

# Resource Summary Report

Generated by [dkNET](#) on Aug 25, 2026

## MetAssign

RRID:SCR\_000092

Type: Tool

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### Proper Citation

MetAssign (RRID:SCR\_000092)

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### 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:** 20260821T193607+0000

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### Ratings and Alerts

No rating or validation information has been found for MetAssign.

No alerts have been found for MetAssign.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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.