# Package 'anyLib'

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Type Package	
Title Install and Load Any Package from CRAN, Bioconductor or Github	
Version 1.0.5	
<b>Description</b> Made to make your life simpler with packages, by installing and loading a list of packages, whether they are on CRAN, Bioconductor or github. For github, if you do not have the full path, with the maintainer name in it (e.g. ``achateigner/topReviGO"), it will be able to load it but not to install it.	
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Encoding UTF-8	
LazyData true	
RoxygenNote 6.1.0	
Imports devtools, withr, BiocManager, httr, curl	
Suggests knitr, rmarkdown, testthat	
VignetteBuilder knitr	
NeedsCompilation no	
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2 anyLib

anyLib	Install and load any library

#### Description

Made to make your life simpler with packages/libraries, by installing and loading a list of packages, whether they are on CRAN, Bioconductor or github. For github, if you do not have the full path, with the maintainer name in it (e.g. "achateigner/topReviGO"), it will not be able to install it. However, once installed you only need the name of the package. For more details see the help vignette: vignette("help", package = "anyLib")

#### Usage

```
anyLib(pkg, force = FALSE, autoUpdate = TRUE, lib = .libPaths()[1],
loadLib = .libPaths(), source = FALSE)
```

## **Arguments**

pkg The package name or the list containing the packages names.

force To force reinstallation of packages.

autoUpdate To select whether Bioconductor packages auto update or not.

lib Where to install the packages

loadLib From where the packages are loaded

source The package to install is a local source file, on the user disk.

#### **Details**

The source option can be a single TRUE or FALSE, or a vector of TRUE and FALSE corresponding to the vector/list of packages. E.g. if source == c(TRUE, FALSE), the first package will be considered as a source file. The file has to be a tar.gz source file, not a binary.

### Value

A named vector of booleans showing if the package is loaded properly

## **Examples**

```
# Install and load 1 package from a local source file, which name is in an object:
lib <- normalizePath(tempdir(), "/")
listOfPackages <- system.file("dummyPackage_0.1.0.tar.gz", package="anyLib")
anyLib(listOfPackages, force = TRUE, autoUpdate = FALSE, lib = lib,
loadLib = lib, source = TRUE)
```

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