

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

MetAssign

RRID:SCR_000092

Type: Tool

Proper Citation

MetAssign (RRID:SCR_000092)

Resource Information

URL: <https://mzmatch.sourceforge.net/MetAssign.php>

Proper Citation: MetAssign (RRID:SCR_000092)

Description: Software that combines information from the mass-to-charge ratio, retention time and intensity of each peak, together with a model of the inter-peak dependency structure, to increase the accuracy of peak annotation. The software has been implemented as part of the mzMatch metabolomics analysis pipeline, which is available for download.

Resource Type: software application, software resource, standalone software

Defining Citation: [PMID:24916385](#)

Keywords: peak annotation, peaks probabilistic identifications,

Availability: Free, Available for download, Freely available

Resource Name: MetAssign

Resource ID: SCR_000092

Alternate IDs: OMICS_04679

Old URLs: <http://mzmatch.sourceforge.net/>

Record Creation Time: 20220129T080159+0000

Record Last Update: 20260727T040243+0000

Ratings and Alerts

No rating or validation information has been found for MetAssign.

No alerts have been found for MetAssign.

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 [nidm-terms](#) .

Daskalaki E, et al. (2015) A study of the effects of exercise on the urinary metabolome using normalisation to individual metabolic output. *Metabolites*, 5(1), 119.