

Resource Summary Report

Generated by [dkNET](#) on Aug 21, 2026

[bedr](#)

RRID:SCR_024526

Type: Tool

Proper Citation

bedr (RRID:SCR_024526)

Resource Information

URL: <https://cran.r-project.org/web/packages/bedr/vignettes/Using-bedr.html>

Proper Citation: bedr (RRID:SCR_024526)

Description: Software R package for genomic interval processing.Genomic regions processing using open-source command line tools to perform genome arithmetic e.g indexing, formatting and merging.

Resource Type: software resource, software toolkit

Keywords: genomic regions processing

Availability: Free, Available for download, Freely available

Resource Name: bedr

Resource ID: SCR_024526

Alternate URLs: <https://CRAN.R-project.org/package=bedr>

License: GNU GPL v2

Record Creation Time: 20231002T161337+0000

Record Last Update: 20260821T194302+0000

Ratings and Alerts

No rating or validation information has been found for bedr.

No alerts have been found for bedr.

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 [dkNET](#) .

Vu HTH, et al. (2023) Unsupervised contrastive peak caller for ATAC-seq. Genome research, 33(7), 1133.