

Resource Summary Report

Generated by [NIF](#) on Sep 12, 2026

[metaMA](#)

RRID:SCR_000408

Type: Tool

Proper Citation

metaMA (RRID:SCR_000408)

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: 20260905T133221+0000

Ratings and Alerts

No rating or validation information has been found for metaMA.

No alerts have been found for metaMA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.