

Resource Summary Report

Generated by [NIF](#) on Aug 30, 2026

JMP Genomics

RRID:SCR_011857

Type: Tool

Proper Citation

JMP Genomics (RRID:SCR_011857)

Resource Information

URL: <http://www.jmp.com/software/genomics/>

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Description: Provides the tools you need to analyze rare and common variants, detect differential expression patterns, discover reliable biomarker profiles, and incorporate pathway information into your analysis workflows.

Abbreviations: JMP Genomics

Resource Type: commercial organization, software resource

Availability: Commercial license

Resource Name: JMP Genomics

Resource ID: SCR_011857

Alternate IDs: OMICS_01129

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260829T183222+0000

Ratings and Alerts

No rating or validation information has been found for JMP Genomics.

No alerts have been found for JMP Genomics.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Salazar-Noratto GE, et al. (2023) Patient-Specific iPSC-Derived Models Link Aberrant Endoplasmic Reticulum Stress Sensing and Response to Juvenile Osteochondritis Dissecans Etiology. *Stem cells translational medicine*, 12(5), 293.

Mashima H, et al. (2021) Improved safety of induced pluripotent stem cell-derived antigen-presenting cell-based cancer immunotherapy. *Molecular therapy. Methods & clinical development*, 21, 171.

Al-Sayegh M, et al. (2021) Mouse Embryonic Fibroblast Adipogenic Potential: A Comprehensive Transcriptome Analysis. *Adipocyte*, 10(1), 1.

Mashima H, et al. (2020) Generation of GM-CSF-producing antigen-presenting cells that induce a cytotoxic T cell-mediated antitumor response. *Oncoimmunology*, 9(1), 1814620.

Al-Sayegh MA, et al. (2020) ??-actin contributes to open chromatin for activation of the adipogenic pioneer factor CEBPA during transcriptional reprogramming. *Molecular biology of the cell*, 31(23), 2511.

Muleta KT, et al. (2017) Loci associated with resistance to stripe rust (*Puccinia striiformis* f. sp. *tritici*) in a core collection of spring wheat (*Triticum aestivum*). *PloS one*, 12(6), e0179087.

Koh S, et al. (2015) Human Umbilical Tissue-Derived Cells Promote Synapse Formation and Neurite Outgrowth via Thrombospondin Family Proteins. *The Journal of neuroscience : the official journal of the Society for Neuroscience*, 35(47), 15649.

Dissanayake SN, et al. (2010) aeGEPUCI: a database of gene expression in the dengue vector mosquito, *Aedes aegypti*. *BMC research notes*, 3, 248.

Grigorova M, et al. (2008) FSHB promoter polymorphism within evolutionary conserved element is associated with serum FSH level in men. *Human reproduction (Oxford, England)*, 23(9), 2160.

Tempelman RJ, et al. (2008) Statistical analysis of efficient unbalanced factorial designs for two-color microarray experiments. *International journal of plant genomics*, 2008, 584360.

Page GP, et al. (2008) Bioinformatic tools for inferring functional information from plant microarray data: tools for the first steps. *International journal of plant genomics*, 2008, 147563.