

# Package ‘BiocVersion’

November 14, 2024

**Version** 3.20.0

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

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

**biocViews** Infrastructure

**Depends** R (>= 4.4.0)

**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\_20

**git\_last\_commit** 43c716a

**git\_last\_commit\_date** 2024-04-30

**Repository** Bioconductor 3.20

**Date/Publication** 2024-11-14

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

**Maintainer** Bioconductor Package Maintainer <[maintainer@bioconductor.org](mailto:maintainer@bioconductor.org)>

## Contents

BiocVersion-pkg	2
<b>Index</b>	<b>3</b>

---

BiocVersion-pkg

*Determine Bioconductor version in use*

---

**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), [2](#)

BiocVersion-pkg, [2](#)

install, [2](#)