

Resource Summary Report

Generated by [RRID](#) on Aug 8, 2026

[MetaDE](#)

RRID:SCR_000199

Type: Tool

Proper Citation

MetaDE (RRID:SCR_000199)

Resource Information

URL: <https://cran.r-project.org/src/contrib/Archive/MetaDE/>

Proper Citation: MetaDE (RRID:SCR_000199)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on August 30, 2022. Software package that implements 12 major meta-analysis methods for differential expression analysis. Package was removed from the CRAN repository. Formerly available versions can be obtained from the archive. Archived on 2018-01-23 as check problems were not corrected in time.

Synonyms: MetaDE: Microarray meta-analysis for differentially expressed gene detection

Resource Type: software resource

Defining Citation: [PMID:22863766](#)

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

Availability: Free, Available for download, Freely available

Resource Name: MetaDE

Resource ID: SCR_000199

Alternate IDs: OMICS_04033

Old URLs: <http://cran.r-project.org/web/packages/MetaDE/>

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260808T185716+0000

Ratings and Alerts

No rating or validation information has been found for MetaDE.

No alerts have been found for MetaDE.

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 [RRID](#) .

Smith AR, et al. (2016) Increased DNA methylation near TREM2 is consistently seen in the superior temporal gyrus in Alzheimer's disease brain. *Neurobiology of aging*, 47, 35.