

# Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

## DEMI

RRID:SCR\_002291

Type: Tool

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### Proper Citation

DEMI (RRID:SCR\_002291)

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### Resource Information

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

**Proper Citation:** DEMI (RRID:SCR\_002291)

**Description:** R package for estimating differential expression from multiple indicators that capitalizes on the high number of concurrent measurements. It extends to various experimental designs and target categories (transcripts, genes, genomic regions) as well as small sample sizes.

**Synonyms:** demi: Differential Expression from Multiple Indicators, Differential Expression from Multiple Indicators

**Resource Type:** software resource

**Defining Citation:** [PMID:24586062](#)

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

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

**Resource Name:** DEMI

**Resource ID:** SCR\_002291

**Alternate IDs:** OMICS\_03438

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

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

**Record Creation Time:** 20220129T080212+0000

**Record Last Update:** 20260725T190523+0000

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### Ratings and Alerts

No rating or validation information has been found for DEMI.

No alerts have been found for DEMI.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Huang Y, et al. (2024) Malware Identification Method in Industrial Control Systems Based on Opcode2vec and CVAE-GAN. Sensors (Basel, Switzerland), 24(17).

Ruggeri K, et al. (2022) The globalizability of temporal discounting. Nature human behaviour, 6(10), 1386.

Pezone R, et al. (2022) Sensitive Transfer-Free Wafer-Scale Graphene Microphones. ACS applied materials & interfaces, 14(18), 21705.

Davoudi P, et al. (2022) Genome-wide detection of copy number variation in American mink using whole-genome sequencing. BMC genomics, 23(1), 649.

Karimi K, et al. (2021) Population Genomics of American Mink Using Whole Genome Sequencing Data. Genes, 12(2).

Muškoví? M, et al. (2021) Photodynamic Inactivation of Legionella pneumophila Biofilm Formation by Cationic Tetra- and Tripyridylporphyrins in Waters of Different Hardness. International journal of molecular sciences, 22(16).