

# Package ‘MolgenisArmadillo’

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**Type** Package

**Version** 2.9.1

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

**License** LGPL (>= 2.1)

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<https://molgenis.github.io/molgenis-r-armadillo/>

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*.add\_slash\_if\_missing Add a Slash to a URL if Missing*

---

### Description

This function ensures that a given URL string ends with a slash ('/'). If the URL does not end with a slash, it appends one.

### Usage

*.add\_slash\_if\_missing(url)*

### Arguments

url                    A character string representing the URL.

### Value

A character string with a trailing slash if it was missing.

---

.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

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

### Value

the extension of the file

---

.compress\_table        *Helper function for compressing to a parquet file*

---

### 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`      *Formats API posts based on subset definition*

### 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

<code>file</code>	source table parquet file
<code>columns</code>	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

<code>file</code>	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 <ul style="list-style-type: none"> <li>• cannot be empty.</li> <li>• must be no more than 56 characters.</li> <li>• cannot end with a -.</li> <li>• must consist of lowercase letters and numbers.</li> </ul> |
| 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_project_folder  
Delete project folder
```

---

## Description

Delete project folder

## Usage

```
armadillo.delete_project_folder(project, folder)
```

## Arguments

project project to delete the object from  
folder folder to delete the object from

`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**

<code>paths</code>	the path(s) to the package(s), can be a vector or a string
<code>profile</code>	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

<code>armadillo</code>	URL of the Armadillo server
<code>username</code>	the username
<code>password</code>	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,
  strict = FALSE
)
```

## Arguments

input_source	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.
subset_def	R object containing subset definition created by <code>armadillo.subset_definition()</code> . Compulsory if <code>input_source = 'subset_def'</code>
source_project	project from which to subset data
source_folder	folder from which to subset data. Compulsory if <code>input_source = 'arguments'</code> .
source_table	table from which to subset data. Compulsory if <code>input_source = 'arguments'</code> .
target_project	project to upload subset to. Will be created if it doesn't exist.
target_folder	folder to upload subset to. Will be created if it doesn't exist. Compulsory if <code>input_source = 'arguments'</code> .
target_table	table to upload subset to. Compulsory if <code>input_source = 'arguments'</code> .
target_vars	variables from 'source_table' to include in the view. Compulsory if <code>input_source = 'arguments'</code> .
new_project	Deprecated: use <code>target_project</code> instead

<code>dry_run</code>	Defunct: previously enabled dry-run to check which variables are missing
<code>strict</code>	Boolean specifying whether to create subset if one or more target variables do not exist in the target data. Option FALSE will throw an error, option TRUE (default) creates subset and return a warning

**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**

<code>reference_csv</code>	.csv file containing details of the variable to subset. Must contain 5 columns: 'source_folder' specifying the folder from which to subset, 'source_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'.
<code>vars</code>	Deprecated: use <code>reference_csv</code> 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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