

# Package: BiocVersion (via r-universe)

June 17, 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 (>= 4.3.0)

**License** Artistic-2.0

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.0.1

**Repository** <https://bioc.r-universe.dev>

**RemoteUrl** <https://github.com/bioc/BiocVersion>

**RemoteRef** HEAD

**RemoteSha** 43c716aa5d110bc2ac42053033edde0fce3fe9cf

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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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