Resource Summary Report

Generated by dkNET on Apr 28, 2025

Genedata Expressionist

RRID:SCR_003298

Type: Tool

Proper Citation

Genedata Expressionist (RRID:SCR_003298)

Resource Information

URL: http://www.genedata.com/products/expressionist/genomic-profiling.html

Proper Citation: Genedata Expressionist (RRID:SCR_003298)

Description: Software that provides data processing, analysis, management, and reporting of metabolomics, proteomics and biotherapeutics characterization studies based on mass spectrometry. It can process raw data from various MS instruments, serve MS processing, analysis and reporting needs, and ensure reproducibility and traceability of results.

Synonyms: Genedata Expressionist for Genomic Profiling

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

Keywords: mass spectrometry data processing, mass spectrometry data analysis, mass spectrometry data management, omics software

Funding:

Availability: Available for purchase

Resource Name: Genedata Expressionist

Resource ID: SCR_003298

Alternate IDs: OMICS_00761

Record Creation Time: 20220129T080218+0000

Record Last Update: 20250428T053029+0000

Ratings and Alerts

No rating or validation information has been found for Genedata Expressionist.

No alerts have been found for Genedata Expressionist.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 7 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Seinen J, et al. (2021) Sputum Proteome Signatures of Mechanically Ventilated Intensive Care Unit Patients Distinguish Samples with or without Anti-pneumococcal Activity. mSystems, 6(2).

Rangel Pedersen N, et al. (2021) Multicomponent carbohydrase system from Trichoderma reesei: A toolbox to address complexity of cell walls of plant substrates in animal feed. PloS one, 16(6), e0251556.

Mor N, et al. (2018) Neutralizing Gatad2a-Chd4-Mbd3/NuRD Complex Facilitates Deterministic Induction of Naive Pluripotency. Cell stem cell, 23(3), 412.

Yoon S, et al. (2017) MLKL, the Protein that Mediates Necroptosis, Also Regulates Endosomal Trafficking and Extracellular Vesicle Generation. Immunity, 47(1), 51.

Shigemura T, et al. (2016) Novel heterozygous C243Y A20/TNFAIP3 gene mutation is responsible for chronic inflammation in autosomal-dominant Behçet's disease. RMD open, 2(1), e000223.

Bajaj I, et al. (2014) Functional characterization of a Penicillium chrysogenum mutanase gene induced upon co-cultivation with Bacillus subtilis. BMC microbiology, 14, 114.

Marshall E, et al. (2011) In silico analysis identifies a novel role for androgens in the regulation of human endometrial apoptosis. The Journal of clinical endocrinology and metabolism, 96(11), E1746.