# **Resource Summary Report**

Generated by <u>dkNET</u> on May 17, 2025

# **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

Funding:

Availability: Artistic License, v2

Resource Name: flowMerge

Resource ID: SCR\_002224

Alternate IDs: OMICS\_05605

Record Creation Time: 20220129T080212+0000

Record Last Update: 20250420T014055+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 <u>dkNET</u>.

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.