	36
tx_to_snomed_coding	37
tx_translate	38
Version	39

conditions 3

Index 40

conditions *Synthetic conditions data*

Description

A synthetic data set of simplified and flattened FHIR Condition resources generated by Synthea.

Usage

`conditions`

Format

An object of class `data.frame` with 19 rows and 6 columns.

Details

A data frame with 19 rows and 6 columns:

- `START` - The onset date
- `STOP` - The abatement date
- `PATIENT` - The ID of the patient
- `ENCOUNTER` - The ID of the encounter
- `CODE` - The SNOMED CT code of the condition
- `DESCRIPTION` - The display name of the condition

ds_aggregate *Execute an aggregate query*

Description

Executes an aggregate query over FHIR data. The query calculates summary values based on aggregations and groupings of FHIR resources.

Usage

```
ds_aggregate(  
  ds,  
  subject_resource,  
  aggregations,  
  groupings = NULL,  
  filters = NULL  
)
```

Arguments

ds	The DataSource object containing the data to be queried.
subject_resource	A string representing the type of FHIR resource to aggregate data from.
aggregations	A named list of FHIRPath expressions that calculate a summary value from each grouping. The expressions must be singular.
groupings	An optional named list of FHIRPath expressions that determine which groupings the resources should be counted within.
filters	An optional sequence of FHIRPath expressions that can be evaluated against each resource in the data set to determine whether it is included within the result. The expression must evaluate to a Boolean value. Multiple filters are combined using logical AND operation.

Value

A Spark DataFrame containing the aggregated data.

See Also

[Pathling documentation - Aggregate](#)

Other FHIRPath queries: [ds_extract\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_ndjson(pathling_examples('ndjson'))
data_source %>% ds_aggregate('Patient',
  aggregations = c(patientCount='count()', 'id.count()'),
  groupings = c('gender', givenName='name.given'),
  filters = c('birthDate > @1950-01-01')
)
pathling_disconnect(pc)
```

ds_extract

Execute an extract query

Description

Executes an extract query over FHIR data. This type of query extracts specified columns from FHIR resources in a tabular format.

Usage

```
ds_extract(ds, subject_resource, columns, filters = NULL)
```

Arguments

ds	The DataSource object containing the data to be queried.
subject_resource	A string representing the type of FHIR resource to extract data from.
columns	A named list of FHIRPath expressions that define the columns to include in the extract.
filters	An optional sequence of FHIRPath expressions that can be evaluated against each resource in the data set to determine whether it is included within the result. The expression must evaluate to a Boolean value. Multiple filters are combined using AND logic.

Value

A Spark DataFrame containing the extracted data.

See Also

[Pathling documentation - Extract](#)

Other FHIRPath queries: [ds_aggregate\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_ndjson(pathling_examples('ndjson'))
data_source %>% ds_extract('Patient',
  columns = c('gender', givenName='name.given'),
  filters = c('birthDate > @1950-01-01')
)
pathling_disconnect(pc)
```

ds_read

Get data for a resource type from a data source

Description

Get data for a resource type from a data source

Usage

```
ds_read(ds, resource_code)
```

Arguments

ds	The DataSource object.
resource_code	A string representing the type of FHIR resource to read data from.

Value

A Spark DataFrame containing the data for the given resource type.

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_ndjson(pathling_examples('ndjson'))
data_source %>% ds_read('Patient') %>% sparklyr::sdf_nrow()
data_source %>% ds_read('Condition') %>% sparklyr::sdf_nrow()
pathling_disconnect(pc)
```

ds_write_delta	<i>Write FHIR data to Delta files</i>
----------------	---------------------------------------

Description

Writes the data from a data source to a directory of Delta files.

Usage

```
ds_write_delta(ds, path, import_mode = ImportMode$OVERWRITE)
```

Arguments

ds	The DataSource object.
path	The URI of the directory to write the files to.
import_mode	The import mode to use when writing the data - "overwrite" will overwrite any existing data, "merge" will merge the new data with the existing data based on resource ID.

Value

No return value, called for side effects only.

See Also

[Pathling documentation - Writing Delta](#)

[ImportMode](#)

Other data sink functions: [ds_write_ndjson\(\)](#), [ds_write_parquet\(\)](#), [ds_write_tables\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_ndjson(pathling_examples('ndjson'))

# Write the data to a directory of Delta files.
data_source %>% ds_write_delta(file.path(tempdir(), 'delta'), import_mode = ImportMode$OVERWRITE)

pathling_disconnect(pc)
```

ds_write_ndjson	<i>Write FHIR data to NDJSON files</i>
-----------------	--

Description

Writes the data from a data source to a directory of NDJSON files. The files will be named using the resource type and the ".ndjson" extension.

Usage

```
ds_write_ndjson(ds, path, file_name_mapper = NULL)
```

Arguments

ds	The DataSource object.
path	The URI of the directory to write the files to.
file_name_mapper	An optional function that can be used to customise the mapping of the resource type to the file name. Currently not implemented.

Value

No return value, called for side effects only.

See Also

[Pathling documentation - Writing NDJSON](#)

Other data sink functions: [ds_write_delta\(\)](#), [ds_write_parquet\(\)](#), [ds_write_tables\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_ndjson(pathling_examples('ndjson'))

# Write the data to a directory of NDJSON files.
data_source %>% ds_write_ndjson(file.path(tempdir(), 'ndjson'))
```

```
pathling_disconnect(pc)
```

ds_write_parquet *Write FHIR data to Parquet files*

Description

Writes the data from a data source to a directory of Parquet files.

Usage

```
ds_write_parquet(ds, path)
```

Arguments

ds	The DataSource object.
path	The URI of the directory to write the files to.

Value

No return value, called for side effects only.

See Also

[Pathling documentation - Writing Parquet](#)

Other data sink functions: [ds_write_delta\(\)](#), [ds_write_ndjson\(\)](#), [ds_write_tables\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_ndjson(pathling_examples('ndjson'))

# Write the data to a directory of Parquet files.
data_source %>% ds_write_parquet(file.path(tempdir(), 'parquet'))

pathling_disconnect(pc)
```

ds_write_tables	<i>Write FHIR data to managed tables</i>
-----------------	--

Description

Writes the data from a data source to a set of tables in the Spark catalog.

Usage

```
ds_write_tables(ds, schema = NULL, import_mode = ImportMode$OVERWRITE)
```

Arguments

ds	The DataSource object.
schema	The name of the schema to write the tables to.
import_mode	The import mode to use when writing the data - "overwrite" will overwrite any existing data, "merge" will merge the new data with the existing data based on resource ID.

Value

No return value, called for side effects only.

See Also

[Pathling documentation - Writing managed tables](#)

[ImportMode](#)

Other data sink functions: [ds_write_delta\(\)](#), [ds_write_ndjson\(\)](#), [ds_write_parquet\(\)](#)

Examples

```
# Create a temporary warehouse location, which will be used when we call ds_write_tables().
temp_dir_path <- tempfile()
dir.create(temp_dir_path)
sc <- sparklyr::spark_connect(master = "local[*]", config = list(
  "sparklyr.shell.conf" = c(
    paste0("spark.sql.warehouse.dir=", temp_dir_path),
    "spark.sql.extensions=io.delta.sql.DeltaSparkSessionExtension",
    "spark.sql.catalog.spark_catalog=org.apache.spark.sql.delta.catalog.DeltaCatalog"
  )
), version = pathling_spark_info()$spark_version)

pc <- pathling_connect(sc)
data_source <- pc %>% pathling_read_ndjson(pathling_examples('ndjson'))

# Write the data to a set of Spark tables in the 'default' database.
data_source %>% ds_write_tables("default", import_mode = ImportMode$MERGE)
```

```
pathling_disconnect(pc)
unlink(temp_dir_path, recursive = TRUE)
```

Equivalence

Concept map equivalence types

Description

The following values are supported:

- RELATEDTO - The concepts are related to each other, and have at least some overlap in meaning, but the exact relationship is not known.
- EQUIVALENT - The definitions of the concepts mean the same thing (including when structural implications of meaning are considered) (i.e. extensionally identical).
- EQUAL - The definitions of the concepts are exactly the same (i.e. only grammatical differences) and structural implications of meaning are identical or irrelevant (i.e. intentionally identical).
- WIDER - The target mapping is wider in meaning than the source concept.
- SUBSUMES - The target mapping subsumes the meaning of the source concept (e.g. the source is-a target).
- NARROWER - The target mapping is narrower in meaning than the source concept. The sense in which the mapping is narrower SHALL be described in the comments in this case, and applications should be careful when attempting to use these mappings operationally.
- SPECIALIZES - The target mapping specializes the meaning of the source concept (e.g. the target is-a source).
- INEXACT - There is some similarity between the concepts, but the exact relationship is not known.
- UNMATCHED - This is an explicit assertion that there is no mapping between the source and target concept.
- DISJOINT - This is an explicit assertion that the target concept is not in any way related to the source concept.

Usage

Equivalence

Format

An object of class `list` of length 10.

See Also

[FHIR R4 - ConceptMapEquivalence](#)

ImportMode

ImportMode

Description

The following import modes are supported:

- OVERWRITE: Overwrite any existing data.
- MERGE: Merge the new data with the existing data based on resource ID.

Usage

ImportMode

Format

An object of class `list` of length 2.

LOINC_URI

LOINC system URI

Description

The URI of the LOINC code system: <http://loinc.org>.

Usage

LOINC_URI

Format

An object of class `character` of length 1.

See Also

[Using LOINC with HL7 Standards](#)

MimeType	<i>FHIR MIME types</i>
----------	------------------------

Description

The following MIME types are supported:

- FHIR_JSON: FHIR resources encoded as JSON
- FHIR_XML: FHIR resources encoded as XML

Usage

MimeType

Format

An object of class `list` of length 2.

See Also

[FHIR R4 - Resource Formats](#)

pathling_connect	<i>Create or retrieve the Pathling context</i>
------------------	--

Description

Creates a Pathling context with the given configuration options.

Usage

```
pathling_connect(
  spark = NULL,
  max_nesting_level = 3,
  enable_extensions = FALSE,
  enabled_open_types = c("boolean", "code", "date", "dateTime", "decimal", "integer",
    "string", "Coding", "CodeableConcept", "Address", "Identifier", "Reference"),
  enable_terminology = TRUE,
  terminology_server_url = "https://tx.ontoserver.csiro.au/fhir",
  terminology_verbose_request_logging = FALSE,
  terminology_socket_timeout = 60000,
  max_connections_total = 32,
  max_connections_per_route = 16,
  terminology_retry_enabled = TRUE,
  terminology_retry_count = 2,
  enable_cache = TRUE,
```

```

cache_max_entries = 2e+05,
cache_storage_type = StorageType$MEMORY,
cache_storage_path = NULL,
cache_default_expiry = 600,
cache_override_expiry = NULL,
token_endpoint = NULL,
enable_auth = FALSE,
client_id = NULL,
client_secret = NULL,
scope = NULL,
token_expiry_tolerance = 120,
accept_language = NULL
)

```

Arguments

spark	A pre-configured SparkSession instance, use this if you need to control the way that the session is set up
max_nesting_level	Controls the maximum depth of nested element data that is encoded upon import. This affects certain elements within FHIR resources that contain recursive references, e.g., QuestionnaireResponse.item.
enable_extensions	Enables support for FHIR extensions
enabled_open_types	The list of types that are encoded within open types, such as extensions.
enable_terminology	Enables the use of terminology functions
terminology_server_url	The endpoint of a FHIR terminology service (R4) that the server can use to resolve terminology queries.
terminology_verbose_request_logging	Setting this option to TRUE will enable additional logging of the details of requests to the terminology service.
terminology_socket_timeout	The maximum period (in milliseconds) that the server should wait for incoming data from the HTTP service
max_connections_total	The maximum total number of connections for the client
max_connections_per_route	The maximum number of connections per route for the client
terminology_retry_enabled	Controls whether terminology requests that fail for possibly transient reasons should be retried
terminology_retry_count	The number of times to retry failed terminology requests
enable_cache	Set this to FALSE to disable caching of terminology requests

<code>cache_max_entries</code>	Sets the maximum number of entries that will be held in memory
<code>cache_storage_type</code>	The type of storage to use for the terminology cache
<code>cache_storage_path</code>	The path on disk to use for the cache
<code>cache_default_expiry</code>	The default expiry time for cache entries (in seconds)
<code>cache_override_expiry</code>	If provided, this value overrides the expiry time provided by the terminology server
<code>token_endpoint</code>	An OAuth2 token endpoint for use with the client credentials grant
<code>enable_auth</code>	Enables authentication of requests to the terminology server
<code>client_id</code>	A client ID for use with the client credentials grant
<code>client_secret</code>	A client secret for use with the client credentials grant
<code>scope</code>	A scope value for use with the client credentials grant
<code>token_expiry_tolerance</code>	The minimum number of seconds that a token should have before expiry when deciding whether to send it with a terminology request
<code>accept_language</code>	The default value of the Accept-Language HTTP header passed to the terminology server

Details

If no Spark session is provided and there is not one already present in this process, a new one will be created.

If a SparkSession is not provided, and one is already running within the current process, it will be reused.

It is assumed that the Pathling library API JAR is already on the classpath. If you are running your own cluster, make sure it is on the list of packages.

Value

A Pathling context instance initialized with the specified configuration

See Also

Other context lifecycle functions: [pathling_disconnect_all\(\)](#), [pathling_disconnect\(\)](#), [pathling_spark\(\)](#)

Examples

```
# Create PathlingContext for an existing Spark connecton.
sc <- sparklyr::spark_connect(master = "local")
pc <- pathling_connect(spark = sc)
pathling_disconnect(pc)
```

```
# Create PathlingContext with a new Spark connection.
pc <- pathling_connect()
spark <- pathling_spark(pc)
pathling_disconnect_all()
```

pathling_disconnect *Disconnect from the Spark session*

Description

Disconnects the Spark connection associated with a Pathling context.

Usage

```
pathling_disconnect(pc)
```

Arguments

pc The PathlingContext object.

Value

No return value, called for side effects only.

See Also

Other context lifecycle functions: [pathling_connect\(\)](#), [pathling_disconnect_all\(\)](#), [pathling_spark\(\)](#)

pathling_disconnect_all
Disconnect all Spark connections

Description

Disconnect all Spark connections

Usage

```
pathling_disconnect_all()
```

Value

No return value, called for side effects only.

See Also

Other context lifecycle functions: [pathling_connect\(\)](#), [pathling_disconnect\(\)](#), [pathling_spark\(\)](#)

pathling_encode	<i>Encode FHIR JSON or XML to a dataframe</i>
-----------------	---

Description

Takes a Spark DataFrame with string representations of FHIR resources in the given column and encodes the resources of the given types as Spark DataFrame.

Usage

```
pathling_encode(pc, df, resource_name, input_type = NULL, column = NULL)
```

Arguments

pc	The Pathling context object.
df	A Spark DataFrame containing the resources to encode.
resource_name	The name of the FHIR resource to extract (e.g., "Condition", "Observation").
input_type	The MIME type of input string encoding. Defaults to "application/fhir+json".
column	The column in which the resources to encode are stored. If set to NULL, the input DataFrame is assumed to have one column of type string.

Value

A Spark DataFrame containing the given type of resources encoded into Spark columns.

See Also

Other encoding functions: [pathling_encode_bundle\(\)](#)

Examples

```
pc <- pathling_connect()
json_resources_df <- pathling_spark(pc) %>%
  sparklyr::spark_read_text(path=system.file('extdata', 'ndjson', 'Condition.ndjson',
    package='pathling'))
pc %>% pathling_encode(json_resources_df, 'Condition')
pathling_disconnect(pc)
```

`pathling_encode_bundle`*Encode FHIR Bundles to a dataframe*

Description

Takes a dataframe with string representations of FHIR bundles in the given column and outputs a dataframe of encoded resources.

Usage

```
pathling_encode_bundle(pc, df, resource_name, input_type = NULL, column = NULL)
```

Arguments

<code>pc</code>	A Pathling context object.
<code>df</code>	A Spark DataFrame containing the bundles with the resources to encode.
<code>resource_name</code>	The name of the FHIR resource to extract (Condition, Observation, etc.).
<code>input_type</code>	The MIME type of the input string encoding. Defaults to 'application/fhir+json'.
<code>column</code>	The column in which the resources to encode are stored. If 'NULL', then the input DataFrame is assumed to have one column of type string.

Value

A Spark DataFrame containing the given type of resources encoded into Spark columns.

See Also

Other encoding functions: [pathling_encode\(\)](#)

Examples

```
pc <- pathling_connect()
json_resources_df <- pathling_spark(pc) %>%
  sparklyr::spark_read_text(path=system.file('extdata','bundle-xml', package='pathling'),
    whole = TRUE)
pc %>% pathling_encode_bundle(json_resources_df, 'Condition',
  input_type = MimeType$FHIR_XML, column = 'contents')
pathling_disconnect(pc)
```

pathling_examples *Get path to Pathling example data*

Description

Construct the path to the package example data in a platform-independent way.

Usage

```
pathling_examples(...)
```

Arguments

... character vector of the path components.

Value

The path to the examples data.

See Also

Other example functions: [pathling_example_resource\(\)](#)

Examples

```
pathling_examples('ndjson', 'Condition.ndjson')
```

pathling_example_resource
 Read resource from Pathling example data

Description

Reads a FHIR resource dataframe from the package example data.

Usage

```
pathling_example_resource(pc, resource_name)
```

Arguments

pc The PathlingContext object.
resource_name The name of the resource to read.

Details

The resources are read from the package example data in the `extdata/parquet` directory. Currently the following resources are available: 'Patient' and 'Condition'.

Value

A Spark DataFrame containing the resource data.

See Also

Other example functions: [pathling_examples\(\)](#)

Examples

```
pc <- pathling_connect()
pathling_example_resource(pc, 'Condition')
pathling_disconnect(pc)
```

`pathling_install_spark`
Install Spark

Description

Installs the version of Spark/Hadoop defined in the package metadata using the `sparklyr::spark_install` function.

Usage

```
pathling_install_spark()
```

Value

List with information about the installed version.

See Also

Other installation functions: [pathling_is_spark_installed\(\)](#), [pathling_spark_info\(\)](#), [pathling_version\(\)](#)

pathling_is_spark_installed
Check if Spark is installed

Description

Checks if the version of Spark/Hadoop required by Pathling is installed.

Usage

```
pathling_is_spark_installed()
```

Value

TRUE if the required version of Spark/Hadoop is installed, FALSE otherwise.

See Also

Other installation functions: [pathling_install_spark\(\)](#), [pathling_spark_info\(\)](#), [pathling_version\(\)](#)

pathling_read_bundles *Create a data source from FHIR bundles*

Description

Creates a data source from a directory containing FHIR bundles.

Usage

```
pathling_read_bundles(pc, path, resource_types, mime_type = MimeType$FHIR_JSON)
```

Arguments

pc	The PathlingContext object.
path	The URI of the directory containing the bundles.
resource_types	A sequence of resource type codes that should be extracted from the bundles.
mime_type	The MIME type of the bundles. Defaults to "application/fhir+json".

Value

A DataSource object that can be used to run queries against the data.

See Also

[Pathling documentation - Reading Bundles](#)

Other data source functions: [pathling_read_datasets\(\)](#), [pathling_read_delta\(\)](#), [pathling_read_ndjson\(\)](#), [pathling_read_parquet\(\)](#), [pathling_read_tables\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_bundles(pathling_examples('bundle-xml'),
  c("Patient", "Observation"), MimeType$FHIR_XML)
data_source %>% ds_read('Observation') %>% sparklyr::sdf_nrow()
pathling_disconnect(pc)
```

pathling_read_datasets

Create a data source from datasets

Description

Creates an immutable, ad-hoc data source from a named list of Spark datasets indexed with resource type codes.

Usage

```
pathling_read_datasets(pc, resources)
```

Arguments

pc	The PathlingContext object.
resources	A name list of Spark datasets, where the keys are resource type codes and the values are the data frames containing the resource data.

Value

A DataSource object that can be used to run queries against the data.

See Also

[Pathling documentation - Reading datasets](#)

Other data source functions: [pathling_read_bundles\(\)](#), [pathling_read_delta\(\)](#), [pathling_read_ndjson\(\)](#), [pathling_read_parquet\(\)](#), [pathling_read_tables\(\)](#)

Examples

```
pc <- pathling_connect()
patient_df <- pc %>% pathling_example_resource('Patient')
condition_df <- pc %>% pathling_example_resource('Condition')
data_source <- pc %>% pathling_read_datasets(list(Patient = patient_df, Condition = condition_df))
data_source %>% ds_read('Patient') %>% sparklyr::sdf_nrow()
pathling_disconnect(pc)
```

pathling_read_delta *Create a data source from Delta tables*

Description

pathling_read_delta() creates a data source from a directory containing Delta tables. Each table must be named according to the name of the resource type that it stores.

Usage

```
pathling_read_delta(pc, path)
```

Arguments

pc	The PathlingContext object.
path	The URI of the directory containing the Delta tables.

Value

A DataSource object that can be used to run queries against the data.

See Also

[Pathling documentation - Reading Delta](#)

Other data source functions: [pathling_read_bundles\(\)](#), [pathling_read_datasets\(\)](#), [pathling_read_ndjson\(\)](#), [pathling_read_parquet\(\)](#), [pathling_read_tables\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_delta(pathling_examples('delta'))
data_source %>% ds_read('Patient') %>% sparklyr::sdf_nrow()
pathling_disconnect(pc)
```

pathling_read_ndjson *Create a data source from NDJSON*

Description

Creates a data source from a directory containing NDJSON files. The files must be named with the resource type code and must have the ".ndjson" extension, e.g. "Patient.ndjson" or "Observation.ndjson".

Usage

```
pathling_read_ndjson(pc, path, extension = "ndjson", file_name_mapper = NULL)
```

Arguments

pc	The PathlingContext object.
path	The URI of the directory containing the NDJSON files.
extension	The file extension to use when searching for files. Defaults to "ndjson".
file_name_mapper	An optional function that maps a filename to the set of resource types that it contains. Currently not implemented.

Value

A DataSource object that can be used to run queries against the data.

See Also

[Pathling documentation - Reading NDJSON](#)

Other data source functions: [pathling_read_bundles\(\)](#), [pathling_read_datasets\(\)](#), [pathling_read_delta\(\)](#), [pathling_read_parquet\(\)](#), [pathling_read_tables\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_ndjson(pathling_examples('ndjson'))
data_source %>% ds_read('Patient') %>% sparklyr::sdf_nrow()
pathling_disconnect(pc)
```

pathling_read_parquet *Create a data source from Parquet tables*

Description

pathling_read_parquet() creates a data source from a directory containing Parquet tables. Each table must be named according to the name of the resource type that it stores.

Usage

```
pathling_read_parquet(pc, path)
```

Arguments

pc	The PathlingContext object.
path	The URI of the directory containing the Parquet tables.

Value

A DataSource object that can be used to run queries against the data.

See Also

[Pathling documentation - Reading Parquet](#)

Other data source functions: [pathling_read_bundles\(\)](#), [pathling_read_datasets\(\)](#), [pathling_read_delta\(\)](#), [pathling_read_ndjson\(\)](#), [pathling_read_tables\(\)](#)

Examples

```
pc <- pathling_connect()
data_source <- pc %>% pathling_read_parquet(pathling_examples('parquet'))
data_source %>% ds_read('Patient') %>% sparklyr::sdf_nrow()
pathling_disconnect(pc)
```

pathling_read_tables *Create a data source from managed tables*

Description

pathling_read_tables() creates a data source from a set of Spark tables, where the table names are the resource type codes.

Usage

```
pathling_read_tables(pc, schema = NULL)
```


Arguments

pc	The PathlingContext object.
schema	An optional schema name that should be used to qualify the table names.

Value

A DataSource object that can be used to run queries against the data.

See Also

[Pathling documentation - Reading managed tables](#)

Other data source functions: [pathling_read_bundles\(\)](#), [pathling_read_datasets\(\)](#), [pathling_read_delta\(\)](#), [pathling_read_ndjson\(\)](#), [pathling_read_parquet\(\)](#)

Examples

```
pc <- pathling_connect()
spark <- pathling_spark(pc)
data_source <- pc %>% pathling_read_tables()
data_source %>% ds_read('Patient') %>% sparklyr::sdf_nrow()
pathling_disconnect(pc)
```

pathling_spark	<i>Get the Spark session</i>
----------------	------------------------------

Description

Returns the Spark connection associated with a Pathling context.

Usage

```
pathling_spark(pc)
```

Arguments

pc	The PathlingContext object.
----	-----------------------------

Value

The Spark connection associated with this Pathling context.

See Also

Other context lifecycle functions: [pathling_connect\(\)](#), [pathling_disconnect_all\(\)](#), [pathling_disconnect\(\)](#)

pathling_spark_info *Get versions of Spark and other dependencies*

Description

Returns the versions of Spark and Spark packages used by the Pathling R library.

Usage

```
pathling_spark_info()
```

Value

A list containing the following keys:

- spark_version: The version of Spark used by Pathling.
- scala_version: The version of Scala used by Pathling.
- hadoop_version: The version of Hadoop used by Pathling.
- hadoop_major_version: The major version of Hadoop used by Pathling.
- delta_version: The version of Delta used by Pathling.

See Also

Other installation functions: [pathling_install_spark\(\)](#), [pathling_is_spark_installed\(\)](#), [pathling_version\(\)](#)

pathling_version *Get version of Pathling*

Description

Get version of Pathling

Usage

```
pathling_version()
```

Value

The version of the Pathling R library.

See Also

Other installation functions: [pathling_install_spark\(\)](#), [pathling_is_spark_installed\(\)](#), [pathling_spark_info\(\)](#)

PropertyType	<i>Coding property data types</i>
--------------	-----------------------------------

Description

The following data types are supported:

- STRING - A string value.
- INTEGER - An integer value.
- BOOLEAN - A boolean value.
- DECIMAL - A decimal value.
- DATETIME - A date/time value.
- CODE - A code value.
- CODING - A Coding value.

Usage

PropertyType

Format

An object of class list of length 7.

See Also

[FHIR R4 - Data Types](#)

SNOMED_URI	<i>SNOMED CT system URI</i>
------------	-----------------------------

Description

The URI of the SNOMED CT code system: <http://snomed.info/sct>.

Usage

SNOMED_URI

Format

An object of class character of length 1.

See Also

[Using SNOMED CT with HL7 Standards](#)

StorageType	<i>Terminology cache storage type</i>
-------------	---------------------------------------

Description

The type of storage to use for the terminology cache.

Usage

StorageType

Format

An object of class `list` of length 2.

Details

The following values are supported:

- MEMORY - Use an in-memory cache
- DISK - Use a disk-based cache

to_array	<i>Convert a vector to a SQL array literal</i>
----------	--

Description

Converts a vector to an expression with the corresponding SQL array literal.

Usage

to_array(value)

Arguments

value A character or numeric vector to be converted

Value

The quosure with the SQL array literal that can be used in `dplyr::mutate`.

tx_designation	<i>Get designations for codings</i>
----------------	-------------------------------------

Description

Takes a Coding column as its input. Returns a Column that contains the values of designations (strings) for this coding that match the specified use and language. If the language is not provided, then all designations with the specified type are returned regardless of their language.

Usage

```
tx_designation(coding, use = NULL, language = NULL)
```

Arguments

coding	A Column containing a struct representation of a Coding.
use	The code with the use of the designations.
language	The language of the designations.

Value

The Column containing the result of the operation (array of strings with designation values).

See Also

[Pathling documentation - Retrieving designations](#)

Examples

```
pc <- pathling_connect()

# Get the (first) SNOMED CT "Fully specified name" ('900000000000003001')
# for the first coding of the Condition resource, in the 'en' language.
pc %>% pathling_example_resource('Condition') %>%
  sparklyr::mutate(
    id,
    designation = (!!tx_designation(code[['coding']] [[0]],
      !!tx_to_snomed_coding('900000000000003001'), language = 'en')) [[0]],
    .keep='none')
pathling_disconnect(pc)
```

tx_display	<i>Get the display text for codings</i>
------------	---

Description

Takes a Coding column as its input. Returns a Column that contains the canonical display name associated with the given code.

Usage

```
tx_display(coding, accept_language = NULL)
```

Arguments

coding	A Column containing a struct representation of a Coding.
accept_language	The optional language preferences for the returned display name. Overrides the parameter 'accept_language' in pathling_connect .

Value

A Column containing the result of the operation (String).

See Also

[Pathling documentation - Multi-language support](#)

Other terminology functions: [tx_member_of\(\)](#), [tx_property_of\(\)](#), [tx_subsumed_by\(\)](#), [tx_subsumes\(\)](#), [tx_translate\(\)](#)

Examples

```
pc <- pathling_connect()

# Get the display name of the first coding of the Condition resource, with the default language.
pc %>% pathling_example_resource('Condition') %>%
  sparklyr::mutate(
    id,
    display = !!tx_display(code[['coding']][[0]]),
    .keep='none')

pathling_disconnect(pc)
```

tx_member_of	<i>Test membership within a value set</i>
--------------	---

Description

Takes a Coding or array of Codings column as its input. Returns the column which contains a Boolean value, indicating whether any of the input Codings is a member of the specified FHIR ValueSet.

Usage

```
tx_member_of(codings, value_set_uri)
```

Arguments

codings	A Column containing a struct representation of a Coding or an array of such structs.
value_set_uri	An identifier for a FHIR ValueSet.

Value

A Column containing the result of the operation.

See Also

[Pathling documentation - Value set membership](#)

Other terminology functions: [tx_display\(\)](#), [tx_property_of\(\)](#), [tx_subsumed_by\(\)](#), [tx_subsumes\(\)](#), [tx_translate\(\)](#)

tx_property_of	<i>Get properties for codings</i>
----------------	-----------------------------------

Description

Takes a Coding column as its input. Returns a Column that contains the values of properties for this coding with specified names and types. The type of the result column depends on the types of the properties. Primitive FHIR types are mapped to their corresponding SQL primitives. Complex types are mapped to their corresponding structs.

Usage

```
tx_property_of(  
  coding,  
  property_code,  
  property_type = "string",  
  accept_language = NULL  
)
```

Arguments

coding	A Column containing a struct representation of a Coding.
property_code	The code of the property to retrieve.
property_type	The type of the property to retrieve.
accept_language	The optional language preferences for the returned property values. Overrides the parameter 'accept_language' in 'PathlingContext.create'.

Value

The Column containing the result of the operation (array of property values).

See Also

[PropertyType](#)

[Pathling documentation - Retrieving properties](#)

Other terminology functions: [tx_display\(\)](#), [tx_member_of\(\)](#), [tx_subsumed_by\(\)](#), [tx_subsumes\(\)](#), [tx_translate\(\)](#)

Examples

```
pc <- pathling_connect()

# Get the (first) value of the `inactive` property of the first coding of the Condition resource.
pc %>% pathling_example_resource('Condition') %>%
  sparklyr::mutate(id,
    is_inavtive = (!!tx_property_of(code[['coding']][[0]],
      "inactive",PropertyType$BOOLEAN))[[0]],
    .keep='none'
  )

pathling_disconnect(pc)
```

tx_subsumed_by	<i>Test subsumption between codings</i>
----------------	---

Description

Takes two Coding columns as input. Returns a Column that contains a Boolean value, indicating whether the left Coding is subsumed by the right Coding.

Usage

```
tx_subsumed_by(left_codings, right_codings)
```


Arguments

`left_codings` A Column containing a struct representation of a Coding or an array of Codings.
`right_codings` A Column containing a struct representation of a Coding or an array of Codings.

Value

A Column containing the result of the operation (boolean).

See Also

[Pathling documentation - Subsumption testing](#)

Other terminology functions: [tx_display\(\)](#), [tx_member_of\(\)](#), [tx_property_of\(\)](#), [tx_subsumes\(\)](#), [tx_translate\(\)](#)

Examples

```
pc <- pathling_connect()

# Test the codings of the Condition `code` for subsumption by a SNOMED CT code.
pc %>% pathling_example_resource('Condition') %>%
  sparklyr::mutate(
    id,
    is_subsumed_by = !!tx_subsumed_by(code[['coding']],
    !!tx_to_snomed_coding('444814009')),
    .keep='none')

pathling_disconnect(pc)
```

tx_subsumes

Test subsumption between codings

Description

Takes two Coding columns as input. Returns a Column that contains a Boolean value, indicating whether the left Coding subsumes the right Coding.

Usage

```
tx_subsumes(left_codings, right_codings)
```

Arguments

`left_codings` A Column containing a struct representation of a Coding or an array of Codings.
`right_codings` A Column containing a struct representation of a Coding or an array of Codings.

Value

A Column containing the result of the operation (boolean).

See Also

[Pathling documentation - Subsumption testing](#)

Other terminology functions: [tx_display\(\)](#), [tx_member_of\(\)](#), [tx_property_of\(\)](#), [tx_subsumed_by\(\)](#), [tx_translate\(\)](#)

Examples

```
pc <- pathling_connect()

# Test the codings of the Condition `code` for subsumption of a SNOMED CT code.
pc %>% pathling_example_resource('Condition') %>%
  sparklyr::mutate(
    id,
    subsumes = !!tx_subsumes(code[['coding']],
      !!tx_to_snomed_coding('444814009')),
    .keep='none')

pathling_disconnect(pc)
```

tx_to_coding

Convert codes to Coding structures

Description

Converts a Column containing codes into a Column that contains a Coding struct.

Usage

```
tx_to_coding(coding_column, system, version = NULL)
```

Arguments

coding_column	The Column containing the codes.
system	The URI of the system the codes belong to.
version	The version of the code system.

Details

The Coding struct Column can be used as an input to terminology functions such as [tx_member_of](#) and [tx_translate](#). Please note that inside sparklyr verbs such as mutate the functions calls need to be preceded with !!, e.g: !!tx_to_coding(CODE, SNOMED_URI).

Value

A Column containing a Coding struct.

See Also

[FHIR R4 - Coding](#)

Other terminology helpers: [tx_to_ecl_value_set\(\)](#), [tx_to_loinc_coding\(\)](#), [tx_to_snomed_coding\(\)](#)

Examples

```
pc <- pathling_connect()
condition_df <- pathling_spark(pc) %>% sparklyr::copy_to(conditions)

# Convert codes to ICD-10 codings.
condition_df %>% sparklyr::mutate(
  icdCoding = !!tx_to_coding(CODE, "http://hl7.org/fhir/sid/icd-10"), .keep = 'none'
)

pathling_disconnect(pc)
```

tx_to_ecl_value_set *Convert a SNOMED CT ECL expression to a ValueSet URI*

Description

Converts a SNOMED CT ECL expression into a FHIR ValueSet URI. It can be used with the [tx_member_of](#) function.

Usage

```
tx_to_ecl_value_set(ecl)
```

Arguments

ecl The ECL expression.

Value

The ValueSet URI.

See Also

[Using SNOMED CT with HL7 Standards - Implicit Value Sets](#)

Other terminology helpers: [tx_to_coding\(\)](#), [tx_to_loinc_coding\(\)](#), [tx_to_snomed_coding\(\)](#)

Examples

```
# Example usage of tx_to_ecl_value_set function
tx_to_ecl_value_set('<<373265006 |Analgesic (substance)|')
```

tx_to_loinc_coding	<i>Convert LOINC codes to Coding structures</i>
--------------------	---

Description

Converts a Column containing codes into a Column that contains a LOINC Coding struct.

Usage

```
tx_to_loinc_coding(coding_column, version = NULL)
```

Arguments

coding_column	The Column containing the codes.
version	The version of the code system.

Details

The Coding struct Column can be used as an input to terminology functions such as [tx_member_of](#) and [tx_translate](#). Please note that inside sparklyr verbs such as `mutate` the functions calls need to be preceded with `!!`, e.g: `!!tx_to_coding(CODE, SNOMED_URI)`.

Value

A Column containing a Coding struct.

See Also

Other terminology helpers: [tx_to_coding\(\)](#), [tx_to_ecl_value_set\(\)](#), [tx_to_snomed_coding\(\)](#)

Examples

```
pc <- pathling_connect()
condition_df <- pathling_spark(pc) %>% sparklyr::copy_to(conditions)

# Convert codes to LOINC codings.
# Equivalent to: tx_to_coding(CODE, "http://loinc.org")
condition_df %>% sparklyr::mutate(loincCoding = !!tx_to_loinc_coding(CODE), .keep = 'none')

pathling_disconnect(pc)
```

tx_to_snomed_coding *Convert SNOMED CT codes to Coding structures*

Description

Converts a Column containing codes into a Column that contains a SNOMED Coding struct.

Usage

```
tx_to_snomed_coding(coding_column, version = NULL)
```

Arguments

coding_column The Column containing the codes.
version The version of the code system.

Details

The Coding struct Column can be used as an input to terminology functions such as [tx_member_of](#) and [tx_translate](#). Please note that inside sparklyr verbs such as `mutate` the functions calls need to be preceded with `!!`, e.g: `!!tx_to_coding(CODE, SNOMED_URI)`.

Value

A Column containing a Coding struct.

See Also

Other terminology helpers: [tx_to_coding\(\)](#), [tx_to_ecl_value_set\(\)](#), [tx_to_loinc_coding\(\)](#)

Examples

```
pc <- pathling_connect()
condition_df <- pathling_spark(pc) %>% sparklyr::copy_to(conditions)

# Convert codes to SNOMED CT codings.
# Equivalent to: tx_to_coding(CODE, "http://snomed.info/sct")
condition_df %>% sparklyr::mutate(snomedCoding = !!tx_to_snomed_coding(CODE), .keep = 'none')

pathling_disconnect(pc)
```

tx_translate	<i>Translate between value sets</i>
--------------	-------------------------------------

Description

Takes a Coding column as input. Returns the Column which contains an array of Coding value with translation targets from the specified FHIR ConceptMap. There may be more than one target concept for each input concept. Only the translation with the specified equivalences are returned.

Usage

```
tx_translate(
  codings,
  concept_map_uri,
  reverse = FALSE,
  equivalences = NULL,
  target = NULL
)
```

Arguments

codings	A Column containing a struct representation of a Coding.
concept_map_uri	An identifier for a FHIR ConceptMap.
reverse	The direction to traverse the map. FALSE results in "source to target" mappings, while TRUE results in "target to source".
equivalences	A value of a collection of values from the ConceptMapEquivalence ValueSet.
target	Identifies the value set in which a translation is sought. If there's no target specified, the server should return all known translations.

Value

A Column containing the result of the operation (an array of Coding structs).

See Also

[Equivalence](#)

[Pathling documentation - Concept translation](#)

Other terminology functions: [tx_display\(\)](#), [tx_member_of\(\)](#), [tx_property_of\(\)](#), [tx_subsumed_by\(\)](#), [tx_subsumes\(\)](#)

Examples

```
pc <- pathling_connect()

# Translates the codings of the Condition `code` using a SNOMED implicit concept map.
pc %>% pathling_example_resource('Condition') %>%
  sparklyr::mutate(
    id,
    translation = !!tx_translate(code[['coding']],
      'http://snomed.info/sct?fhir_cm=900000000000527005'),
    .keep='none')

pathling_disconnect(pc)
```

Version

FHIR versions

Description

The following FHIR versions are supported:

- R4: FHIR R4

Usage

Version

Format

An object of class `list` of length 1.

Index

- * **FHIRPath queries**
 - ds_aggregate, 3
 - ds_extract, 4
- * **Terminology functions**
 - tx_designation, 29
- * **context lifecycle functions**
 - pathling_connect, 12
 - pathling_disconnect, 15
 - pathling_disconnect_all, 15
 - pathling_spark, 25
- * **data sink functions**
 - ds_write_delta, 6
 - ds_write_ndjson, 7
 - ds_write_parquet, 8
 - ds_write_tables, 9
- * **data source functions**
 - pathling_read_bundles, 20
 - pathling_read_datasets, 21
 - pathling_read_delta, 22
 - pathling_read_ndjson, 23
 - pathling_read_parquet, 24
 - pathling_read_tables, 24
- * **datasets**
 - conditions, 3
 - Equivalence, 10
 - ImportMode, 11
 - LOINC_URI, 11
 - MimeType, 12
 - PropertyType, 27
 - SNOMED_URI, 27
 - StorageType, 28
 - Version, 39
- * **encoding functions**
 - pathling_encode, 16
 - pathling_encode_bundle, 17
- * **example functions**
 - pathling_example_resource, 18
 - pathling_examples, 18
- * **installation functions**
 - pathling_install_spark, 19
 - pathling_is_spark_installed, 20
 - pathling_spark_info, 26
 - pathling_version, 26
- * **terminology functions**
 - tx_display, 30
 - tx_member_of, 31
 - tx_property_of, 31
 - tx_subsumed_by, 32
 - tx_subsumes, 33
 - tx_translate, 38
- * **terminology helpers**
 - tx_to_coding, 34
 - tx_to_ecl_value_set, 35
 - tx_to_loinc_coding, 36
 - tx_to_snomed_coding, 37
- conditions, 3
- ds_aggregate, 3, 5
- ds_extract, 4, 4
- ds_read, 5
- ds_write_delta, 6, 7–9
- ds_write_ndjson, 6, 7, 8, 9
- ds_write_parquet, 6, 7, 8, 9
- ds_write_tables, 6–8, 9
- Equivalence, 10, 38
- ImportMode, 6, 9, 11
- LOINC_URI, 11
- MimeType, 12
- pathling_connect, 12, 15, 25, 30
- pathling_disconnect, 14, 15, 15, 25
- pathling_disconnect_all, 14, 15, 15, 25
- pathling_encode, 16, 17
- pathling_encode_bundle, 16, 17
- pathling_example_resource, 18, 18

pathling_examples, [18](#), [19](#)
pathling_install_spark, [19](#), [20](#), [26](#)
pathling_is_spark_installed, [19](#), [20](#), [26](#)
pathling_read_bundles, [20](#), [21–25](#)
pathling_read_datasets, [21](#), [21](#), [22–25](#)
pathling_read_delta, [21](#), [22](#), [23–25](#)
pathling_read_ndjson, [21](#), [22](#), [23](#), [24](#), [25](#)
pathling_read_parquet, [21–23](#), [24](#), [25](#)
pathling_read_tables, [21–24](#), [24](#)
pathling_spark, [14](#), [15](#), [25](#)
pathling_spark_info, [19](#), [20](#), [26](#), [26](#)
pathling_version, [19](#), [20](#), [26](#), [26](#)
PropertyType, [27](#), [32](#)

SNOMED_URI, [27](#)
StorageType, [28](#)

to_array, [28](#)
tx_designation, [29](#)
tx_display, [30](#), [31–34](#), [38](#)
tx_member_of, [30](#), [31](#), [32–38](#)
tx_property_of, [30](#), [31](#), [31](#), [33](#), [34](#), [38](#)
tx_subsumed_by, [30–32](#), [32](#), [34](#), [38](#)
tx_subsumes, [30–33](#), [33](#), [38](#)
tx_to_coding, [34](#), [35–37](#)
tx_to_ecl_value_set, [35](#), [35](#), [36](#), [37](#)
tx_to_loinc_coding, [35](#), [36](#), [37](#)
tx_to_snomed_coding, [35](#), [36](#), [37](#)
tx_translate, [30–34](#), [36](#), [37](#), [38](#)

Version, [39](#)