Package 'mRatools'

May 9, 2023

Type Package	
Title Microbiological risk assessment tools	
Version 0.0.1	
Authors Mayooran Thevaraja [aut, cre], Kondaswamy Govindaraju [aut], Mark Bebbington [aut]	
<pre>URL https://github.com/Mayooran1987/mRatools</pre>	
<pre>BugReports https://github.com/Mayooran1987/mRatools/issues</pre>	
Description This package is designed to make it easy to install and load multiple microbiological risk assessment based R packages in a single step.	
License GPL (>= 2)	
Encoding UTF-8	
LazyData true	
Imports grabsampling, mixingsimulation, uneqmixr, dilutionrisk	
Remotes Mayooran1987/grabsampling, Mayooran1987/dilutionrisk, Mayooran1987/mixingsimulation, Mayooran1987/uneqmixr	
Suggests rstudioapi, rlang, cli, crayon, remotes, spelling, testthat (>= 3.0.0), covr, mockr	
RoxygenNote 7.2.3	
Depends R (>= 4.0)	
Maintainer Mayooran Thevaraja <mayooran@eng.jfn.ac.lk></mayooran@eng.jfn.ac.lk>	
Language en-US	
Config/testthat/edition 3	
R topics documented:	
mRatools_conflicts	2 2
Index	3

2 mRatools_packages

mRatools_conflicts

Conflicts between the mRatools and other packages

Description

This function lists all the conflicts between packages in the mRatools and other packages that you have loaded.

Usage

```
mRatools_conflicts(only = NULL)
```

Arguments

only

Set this to a character vector to restrict to conflicts only with these packages.

Details

There are no conflicts between the mRatools and other packages at this stage.

Examples

```
mRatools_conflicts()
```

 ${\tt mRatools_packages}$

List all packages in the mRatools

Description

List all packages in the mRatools

Usage

```
mRatools_packages(include_self = TRUE)
```

Arguments

Examples

```
mRatools_packages()
```

Index

mRatools_conflicts, 2
mRatools_packages, 2