## **Resource Summary Report**

Generated by dkNET on May 5, 2025

# **JMP Genomics**

RRID:SCR\_011857

Type: Tool

### **Proper Citation**

JMP Genomics (RRID:SCR\_011857)

#### **Resource Information**

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

**Proper Citation:** JMP Genomics (RRID:SCR\_011857)

**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

**Funding:** 

Availability: Commercial license

Resource Name: JMP Genomics

Resource ID: SCR\_011857

Alternate IDs: OMICS\_01129

Record Creation Time: 20220129T080307+0000

Record Last Update: 20250505T054122+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 dkNET.

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 antigenpresenting 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 reprograming. 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.