

Resource Summary Report

Generated by [Kravitz](#) on Sep 11, 2026

Rdisop

RRID:SCR_000453

Type: Tool

Proper Citation

Rdisop (RRID:SCR_000453)

Resource Information

URL: <http://bioconductor.org/packages/release/bioc/html/Rdisop.html>

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Description: Software for identification of metabolites using high precision mass spectrometry. MS Peaks are used to derive a ranked list of sum formulae, alternatively for a given sum formula the theoretical isotope distribution can be calculated to search in MS peak lists.

Synonyms: Rdisop - Decomposition of Isotopic Patterns, Rdisop: Decomposition of Isotopic Patterns, Decomposition of Isotopic Patterns

Resource Type: software resource

Keywords: standalone software, mac os x, unix/linux, windows, r, mass spectrometry, metabolomics, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: Rdisop

Resource ID: SCR_000453

Alternate IDs: OMICS_02416, biotools:rdisop

Alternate URLs: <https://github.com/sneumann/Rdisop>, <https://bio.tools/rdisop>

License: GNU General Public License, v2

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260905T132419+0000

Ratings and Alerts

No rating or validation information has been found for Rdisop.

No alerts have been found for Rdisop.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.