

Resource Summary Report

Generated by SPARC SAWG on Sep 18, 2026

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](https://pubmed.ncbi.nlm.nih.gov/22595209/)

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

Availability: Free, Available for download, Freely available

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>

License: Artistic License, V1

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260912T195508+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.