

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

[plyranges](#)

RRID:SCR_021324

Type: Tool

Proper Citation

plyranges (RRID:SCR_021324)

Resource Information

URL: <https://bioconductor.org/packages/plyranges/>

Proper Citation: plyranges (RRID:SCR_021324)

Description: Open source software R package for interacting with common Bioconductor classes Ranges and GenomicRanges. Grammar of genomic data transformation. Provides grammatical and consistent way of manipulating these classes to increase accessibility for new Bioconductor users.

Resource Type: software toolkit, software resource

Defining Citation: [DOI:10.1186/s13059-018-1597-8](https://doi.org/10.1186/s13059-018-1597-8)

Keywords: genomic data transformation, genomic data transformation grammar, genomic data, data transformation,

Availability: Free, Available for download, Freely available

Resource Name: plyranges

Resource ID: SCR_021324

Alternate URLs: <https://github.com/sa-lee/plyranges>

License: Artistic 2.0

Record Creation Time: 20220129T080354+0000

Record Last Update: 20260810T040802+0000

Ratings and Alerts

No rating or validation information has been found for plyranges.

No alerts have been found for plyranges.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Lehle JD, et al. (2024) An in vitro approach reveals molecular mechanisms underlying endocrine disruptor-induced epimutagenesis. eLife, 13.