

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 20, 2026

## [flowPeaks](#)

RRID:SCR\_000407

Type: Tool

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### Proper Citation

flowPeaks (RRID:SCR\_000407)

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### 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

**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:** 20260919T194922+0000

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### Ratings and Alerts

No rating or validation information has been found for flowPeaks.

No alerts have been found for flowPeaks.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.