

[Home](#) » [BiocViews](#) ← **Rule 2**

All Packages

Bioconductor version 3.12 (Development)

Packages found under Software:

Rank based on number of downloads: lower numbers are more frequently downloaded.

Show entries

Search table:

Package	Maintainer	Title	Rank
BiocVersion	Bioconductor Package Maintainer	Set the appropriate version of Bioconductor packages	1
BiocGenerics	Bioconductor Package Maintainer	S4 generic functions used in Bioconductor	2
S4Vectors	Bioconductor Package Maintainer	Foundation of vector-like and list-like containers in Bioconductor	3
IRanges	Bioconductor Package Maintainer	Foundation of integer range manipulation in Bioconductor	4
Biobase	Bioconductor Package Maintainer	Biobase: Base functions for Bioconductor	5
zlibbioc	Bioconductor Package Maintainer	An R packaged zlib-1.2.5	6
AnnotationDbi	Bioconductor Package Maintainer	Manipulation of SQLite-based annotations in Bioconductor	7
BiocParallel	Bioconductor Package Maintainer	Bioconductor facilities for parallel evaluation	8
XVector	Hervé Pagès	Foundation of external vector representation and manipulation in Bioconductor	9

☐ Developers: check this box to toggle the visibility of childless biocViews.

Autocomplete biocViews search:

▼ Software (1897)

- ▶ AssayDomain (767)
- ▶ BiologicalQuestion (789)
- ▶ Infrastructure (433)
- ▶ ResearchField (856)
- ▶ StatisticalMethod (698)
- ▶ Technology (1205)
- ▶ WorkflowStep (1039)
- ▶ AnnotationData (965)
- ▶ ExperimentData (394)
- ▶ Workflow (27)