

# Package ‘MolgenisArmadillo’

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**Type** Package

**Version** 2.6.3

**Title** Armadillo Client for the Armadillo Service

**Description** A set of functions to manage data shared on a  
‘MOLGENIS Armadillo’ server.

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**Depends** R (>= 3.6)

**Imports** base64enc, httr, urltools, dplyr, purrr, stringr, tidyR,  
tibble, MolgenisAuth (>= 0.0.25), arrow, rlist, httr2, readr,  
cli

**Suggests** stringi, withr, knitr, testthat, webmockr, mockery, datasets,  
rmarkdown, DSMolgenisArmadillo

**License** LGPL (>= 2.1)

**URL** <https://github.com/molgenis/molgenis-r-armadillo/>,  
<https://molgenis.github.io/molgenis-r-armadillo/>

**BugReports** <https://github.com/molgenis/molgenis-r-armadillo/issues/>

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`.compress_resource`      *Helper function for compressing to an RDS file*

---

### Description

Helper function for compressing to an RDS file

### Usage

```
.compress_resource(resource, file)
```

### Arguments

<code>resource</code>	the resource to write to file
<code>file</code>	the name of the file (without extension)

**Value**

the extension of the file

---

.compress_table	<i>Helper function for compressing to a parquet file</i>
-----------------	--

---

**Description**

Helper function for compressing to a parquet file

**Usage**

.compress\_table(table, file)

**Arguments**

table	the table to write to file
file	the name of the file (without extension)

**Value**

the extension of the file

---

.format_api_posts	<i>Formats API posts based on subset definition</i>
-------------------	---

---

**Description**

Formats API posts based on subset definition

**Usage**

.format\_api\_posts(posts, subset\_def)

**Arguments**

posts	A list of API posts
subset_def	A tibble containing subset definition

**Value**

A tibble consisting of original subset\_def with columns 'posts' and 'status' appended.

---

`.get_linkfile_content` *Helper function to get the contents of a linkfile*

---

### Description

Helper function to get the contents of a linkfile

### Usage

`.get_linkfile_content(project, object_name)`

### Arguments

`project` projectname where the linkfile is stored  
`object_name` folder/name of linkfile

### Value

the contents of the linkfile

---

`.load_linked_table` *Helper function to extract the source parquet file in a linkfile*

---

### Description

Helper function to extract the source parquet file in a linkfile

### Usage

`.load_linked_table(file, columns)`

### Arguments

`file` source table parquet file  
`columns` character list of columns to select from source file

### Value

the contents of the file, as data frame

---

`.load_resource`      *Helper function to extract an RDS file*

---

### Description

Helper function to extract an RDS file

### Usage

`.load_resource(file)`

### Arguments

`file`      file to extract

### Value

the contents of the file

---

`.load_table`      *Helper function to extract a parquet file*

---

### Description

Helper function to extract a parquet file

### Usage

`.load_table(file)`

### Arguments

`file`      file to extract

### Value

the contents of the file, as data frame

---

`armadillo.copy_resource`  
*Copy resource*

---

### Description

Copy resource

### Usage

```
armadillo.copy_resource(  
  project,  
  folder,  
  name,  
  new_folder = folder,  
  new_name = name  
)
```

### Arguments

project	study or other variable collection
folder	the folder containing the resource
name	specific resource for copy action
new_folder	name of the folder in which to place the copy, defaults to folder
new_name	name of the copy, defaults to name

### Value

the response from the server

### Examples

```
## Not run:  
armadillo.copy_resource(  
  project = "gecko",  
  folder = "core_all",  
  name = "table1",  
  new_folder = "core_all_v2",  
)  
## End(Not run)
```

---

armadillo.copy\_table    *Copy table*

---

## Description

Copy table

## Usage

```
armadillo.copy_table(  
  project,  
  folder,  
  name,  
  new_folder = folder,  
  new_name = name  
)
```

## Arguments

project	study or other variable collection
folder	the folder containing the table
name	specific table for copy action
new_folder	name of the folder in which to place the copy, defaults to folder
new_name	name of the copy, defaults to name

## Value

the response from the server

## Examples

```
## Not run:  
armadillo.copy_table(  
  project = "gecko",  
  folder = "core_all",  
  name = "table1",  
  new_folder = "core_all_v2",  
)  
## End(Not run)
```

**armadillo.create\_project***Create a project for a variable collection*

---

**Description**

Create a project for a variable collection

**Usage**

```
armadillo.create_project(  
    project_name = NULL,  
    users = NULL,  
    overwrite_existing = "choose"  
)
```

**Arguments**

`project_name` The name of the project to create. The project name

- cannot be empty.
- must be no more than 56 characters.
- cannot end with a -.
- must consist of lowercase letters and numbers.

`users` A list collection of the users that should have access to the project

`overwrite_existing`

Character, specifying action to take if project still exists: 'choose' (default) displays a menu giving the option to overwrite or not, 'yes' overwrites the existing project and 'no' exists the function with a message.

**Examples**

```
## Not run:  
armadillo.create_project("gecko")  
  
## End(Not run)
```

---

```
armadillo.delete_project  
Delete project
```

---

### Description

A project represents usually a study or collection of variables

### Usage

```
armadillo.delete_project(project_name)
```

### Arguments

project\_name the name of the study or collection of variables name

### Examples

```
## Not run:  
armadillo.delete_project(project_name = "gecko")  
## End(Not run)
```

---

```
armadillo.delete_resource  
Delete resource
```

---

### Description

Delete resource

### Usage

```
armadillo.delete_resource(project, folder, name)
```

### Arguments

project project to delete the resource from  
folder folder to delete the resource from  
name resource name

**Examples**

```
## Not run:
armadillo.delete_resource(
  project = "gecko",
  folder = "core_all",
  name = "table1"
)

## End(Not run)
```

**armadillo.delete\_table**  
*Delete table*

**Description**

Delete table

**Usage**

```
armadillo.delete_table(project, folder, name)
```

**Arguments**

project	project to delete the table from
folder	folder to delete the table from
name	table name

**Examples**

```
## Not run:
armadillo.delete_table(
  project = "gecko",
  folder = "core_all",
  name = "table1"
)

## End(Not run)
```

---

```
armadillo.get_projects_info  
Gets the Projects information
```

---

## Description

Gets the Projects information

## Usage

```
armadillo.get_projects_info()
```

## Value

the projects and their information

## Examples

```
## Not run:  
armadillo.get_projects_info()  
  
## End(Not run)
```

---

```
armadillo.get_project_users  
Gets the users of an given project name
```

---

## Description

Gets the users of an given project name

## Usage

```
armadillo.get_project_users(project_name)
```

## Arguments

project\_name the name of the project to extract the users from

## Value

List of all users within "project\_name"

**Examples**

```
## Not run:
armadillo.get_project_users("some-project")

## End(Not run)
```

**armadillo.install\_packages**  
*Install package*

**Description**

Installs a user defined package into the provided profile. The package is automatically whitelisted after installation. Only available during development.

**Usage**

```
armadillo.install_packages(paths, profile = "default")
```

**Arguments**

paths	the path(s) to the package(s), can be a vector or a string
profile	the selected profile

**armadillo.list\_projects**  
*List the projects*

**Description**

List the projects

**Usage**

```
armadillo.list_projects()
```

**Value**

the projects

**Examples**

```
## Not run:
armadillo.list_projects()

## End(Not run)
```

---

```
armadillo.list_resources
```

*List the resources in a project*

---

## Description

List the resources in a project

## Usage

```
armadillo.list_resources(project)
```

## Arguments

project            the shared project in which the resources are located

## Value

the resources in the project

## Examples

```
## Not run:  
armadillo.list_resources("gecko")  
  
## End(Not run)
```

---

```
armadillo.list_tables List the tables in a project
```

---

## Description

List the tables in a project

## Usage

```
armadillo.list_tables(project)
```

## Arguments

project            the shared project in which the tables are located

## Value

the table names, without the extension

## Examples

```
## Not run:
armadillo.list_tables("gecko")

## End(Not run)
```

`armadillo.load_resource`

*Load a resource from a project*

## Description

Load a resource from a project

## Usage

```
armadillo.load_resource(project, folder, name)
```

## Arguments

project	study or collection variables
folder	the folder containing the resource
name	name of the resource

## Value

the loaded resource

## Examples

```
## Not run:
armadillo.load_resource(
  project = "gecko",
  folder = "core_all",
  name = "lc_core_1"
)

## End(Not run)
```

---

armadillo.load\_table *Load a table from a project*

---

### Description

Load a table from a project

### Usage

```
armadillo.load_table(project, folder, name)
```

### Arguments

project	study or collection variables
folder	the folder containing the table
name	name of the table

### Value

the contents of the table file, as data frame

### Examples

```
## Not run:  
armadillo.load_table(  
  project = "gecko",  
  folder = "core_all",  
  name = "lc_core_1"  
)  
  
## End(Not run)
```

---

armadillo.login *Login*

---

### Description

Interactively obtains an id token and uses it to create a session token for an Armadillo Service

### Usage

```
armadillo.login(armadillo)
```

### Arguments

armadillo	URL of the Armadillo server,
-----------	------------------------------

**Value**

the id token

**Examples**

```
## Not run:
armadillo.login(
  "https://armadillo.dev.molgenis.org"
)
armadillo.login("http://localhost:8080")

## End(Not run)
```

**armadillo.login\_basic** *Login with username / password (meant for dev and test environments)*

**Description**

Login with username / password (meant for dev and test environments)

**Usage**

```
armadillo.login_basic(armadillo, username, password)
```

**Arguments**

armadillo	URL of the Armadillo server
username	the username
password	the password

**Examples**

```
## Not run:
armadillo.login(
  "https://armadillo.dev.molgenis.org", "admin", "admin"
)
armadillo.login("http://localhost:8080", "admin", "admin")

## End(Not run)
```

---

```
armadillo.move_resource
  Move the resource
```

---

## Description

Move the resource

## Usage

```
armadillo.move_resource(
  project,
  folder,
  name,
  new_folder = folder,
  new_name = name
)
```

## Arguments

project	a study or collection of variables
folder	the folder containing the resource to move
name	a resource to move
new_folder	the folder to move the resource to, defaults to folder
new_name	use to rename the file, defaults to name

## Value

NULL, invisibly

## Examples

```
## Not run:
armadillo.move_resource(
  project = "gecko",
  folder = "core_all",
  name = "table1",
  new_folder = "core_all_v2",
)
## End(Not run)
```

**armadillo.move\_table** *Move the table*

## Description

Move the table

## Usage

```
armadillo.move_table(
  project,
  folder,
  name,
  new_folder = folder,
  new_name = name
)
```

## Arguments

project	a study or collection of variables
folder	the folder containing the table to move
name	a table to move
new_folder	the folder to move the table to, defaults to folder
new_name	use to rename the file, defaults to name

## Value

NULL, invisibly

## Examples

```
## Not run:
armadillo.move_table(
  project = "gecko",
  folder = "core_all",
  name = "table1",
  new_folder = "core_all_v2",
)
## End(Not run)
```

---

armadillo.subset	<i>Describes data available to subset and makes subset</i>
------------------	--

---

## Description

This automates the process of:

1. Checking what data is available to create subsets
2. Make the subset

## Usage

```
armadillo.subset(
  input_source = NULL,
  subset_def = NULL,
  source_project = NULL,
  source_folder = NULL,
  source_table = NULL,
  target_project = NULL,
  target_folder = NULL,
  target_table = NULL,
  target_vars = NULL,
  new_project = NULL,
  dry_run = NULL
)
```

## Arguments

<code>input_source</code>	Character specifying how information about the target view is provided: choose 'subset_def' if providing a subset definition object, or 'arguments' if providing information directly.
<code>subset_def</code>	R object containing subset definition created by <code>armadillo.subset_definition()</code> . Compulsory if <code>input_source = 'subset_def'</code>
<code>source_project</code>	project from which to subset data
<code>source_folder</code>	folder from which to subset data. Compulsory if <code>input_source = 'arguments'</code> .
<code>source_table</code>	table from which to subset data. Compulsory if <code>input_source = 'arguments'</code> .
<code>target_project</code>	project to upload subset to. Will be created if it doesn't exist.
<code>target_folder</code>	folder to upload subset to. Will be created if it doesn't exist. Compulsory if <code>input_source = 'arguments'</code> .
<code>target_table</code>	table to upload subset to. Compulsory if <code>input_source = 'arguments'</code> .
<code>target_vars</code>	variables from 'source_table' to include in the view. Compulsory if <code>input_source = 'arguments'</code> .
<code>new_project</code>	Deprecated: use <code>target_project</code> instead
<code>dry_run</code>	Defunct: previously enabled dry-run to check which variables are missing

**Value**

missing variables provided in the subset definition

**Examples**

```
## Not run:
armadillo.subset(
  source_project = "gecko",
  target_project = "study1",
  subset_def = local_subset
)
## End(Not run)
```

**armadillo.subset\_definition**

*Builds an R object containing info required to make subsets*

**Description**

Builds an R object containing info required to make subsets

**Usage**

```
armadillo.subset_definition(reference_csv = NULL, vars = NULL)
```

**Arguments**

reference_csv	.csv file containing details of the variable to subset. Must contain 5 columns: 'source_folder' specifying the folder from which to subset, 'souce_table' specifying the table from which to subset, 'target_folder' specifying the folder in which to create the subset 'target_table' specifying the name of the subset and 'variable' specifying the variable(s) to include in the subset. Note that 'source_project' and 'target_project' are specified as arguments to 'armadillo.subset'.
vars	Deprecated: use reference_csv instead

**Value**

A dataframe containing variables that is used for input in the `armadillo.subset()` method

**Examples**

```
## Not run:
armadillo.subset_definition(
  reference_csv = "C:/tmp/vars.csv"
)
```

```
## End(Not run)
```

---

**armadillo.upload\_resource**

*Uploads a resource to a folder in a project*

---

**Description**

Uploads a resource to a folder in a project

**Usage**

```
armadillo.upload_resource(project, folder, resource, name = NULL)
```

**Arguments**

project	the project to upload to
folder	the folder to upload to
resource	the resource to upload
name	name of the resource (optional)

**Examples**

```
## Not run:  
armadillo.upload_table(  
  project = "gecko",  
  folder = "core_all",  
  table1  
)  
  
## End(Not run)
```

---

**armadillo.upload\_table**

*Uploads a table to a folder in a project*

---

**Description**

Uploads a table to a folder in a project

**Usage**

```
armadillo.upload_table(project, folder, table, name = NULL)
```

**Arguments**

project	the project to upload to
folder	the folder to upload to
table	the table to upload
name	name of the table (optional)

**Examples**

```
## Not run:  
armadillo.upload_table(  
  project = "gecko",  
  folder = "core_all",  
  table1  
)  
  
## End(Not run)
```

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