### **Resource Summary Report**

Generated by dkNET on Apr 24, 2025

# **lapmix**

RRID:SCR\_001347

Type: Tool

### **Proper Citation**

lapmix (RRID:SCR\_001347)

#### **Resource Information**

URL: http://www.bioconductor.org/packages/release/bioc/html/lapmix.html

**Proper Citation:** lapmix (RRID:SCR\_001347)

**Description:** Software to identify differentially expressed genes. A hierarchical Bayesian

approach is used, and the hyperparameters are estimated using empirical Bayes.

**Abbreviations:** lapmix

Synonyms: Laplace Mixture Model in Microarray Experiments

Resource Type: software resource

**Keywords:** differential expression, microarray, bio.tools

**Funding:** 

Availability: GNU General Public License, v2 or newer

Resource Name: lapmix

Resource ID: SCR\_001347

Alternate IDs: biotools:lapmix, OMICS\_01992

Alternate URLs: https://bio.tools/lapmix

**Record Creation Time:** 20220129T080207+0000

**Record Last Update:** 20250420T014027+0000

### Ratings and Alerts

No rating or validation information has been found for lapmix.

No alerts have been found for lapmix.

### Data and Source Information

Source: SciCrunch Registry

## Usage and Citation Metrics

We have not found any literature mentions for this resource.