

# Resource Summary Report

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

## [metaMA](#)

RRID:SCR\_000408

Type: Tool

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### Proper Citation

metaMA (RRID:SCR\_000408)

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### Resource Information

**URL:** <http://cran.r-project.org/web/packages/metaMA/>

**Proper Citation:** metaMA (RRID:SCR\_000408)

**Description:** Software R package for meta-analysis for microarrays. It combines either p-values or modified effect sizes from different studies to find differentially expressed genes.

**Synonyms:** Meta-analysis for MicroArrays, Meta analysis for MicroArrays

**Resource Type:** software resource, software toolkit

**Defining Citation:** [PMID:19628502](#)

**Keywords:** standalone software, mac os x, unix/linux, windows, r

**Funding:** Scottish Government Rural and Environment Research and Analysis Directorate

**Availability:** Free, Available for download, Freely available

**Resource Name:** metaMA

**Resource ID:** SCR\_000408

**Alternate IDs:** OMICS\_03524

**License:** GNU General Public License, v2, v3

**Record Creation Time:** 20220129T080201+0000

**Record Last Update:** 20260821T193613+0000

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### Ratings and Alerts

No rating or validation information has been found for metaMA.

No alerts have been found for metaMA.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.