

# Package ‘BiocVersion’

May 14, 2026

**Version** 3.24.0

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

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

**biocViews** Infrastructure

**Depends** R (>= 4.6.0)

**License** Artistic-2.0

**Encoding** UTF-8

**RoxygenNote** 6.0.1

**git\_url** <https://git.bioconductor.org/packages/BiocVersion>

**git\_branch** devel

**git\_last\_commit** 84a87a5

**git\_last\_commit\_date** 2026-04-28

**Repository** Bioconductor 3.24

**Date/Publication** 2026-05-14

**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 . . . . . | 1        |
| <b>Index</b>              | <b>3</b> |

---

|                 |  |
|-----------------|--|
| 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, [/](#)