

Package ‘influxdbclient’

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Title 'InfluxDB' 2.x Client

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Description

'InfluxDB' 2.x time-series database client. Supports both 'InfluxDB' OSS (<<https://portal.influxdata.com/downloads/>>) and Cloud (<<https://cloud2.influxdata.com/>>) version.

URL <https://github.com/influxdata/influxdb-client-r>

BugReports <https://github.com/influxdata/influxdb-client-r/issues>

Depends R (>= 3.4)

Encoding UTF-8

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Imports R6, htr, jsonlite, bit64, nanotime (>= 0.3.0), plyr

RoxygenNote 7.1.1

Suggests testthat, httpertest

NeedsCompilation no

Repository CRAN

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InfluxDBClient*InfluxDBClient*

Description

Client for querying from and writing to InfluxDB 2.x.

Format

An R6Class object

Public fields

url Database URL
token Authentication token
org Organization name
dialect Flux dialect
retryOptions Retry options

Methods**Public methods:**

- [InfluxDBClient\\$new\(\)](#)
- [InfluxDBClient\\$health\(\)](#)
- [InfluxDBClient\\$query\(\)](#)
- [InfluxDBClient\\$ready\(\)](#)
- [InfluxDBClient\\$write\(\)](#)
- [InfluxDBClient\\$clone\(\)](#)

Method new(): Creates instance of InfluxDBClient.

Usage:

```
InfluxDBClient$new(url, token, org, retryOptions = NULL)
```

Arguments:

url InfluxDB instance URL
token Authentication token
org Organization name
org Retry options. See `RetryOptions` for details. Set to TRUE for default retry options. Default is NULL which disables retries.

Method health(): Gets health info of the InfluxDB instance.

Usage:

```
InfluxDBClient$health()
```

Returns: Named list with name, message, status, version, commit elements or error

Method `query()`: Queries data in the InfluxDB instance.

Usage:

```
InfluxDBClient$query(
  text,
  POSIXctCol = c(`_time` = "time"),
  flatSingleResult = TRUE
)
```

Arguments:

`text` Flux query

`POSIXctCol` Flux time to (new) POSIXct column mapping (named list). Default is `c("_time" = "time")`.
Use `NULL` to skip it.

`flatSingleResult` Whether to return simple list when response contains only one result. Default is `TRUE`.

Returns: List of data frames. Data frame represents Flux table. It can be a named list of nested lists of data frames when query response contains multiple results (see Flux `yield`), or a simple list of data frames for single result response.

Method `ready()`: Gets readiness status of the InfluxDB instance.

Usage:

```
InfluxDBClient$ready()
```

Returns: Named list with `status`, `started` and `up` elements or error

Method `write()`: Writes data to the InfluxDB instance.

Usage:

```
InfluxDBClient$write(
  x,
  bucket,
  batchSize = 5000,
  precision = c("ns", "us", "ms", "s"),
  measurementCol = "_measurement",
  tagCols = NULL,
  fieldCols = c(`_field` = "_value"),
  timeCol = "_time",
  object = NULL,
  ...
)
```

Arguments:

`x` Data as (list of) `data.frame`

`bucket` Target bucket name

`batchSize` Batch size. Positive number or `FALSE` to disable. Default is `5000`.

`precision` Time precision

`measurementCol` Name of measurement column. Default is `"_measurement"`.

`tagCols` Names of tag (index) columns

`fieldCols` Names of field columns. In case of unpivoted data previously retrieved from InfluxDB, use default value ie. named list `c("_field" = "_value")`. For all other cases, just use simple vector of column names (see Examples).

`timeCol` Name of time column. The column values should be either of `nanotime` or `POSIXct` type. Default is `"_time"`.

`object` *Output object name.* For dry-run operation, specify the name of the object to receive the output. Default is `NULL`. For debugging purposes.

Method `clone()`: The objects of this class are cloneable with this method.

Usage:

```
InfluxDBClient$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

Examples

```
## Not run:
# Instantiation
client <- InfluxDBClient$new(url = "http://localhost:8086",
                             token = "my-token",
                             org = "my-org")

# Query
data <- client$query('from(bucket: "my-bucket") |> range(start: -1h)')

# Write
data <- data.frame(...)
client$write(data, bucket = "my-bucket", precision = "us",
             measurementCol = "name",
             tagCols = c("location", "id"),
             fieldCols = c("altitude", "temperature"),
             timeCol = "time")

# Ready status
ready <- client$ready()

# Healt info
ready <- client$health()

## End(Not run)
```

Description

Retry options may be specified as optional argument to `write`.

Format

An R6Class object

Public fields

`retryJitter` Maximum number of random milliseconds included in delay. Default is 0.
`retryInterval` First retry delay in seconds. Default is 5.
`maxDelay` Maximum delay between retries in seconds. Default is 125.
`maxRetryTime` Maximum time to spend retrying in seconds. Default is 180.
`maxAttempts` Number of retry attempts. Default is 5.
`exponentialBase` Base for exponential backoff strategy. Default is 2.

Methods

Public methods:

- `RetryOptions$new()`
- `RetryOptions$clone()`

Method new(): Creates instance of RetryOptions.

Usage:

```
RetryOptions$new(  
  retryJitter = 0,  
  retryInterval = 5,  
  maxDelay = 125,  
  maxRetryTime = 180,  
  maxAttempts = 5,  
  exponentialBase = 2,  
  ...  
)
```

Method clone(): The objects of this class are cloneable with this method.

Usage:

```
RetryOptions$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

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