

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

FluxModeCalculator

RRID:SCR_016290

Type: Tool

Proper Citation

FluxModeCalculator (RRID:SCR_016290)

Resource Information

URL: <https://omictools.com/fluxmodecalculator-tool>

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Description: Software for performing flux mode analysis in stoichiometric models. FluxModeCalculator enables large-scale elementary flux mode (EFM) computation and uses the OpenMP API to optimally exploit processor architectures with multiple cores., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Synonyms: Flux Mode Calculator

Resource Type: data analysis software, data processing software, software application, software resource

Defining Citation: [PMID:26685305](#)

Keywords: efm, flux, stoichiometry, algorithm, matlab, linux, model, magnitude, bio.tools

Funding: Biobanking and Biomolecular Resources Research Infrastructure (BBMRI) ; CardioVascular Research Netherlands (CVON-ENERGISE) ; Center of Medical Systems Biology (CMSB) ; European Network for Genetic and Genomic Epidemiology (ENGAGE) ; Netherlands Consortium for Systems Biology (NCSB)

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: FluxModeCalculator

Resource ID: SCR_016290

Alternate IDs: biotools:fluxmodecalculator, OMICS_10894

Alternate URLs: <https://bio.tools/fluxmodecalculator>

License: GNU General Public License version 3.0

Record Creation Time: 20220129T080329+0000

Record Last Update: 20260905T132806+0000

Ratings and Alerts

No rating or validation information has been found for FluxModeCalculator .

No alerts have been found for FluxModeCalculator .

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [Kravitz](#) .

Averesch NJH, et al. (2019) Metabolic engineering of Bacillus subtilis for production of para-aminobenzoic acid - unexpected importance of carbon source is an advantage for space application. Microbial biotechnology, 12(4), 703.