

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

flowPhyto

RRID:SCR_002183

Type: Tool

Proper Citation

flowPhyto (RRID:SCR_002183)

Resource Information

URL: <https://bioconductor.org/packages/2.11/bioc/html/flowPhyto.html>

Proper Citation: flowPhyto (RRID:SCR_002183)

Description: An R package that performs aggregate statistics on virtually unlimited collections of raw flow cytometry files and provides a memory efficient, parallelized solution for analyzing high-throughput flow cytometric data.

Synonyms: flowPhyto - Methods for Continuous Flow Cytometry

Resource Type: software resource

Defining Citation: [PMID:21208987](#)

Keywords: software package, mac os x, unix/linux, windows, r, classification, clustering, data import, flow cytometry, quality control, visualization

Availability: Free, Available for download, Freely available

Resource Name: flowPhyto

Resource ID: SCR_002183

Alternate IDs: OMICS_05606

Old URLs: <http://www.bioconductor.org/packages/release/bioc/html/flowPhyto.html>

License: Artistic License, v2

Record Creation Time: 20220129T080212+0000

Record Last Update: 20260725T190518+0000

Ratings and Alerts

No rating or validation information has been found for flowPhyto.

No alerts have been found for flowPhyto.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.