## **Resource Summary Report**

Generated by dkNET on Apr 30, 2025

# 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: web site, software resource

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

**Funding:** 

Availability: GNU General Public License, v2, v3

Resource Name: enviPick

Resource ID: SCR\_003059

Alternate IDs: OMICS\_05010

**Record Creation Time:** 20220129T080216+0000

Record Last Update: 20250429T054818+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 dkNET.

Ani?i? N, et al. (2023) Functional iridoid synthases from iridoid producing and non-producing Nepeta species (subfam. Nepetoidae, 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).