

# Package: cloudosR (via r-universe)

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**Type** Package

**Title** 'Lifebit' Platform 'API' Client

**Version** 0.2.0

**Description** Interacts with the 'Lifebit' Platform Cohort Browser 'API' <<https://cloudos.lifebit.ai>>. Enables schema discovery, table exploration, and read-only 'SQL' query execution with policy-aware behavior and team-based access control for cohort data analysis. Requires bastion-enabled workspaces for 'API' access.

**License** MIT + file LICENSE

**Encoding** UTF-8

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**Imports** htr2, jsonlite, utils

**Suggests** testthat (>= 3.0.0), knitr, rmarkdown

**VignetteBuilder** knitr

**RoxygenNote** 7.3.3

**NeedsCompilation** no

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cloudos.cohort\_tables *List Cohort Tables*

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### Description

Retrieves the list of available database schemas and tables for a cohort.

### Usage

```
cloudos.cohort_tables(profilename = "", cohort_id = "")
```

### Arguments

profilename	Character. Name of the configured profile to use. If empty or NULL, uses the default profile.
cohort_id	Character. ID of the cohort to query schemas for.

### Value

List with schema information including databases, tables, and columns.

### Examples

```
## Not run:
# Get available schemas for a cohort
schemas <- cloudos.cohort_tables(
  profilename = "production",
  cohort_id = "your-cohort-id"
)

# Display databases
cat("Available databases:\n")
for (db in schemas) {
  cat("  ", db$database, "\n")
}

## End(Not run)
```

---

cloudos.configure	<i>Configure Lifebit Platform Profile</i>
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---

### Description

Stores API credentials and workspace context for a named profile. This is the required first step before any API wrapper call.

### Usage

```
cloudos.configure(  
  profilename = "",  
  apikey = "",  
  workspace_id = "",  
  base_url = "https://cloudos.lifebit.ai",  
  set_default = FALSE  
)
```

### Arguments

profilename	Character. Name of the profile to create or update.
apikey	Character. API key for authentication.
workspace_id	Character. Workspace/team ID for API requests.
base_url	Character. Base URL for Lifebit Platform API (default: "https://cloudos.lifebit.ai").
set_default	Logical. If TRUE, sets this profile as the default (default: FALSE).

### Value

Invisible NULL. Prints a success message.

### Examples

```
## Not run:  
cloudos.configure(  
  profilename = "production",  
  apikey = "your-api-key",  
  workspace_id = "5c6d3e9bd954e800b23f8c62",  
  set_default = TRUE  
)  
  
## End(Not run)
```

---

cloudos.profile\_list    *List Lifebit Platform Profiles*

---

### Description

Lists configured profiles so users can confirm available environments.

### Usage

```
cloudos.profile_list()
```

### Value

Data frame with profile names and metadata (workspace\_id, default, created\_at, updated\_at). Returns empty data frame if no profiles are configured.

### Examples

```
## Not run:
profiles <- cloudos.profile_list()
print(profiles)

## End(Not run)
```

---

cloudos.query            *Execute SQL Query (Orchestrator)*

---

### Description

High-level function that orchestrates the full query lifecycle: submit -> poll status -> fetch results as dataframe.

### Usage

```
cloudos.query(
  profilename = "",
  cohort_id = "",
  sql = "",
  poll_interval = 2,
  max_wait = 600,
  page_size = 1000,
  all_pages = TRUE
)
```

## Arguments

profilename	Character. Name of the configured profile to use. If empty or NULL, uses the default profile.
cohort_id	Character. ID of the cohort to query.
sql	Character. SQL query to execute.
poll_interval	Integer. Seconds between status checks (default: 2).
max_wait	Integer. Maximum seconds to wait for completion (default: 600).
page_size	Integer. Number of rows per page (default: 1000).
all_pages	Logical. Fetch all result pages automatically (default: TRUE). When TRUE, submits multiple async tasks (one per page) to fetch complete results.

## Details

**IMPORTANT:** Pagination works by submitting separate tasks for each page. When `all_pages=TRUE`, this function submits multiple async tasks (one per page), waits for all to complete, and combines the results. This may take longer for large result sets.

## Value

Data frame with query results.

## Examples

```
## Not run:
# Fetch all results
results <- cloudos.query(
  profilename = "production",
  cohort_id = "699edb4380a6867895f0c9e1",
  sql = "SELECT person_id, gender_concept_id FROM person LIMIT 500"
)

# Fetch only first page
results_page1 <- cloudos.query(
  profilename = "production",
  cohort_id = "699edb4380a6867895f0c9e1",
  sql = "SELECT person_id FROM person",
  all_pages = FALSE,
  page_size = 100
)

## End(Not run)
```

---

cloudos.query\_results *Fetch Async Query Results*

---

**Description**

Fetches results from a completed async SQL task and returns as a dataframe.

**Usage**

```
cloudos.query_results(profilename = "", task_id = "")
```

**Arguments**

profilename	Character. Name of the configured profile to use. If empty or NULL, uses the default profile.
task_id	Character. Task ID returned from cloudos.query_submit_async().

**Details**

Note: Pagination is controlled when submitting the query via cloudos.query\_submit\_async(), not when fetching results. This function returns whatever page the task was configured for.

**Value**

Data frame with query results from the page configured at submission time.

**Examples**

```
## Not run:  
# Fetch results for a task (returns the page configured when query was submitted)  
results <- cloudos.query_results(  
  profilename = "production",  
  task_id = "69a5c58d626fe626da0025ce"  
)  
  
## End(Not run)
```

---

cloudos.query\_status *Check Async Query Status*

---

**Description**

Returns the current status and metadata for a submitted async SQL task.

**Usage**

```
cloudos.query_status(profilename = "", task_id = "")
```

**Arguments**

profilename	Character. Name of the configured profile to use. If empty or NULL, uses the default profile.
task_id	Character. Task ID returned from cloudos.query_submit_async().

**Value**

List with task status, count of results, and other metadata.

**Examples**

```
## Not run:
status <- cloudos.query_status(
  profilename = "production",
  task_id = "69a5c58d626fe626da0025ce"
)
print(status$status)
print(status$count_of_results)

## End(Not run)
```

---

cloudos.query\_submit\_async  
*Submit Async SQL Query*

---

**Description**

Starts async SQL execution for a cohort and returns a task ID for tracking.

**Usage**

```
cloudos.query_submit_async(
  profilename = "",
  cohort_id = "",
  sql = "",
  pagination = NULL
)
```

**Arguments**

profilename	Character. Name of the configured profile to use. If empty or NULL, uses the default profile.
cohort_id	Character. ID of the cohort to query.
sql	Character. SQL query to execute.
pagination	List (optional). Pagination settings with pageNumber and pageSize. Example: list(pageNumber = 0, pageSize = 100). If NULL, API returns default page.

**Value**

List with task metadata including task\_id, status, and full response.

**Examples**

```
## Not run:
# Submit query without pagination
task <- cloudos.query_submit_async(
  profilename = "production",
  cohort_id = "699edb4380a6867895f0c9e1",
  sql = "SELECT person_id FROM person LIMIT 100"
)

# Submit query with pagination for page 2 with 50 rows per page
task <- cloudos.query_submit_async(
  profilename = "production",
  cohort_id = "699edb4380a6867895f0c9e1",
  sql = "SELECT person_id FROM person",
  pagination = list(pageNumber = 2, pageSize = 50)
)
print(task$task_id)

## End(Not run)
```

---

cloudos.sql\_validate    *Validate SQL Query*

---

**Description**

Validates SQL syntax and references before execution.

**Usage**

```
cloudos.sql_validate(profilename = "", sql = "")
```

**Arguments**

profilename	Character. Name of the configured profile to use. If empty or NULL, uses the default profile.
sql	Character. SQL query to validate.

**Value**

List with validation results including isValid, tableReferences, and columnReferences.

## Examples

```
## Not run:
# Validate a SQL query
validation <- cloudos.sql_validate(
  profilename = "production",
  sql = "SELECT person_id FROM person WHERE year_of_birth >= 1960"
)

if (validation$isValid) {
  cat("SQL is valid\n")
  cat("Tables:", paste(validation$tableReferences, collapse = ", "), "\n")
} else {
  cat("SQL is invalid:", validation$message, "\n")
}

## End(Not run)
```

---

print.cloudos\_tables *Print method for cloudos\_tables*

---

## Description

Print method for cloudos\_tables

## Usage

```
## S3 method for class 'cloudos_tables'
print(x, ...)
```

## Arguments

x	A cloudos_tables object
...	Additional arguments (unused)

## Value

The cloudos\_tables object x, returned invisibly. Called primarily for its side effect of printing a formatted list of schemas and tables to the console.

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