**flowPeaks**

RRID:SCR_000407  
Type: Tool

**Proper Citation**

flowPeaks (RRID:SCR_000407)

**Resource Information**

**URL:** [https://github.com/yongchao/flowPeaks](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://www.ncbi.nlm.nih.gov/pubmed/22595209)

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

**Availability:** Artistic License, V1

**Resource Name:** flowPeaks

**Resource ID:** SCR_000407

**Alternate IDs:** OMICS_05604, biotools:flowpeaks


**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.