

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 7, 2026

[metabnorm](#)

RRID:SCR_001266

Type: Tool

Proper Citation

metabnorm (RRID:SCR_001266)

Resource Information

URL: <http://sourceforge.net/projects/metabnorm/>

Proper Citation: metabnorm (RRID:SCR_001266)

Description: Software tool as mixed model normalization method for metabolomics data. Uses normalization approach based on mixed model, with simultaneous estimation of correlation matrix.

Resource Type: standalone software, software application, software resource

Defining Citation: [PMID:24711654](#)

Keywords: Metabolomics datasets, corelation, normalization, identifying metabolites, bio.tools

Funding: Cancer Research UK Cambridge Institute ;
Erik and Edith Fernstr m foundation ;
Cancer Research UK

Availability: Free, Available for download, Freely available

Resource Name: metabnorm

Resource ID: SCR_001266

Alternate IDs: OMICS_03548, biotools:metabnorm

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

License: GNU General Public License

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260807T042522+0000

Ratings and Alerts

No rating or validation information has been found for metabnorm.

No alerts have been found for metabnorm.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.