

# Package ‘BiocVersion’

November 27, 2023

**Version** 3.18.1

**Title** Set the appropriate version of Bioconductor packages

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

**biocViews** Infrastructure

**Depends** R (>= 4.3.0)

**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\_18

**git\_last\_commit** 70680b8

**git\_last\_commit\_date** 2023-11-15

**Repository** Bioconductor 3.18

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## R topics documented:

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