

Package ‘BiocVersion’

October 15, 2018

Version 3.7.4

Title Set the appropriate version of Bioconductor packages

Description This package provides repository information for the appropriate version of Bioconductor.

biocViews Infrastructure

Depends R (>= 3.5.0), R (< 3.6.0)

License Artistic-2.0

Encoding UTF-8

LazyData true

RoxygenNote 6.0.1

git_url <https://git.bioconductor.org/packages/BiocVersion>

git_branch RELEASE_3_7

git_last_commit 470a683

git_last_commit_date 2018-03-21

Date/Publication 2018-10-15

Author Martin Morgan [aut],
Marcel Ramos [ctb],
Bioconductor Package Maintainer [ctb, cre]

Maintainer Bioconductor Package Maintainer <maintainer@bioconductor.org>

R topics documented:

BiocVersion-pkg	1
Index	3

BiocVersion-pkg	<i>Determine Bioconductor version in use</i>
-----------------	--

Description

The ‘major’ and ‘minor’ version of this package determines the version of Bioconductor software packages to be installed by the `install` function.

Examples

```
packageVersion("BiocVersion")[,1:2]
```

Index

BiocVersion (BiocVersion-pkg), [1](#)

BiocVersion-pkg, [1](#)

install, [/](#)