LIMMA
RRID:SCR_010943
Type: Tool

Proper Citation

LIMMA (RRID:SCR_010943)

Resource Information

URL: http://bioinf.wehi.edu.au/limma/

Description: Software package for the analysis of gene expression microarray data, especially the use of linear models for analyzing designed experiments and the assessment of differential expression.

Resource Name: LIMMA

Proper Citation: LIMMA (RRID:SCR_010943)

Resource Type: Resource, software resource, software application, data analysis software, data processing software

Keywords: analysis, gene, expression, microarray, data, linear, model

Resource ID: SCR_010943

Related resources: GEO2R, Bioconductor

Availability: Free, Available for download, Freely available

Website Status: Last checked up

Alternate IDs: OMICS_00769

Alternate URLs: https://omictools.com/limma-tool

Abbreviations: LIMMA

Mentions Count: 7362
Ratings and Alerts

No rating or validation information has been found for LIMMA.

No alerts have been found for LIMMA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 7362 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](https://www.fdi-lab.org/scicrunch).


Freihen V, et al. (2020) SNAIL1 employs ?-Catenin-LEF1 complexes to control colorectal cancer cell invasion and proliferation. International journal of cancer, 146(8), 2229-2242.


Niu B, et al. (2020) Glioma stages prediction based on machine learning algorithm combined


