Resource Summary Report

Generated by NIF on May 25, 2025

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

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE.

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: 20250525T030527+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 NIF.

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.