## **Resource Summary Report**

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# 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: software resource, software application, standalone software

Defining Citation: PMID:24711654

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

**Funding:** Cancer Research UK Cambridge Institute ; Erik and Edith Fernström 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: 20250519T204853+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: <u>SciCrunch Registry</u>

### **Usage and Citation Metrics**

We have not found any literature mentions for this resource.