

Package ‘crisprVerse’

January 23, 2026

Title Easily install and load the crisprVerse ecosystem for CRISPR
gRNA design

Version 1.13.1

Description The crisprVerse is a modular ecosystem of R packages developed for the design and manipulation of CRISPR guide RNAs (gRNAs). All packages share a common language and design principles. This package is designed to make it easy to install and load the crisprVerse packages in a single step. To learn more about the crisprVerse, visit <<https://www.github.com/crisprVerse>>.

Depends R (>= 4.2.0)

Imports BiocManager, cli, crisprBase, crisprBowtie, crisprScore, crisprScoreData, crisprDesign, crisprViz, rlang, tools, utils

Suggests BiocStyle, knitr, testthat

biocViews CRISPR, FunctionalGenomics, GeneTarget

License MIT + file LICENSE

Encoding UTF-8

RoxygenNote 7.3.3

VignetteBuilder knitr

BugReports <https://github.com/crisprVerse/crisprVerse/issues>

URL <https://github.com/crisprVerse/crisprVerse>

git_url <https://git.bioconductor.org/packages/crisprVerse>

git_branch devel

git_last_commit 4ea8be4

git_last_commit_date 2025-11-21

Repository Bioconductor 3.23

Date/Publication 2026-01-23

Author Jean-Philippe Fortin [aut, cre]

Maintainer Jean-Philippe Fortin <fortin946@gmail.com>

Contents

crisprVerse-package	2
crisprVerse_deps	2
crisprVerse_packages	3
crisprVerse_update	3

Index	5
--------------	----------

crisprVerse-package	<i>crisprVerse: Easily install and load the crisprVerse ecosystem for CRISPR gRNA design</i>
---------------------	--

Description

The crisprVerse is a modular ecosystem of R packages developed for the design and manipulation of CRISPR guide RNAs (gRNAs). All packages share a common language and design principles. This package is designed to make it easy to install and load the crisprVerse packages in a single step. To learn more about the crisprVerse, visit <https://www.github.com/crisprVerse>.

Author(s)

Maintainer: Jean-Philippe Fortin <fortin946@gmail.com>

See Also

Useful links:

- <https://github.com/crisprVerse/crisprVerse>
- Report bugs at <https://github.com/crisprVerse/crisprVerse/issues>

crisprVerse_deps	<i>List all crisprVerse dependencies</i>
------------------	--

Description

List all crisprVerse dependencies

Usage

```
crisprVerse_deps(recursive = FALSE)
```

Arguments

`recursive` If TRUE, will also list all dependencies of crisprVerse packages.

Value

A data.frame describing crisprVerse dependencies

Examples

```
if (interactive()){  
  crisprVerse_deps()  
}
```

crisprVerse_packages *List all packages in the crisprVerse*

Description

List all packages in the crisprVerse

Usage

```
crisprVerse_packages(include_self = TRUE)
```

Arguments

include_self Include crisprVerse in the list?

Value

A character vector of packages in the crisprVerse

Examples

```
crisprVerse_packages()
```

crisprVerse_update *Update crisprVerse packages*

Description

This will check to see if all crisprVerse packages (and optionally, their dependencies) are up-to-date, and will install after an interactive confirmation.

Usage

```
crisprVerse_update(recursive = FALSE)
```

Arguments

recursive If TRUE, will also list all dependencies of crisprVerse packages.

Value

Print messages to console as a side effect.

Examples

```
if (interactive()){  
  crisprVerse_update()  
}
```

Index

* **internal**

 crisprVerse-package, [2](#)

crisprVerse (crisprVerse-package), [2](#)

crisprVerse-package, [2](#)

crisprVerse_deps, [2](#)

crisprVerse_packages, [3](#)

crisprVerse_update, [3](#)