

Resource Summary Report

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bulkAnalyseR

RRID:SCR_027647

Type: Tool

Proper Citation

bulkAnalyseR (RRID:SCR_027647)

Resource Information

URL: <https://github.com/Core-Bioinformatics/bulkAnalyseR>

Proper Citation: bulkAnalyseR (RRID:SCR_027647)

Description: Software R package for most bulk sequencing datasets. Creates shiny app for interactive data analysis and visualisation. Used for analysing and sharing bulk sequencing results.

Resource Type: software resource, software toolkit

Defining Citation: [PMID:36583521](#)

Keywords: bulk sequencing results, analysing and sharing results, sequencing data,

Funding: Wellcome Trust

Availability: Free, Available for download, Freely available

Resource Name: bulkAnalyseR

Resource ID: SCR_027647

Alternate URLs: <https://cran.r-project.org/web/packages/bulkAnalyseR/>

License: GNU GPL v3

Record Creation Time: 20251111T060300+0000

Record Last Update: 20260808T190816+0000

Ratings and Alerts

No rating or validation information has been found for bulkAnalyseR.

No alerts have been found for bulkAnalyseR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.