Resource Summary Report

Generated by RRID on Aug 6, 2026

METAFlux

RRID:SCR_028022
Type: Tool

Proper Citation

METAFlux (RRID:SCR_028022)

Resource Information

URL: https://github.com/KChen-lab/METAFlux?tab=readme-ov-file

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Description: Software tool that predicts cancer metabolic fluxes from bulk RNA-seq and scRNA-seq data to address these analytic gaps. Used for characterizing metabolic circuits and output non-degenerative fluxes using cancer gene expression data.

Resource Type: software application, source code, software resource

Defining Citation: PMID:37573313

Keywords: predict cancer metabolic fluxes, RNA-seq data, scRNA-seq data, characterizing metabolic circuits, non-degenerative fluxes, cancer gene expression data,

Funding: NCI U01CA247760;
NCI P30 CA016672;
Chan Zuckerberg Initiative ;
Silicon Valley Community Foundation ;
MD Anderson Moonshot programs

Availability: Free, Available for download, Freely available

Resource Name: METAFlux

Resource ID: SCR_028022

License: MIT license

Record Creation Time: 20260225T055938+0000

Record Last Update: 20260806T054850+0000

Ratings and Alerts

No rating or validation information has been found for METAFlux.

No alerts have been found for METAFlux.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.