

Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

enviPick

RRID:SCR_003059

Type: Tool

Proper Citation

enviPick (RRID:SCR_003059)

Resource Information

URL: <http://cran.r-project.org/web/packages/enviPick/>

Proper Citation: enviPick (RRID:SCR_003059)

Description: Software for sequential partitioning, clustering and peak detection of centroided LC-MS mass spectrometry data (.mzXML). Interactive result and raw data plot.

Synonyms: enviPick: Peak picking for high resolution mass spectrometry data

Resource Type: software resource, web site

Keywords: standalone software, mac os x, unix/linux, windows, r

Availability: Free, Available for download, Freely available

Resource Name: enviPick

Resource ID: SCR_003059

Alternate IDs: OMICS_05010

Alternate URLs: <https://rdr.io/cran/enviPick/>

Record Creation Time: 20220129T080216+0000

Record Last Update: 20260912T200229+0000

Ratings and Alerts

No rating or validation information has been found for enviPick.

No alerts have been found for enviPick.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Ani?i? N, et al. (2023) Functional iridoid synthases from iridoid producing and non-producing *Nepeta* species (subfam. Nepetoideae, fam. Lamiaceae). *Frontiers in plant science*, 14, 1211453.

Peris-Díaz MD, et al. (2019) R-MetaboList 2: A Flexible Tool for Metabolite Annotation from High-Resolution Data-Independent Acquisition Mass Spectrometry Analysis. *Metabolites*, 9(9).