Resource Summary Report

Generated by NIF on May 15, 2025

flowPeaks

RRID:SCR_000407
Type: Tool

Proper Citation

flowPeaks (RRID:SCR_000407)

Resource Information

URL: https://github.com/yongchao/flowPeaks

Proper Citation: flowPeaks (RRID:SCR_000407)

Description: Software for fast and automatic clustering to classify the cells into subpopulations based on finding the peaks from the overall density function generated by K-means.

Resource Type: software resource

Defining Citation: PMID:22595209

Keywords: software package, mac os x, unix/linux, windows, r, clustering, flow cytometry, gating, bio.tools

Funding:

Availability: Artistic License, V1

Resource Name: flowPeaks

Resource ID: SCR_000407

Alternate IDs: biotools:flowpeaks, OMICS_05604

Alternate URLs: http://www.bioconductor.org/packages/devel/bioc/html/flowPeaks.html, https://bio.tools/flowpeaks

Record Creation Time: 20220129T080201+0000

Record Last Update: 20250420T013945+0000

Ratings and Alerts

No rating or validation information has been found for flowPeaks.

No alerts have been found for flowPeaks.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.