

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 18, 2026

ExpressionPlot

RRID:SCR_001904

Type: Tool

Proper Citation

ExpressionPlot (RRID:SCR_001904)

Resource Information

URL: <http://blog.expressionplot.com/>

Proper Citation: ExpressionPlot (RRID:SCR_001904)

Description: Software package consisting of a default back end, which prepares raw sequencing or Affymetrix microarray data, and a web-based front end, which offers a biologically centered interface to browse, visualize, and compare different data sets.

Abbreviations: expressionplot

Resource Type: software resource

Defining Citation: [PMID:21797991](#)

Keywords: analysis, rna-seq, microarray, gene expression, affymetrix, prototype

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ExpressionPlot

Resource ID: SCR_001904

Alternate IDs: OMICS_01939

Record Creation Time: 20220129T080210+0000

Record Last Update: 20260912T195533+0000

Ratings and Alerts

No rating or validation information has been found for ExpressionPlot.

No alerts have been found for ExpressionPlot.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

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

Shen K, et al. (2021) Multiple sclerosis risk gene MERTK is required for microglial activation and subsequent remyelination. *Cell reports*, 34(10), 108835.

Yap K, et al. (2018) A Short Tandem Repeat-Enriched RNA Assembles a Nuclear Compartment to Control Alternative Splicing and Promote Cell Survival. *Molecular cell*, 72(3), 525.

Mendelsohn AI, et al. (2017) Divergent Hox Coding and Evasion of Retinoid Signaling Specifies Motor Neurons Innervating Digit Muscles. *Neuron*, 93(4), 792.

Collins KM, et al. (2017) An RRM-ZnF RNA recognition module targets RBM10 to exonic sequences to promote exon exclusion. *Nucleic acids research*, 45(11), 6761.

Hamid FM, et al. (2017) A mechanism underlying position-specific regulation of alternative splicing. *Nucleic acids research*, 45(21), 12455.

Aaker JD, et al. (2016) Transcriptional Fingerprint of Hypomyelination in Zfp191null and Shiverer (Mbpshi) Mice. *ASN neuro*, 8(5).

Yap K, et al. (2016) Polarizing the Neuron through Sustained Co-expression of Alternatively Spliced Isoforms. *Cell reports*, 15(6), 1316.

Kolumam G, et al. (2015) Sustained Brown Fat Stimulation and Insulin Sensitization by a Humanized Bispecific Antibody Agonist for Fibroblast Growth Factor Receptor 1/Klotho Complex. *EBioMedicine*, 2(7), 730.

Bai Y, et al. (2015) IRcall and IRclassifier: two methods for flexible detection of intron retention events from RNA-Seq data. *BMC genomics*, 16 Suppl 2(Suppl 2), S9.