

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

Fluxfix

RRID:SCR_026259

Type: Tool

Proper Citation

Fluxfix (RRID:SCR_026259)

Resource Information

URL: <https://fluxfix.science/>

Proper Citation: Fluxfix (RRID:SCR_026259)

Description: Web isotopologue analysis tool for metabolomics experiment. Used to compute tracer analysis. Transforms raw mass spec AUC values into the percent representation of each isotopologue measured.

Resource Type: data access protocol, software resource, web service, source code

Keywords: transform raw mass spec AUC values, percent representation, isotopologue measure,

Availability: Free, Freely available

Resource Name: Fluxfix

Resource ID: SCR_026259

Alternate URLs: <https://github.com/sophietrefely/traceranalysis>

Record Creation Time: 20250110T053324+0000

Record Last Update: 20260815T042924+0000

Ratings and Alerts

No rating or validation information has been found for Fluxfix.

No alerts have been found for Fluxfix.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Sharma G, et al. (2025) IGF2BP3 redirects glycolytic flux to promote one-carbon metabolism and RNA methylation. Cell reports, 116330.

Ball AB, et al. (2025) Pro-inflammatory macrophage activation does not require inhibition of oxidative phosphorylation. EMBO reports, 26(4), 982.

Jones AE, et al. (2024) The metabolic cofactor Coenzyme A enhances alternative macrophage activation via MyD88-linked signaling. bioRxiv : the preprint server for biology.