

Resource Summary Report

Generated by [Kravitz](#) on Sep 17, 2026

[flowMerge](#)

RRID:SCR_002224

Type: Tool

Proper Citation

flowMerge (RRID:SCR_002224)

Resource Information

URL: <http://www.bioconductor.org/packages/release/bioc/html/flowMerge.html>

Proper Citation: flowMerge (RRID:SCR_002224)

Description: Software for merging of mixture components for model-based automated gating of flow cytometry data using the flowClust framework.

Synonyms: flowMerge - Cluster Merging for Flow Cytometry Data

Resource Type: software resource

Defining Citation: [PMID:20049161](#)

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

Availability: Free, Available for download, Freely available

Resource Name: flowMerge

Resource ID: SCR_002224

Alternate IDs: OMICS_05605

License: Artistic License, v2

Record Creation Time: 20220129T080212+0000

Record Last Update: 20260912T195538+0000

Ratings and Alerts

No rating or validation information has been found for flowMerge.

No alerts have been found for flowMerge.

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 [Kravitz](#) .

Chen Y, et al. (2024) Benchmarking digital PCR partition classification methods with empirical and simulated duplex data. Briefings in bioinformatics, 25(3).

Rahim A, et al. (2018) High throughput automated analysis of big flow cytometry data. Methods (San Diego, Calif.), 134-135, 164.