

# Resource Summary Report

Generated by [kravitz2](#) on Jul 30, 2026

## flowPhyto

RRID:SCR\_002183

Type: Tool

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### Proper Citation

flowPhyto (RRID:SCR\_002183)

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### Resource Information

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

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**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

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### Ratings and Alerts

No rating or validation information has been found for flowPhyto.

No alerts have been found for flowPhyto.

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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.