**LIMMA**
RRID:SCR_010943
Type: Tool

**Proper Citation**

LIMMA (RRID:SCR_010943)

**Resource Information**

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

**Proper Citation:** LIMMA (RRID:SCR_010943)

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

**Keywords:** analysis, gene, expression, microarray, data, linear, model, bio.tools

**Related resources:** GEO2R, Bioconductor

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

**Website Status:** Last checked up

**Abbreviations:** LIMMA

**Resource Name:** LIMMA

**Resource ID:** SCR_010943

**Alternate IDs:** OMICS_00769, biotools:limma

**Alternate URLs:** https://omictools.com/limma-tool, https://bio.tools/limma

**Ratings and Alerts**
No rating or validation information has been found for LIMMA.

No alerts have been found for LIMMA.

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**Data and Source Information**

**Source**: [SciCrunch Registry](https://www.sci.crunch.org)

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**Usage and Citation Metrics**

We found 9536 mentions in open access literature.

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


Ma M, et al. (2021) Sustained androgen receptor signaling is a determinant of melanoma cell growth potential and tumorigenesis. The Journal of experimental medicine, 218(2).


Jiang M, et al. (2021) Transcriptional network modulated by the prognostic signature transcription factors and their long noncoding RNA partners in primary prostate cancer. EBioMedicine, 63, 103150.