Resource Summary Report

Generated by RRID on Aug 15, 2026

Progenesis QI

RRID:SCR_018923
Type: Tool

Proper Citation

Progenesis QI (RRID:SCR_018923)

Resource Information

URL: http://www.nonlinear.com/progenesis/qi-for-proteomics/

Proper Citation: Progenesis QI (RRID:SCR_018923)

Description: Software tool as next generation in LC-MS proteomics data analysis software by Nonlinear Dynamics.

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

Keywords: Next generation, LC-MS proteomics data, data analysis, Nonlinear Dynamics

Resource Name: Progenesis QI

Resource ID: SCR_018923

Record Creation Time: 20220129T080342+0000

Record Last Update: 20260815T182600+0000

Ratings and Alerts

No rating or validation information has been found for Progenesis QI.

No alerts have been found for Progenesis QI.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1011 mentions in open access literature.

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

Yang J, et al. (2026) Temporal Metabolomics Profiling Reveals Liver Metabolic Control in Single and Repeated Exhaustive Exercise in Murine Models. FASEB journal : official publication of the Federation of American Societies for Experimental Biology, 40(9), e71868.

Wherley TJ, et al. (2026) Myomaker and ether lipids cooperate to promote fusion-competent membrane states. Cell reports, 45(2), 116900.

Ran X, et al. (2026) Multi-omics reveals cholesterol-driven macrophage metabolic reprogramming and inflammation in chronic obstructive pulmonary disease. Genome medicine, 18(1), 17.

Huo R, et al. (2026) Procyanidin Suppresses Tumor Growth by Activating the B-Cell MAPK Pathway through Remodulation of the Gut Microbiota and Metabolites in Hepatocellular Carcinoma. International journal of biological sciences, 22(1), 161.

Bacha AS, et al. (2026) Time-dependent Lactiplantibacillus plantarum (LP) AS21 intervention enhances mesalazine efficacy by modulating gut microbiota and host immunometabolic responses in DSS-induced colitis. Journal of translational medicine, 24(1), 163.

Duan J, et al. (2026) Multi-Omics Analysis Reveals the Potential Preventive Mechanism of Lactobacillus salivarius Li01 Against L-Arginine-Induced Acute Pancreatitis in Mice. Microbial biotechnology, 19(1), e70300.

Kataoka R, et al. (2026) Effects of Aspergillus oryzae-derived rice-koji protein on the sake metabolome. Applied and environmental microbiology, 92(3), e0195525.

Tieze SM, et al. (2026) Deficient de-S-acylation in aging and CLN1 contributes to lyso-mitochondrial dysfunction, lipid dyshomeostasis, and resultant lipofuscin biogenesis. Research square.

Shi W, et al. (2026) Metabolomics Analysis Reveals Gut Microbiota-Associated Sakuranin Modulates Endometrial Stem Cell Differentiation and Inflammation to Alleviate Pain in Endometriosis. Journal of pain research, 19, 557430.

Mao ZA, et al. (2026) Systematic Investigation of Tumor Immune Microenvironment Modulation by Cynomorium songaricum Against Breast Cancer Through Integrated Chemomics, Network Pharmacology and Molecular Docking. Pharmaceuticals (Basel, Switzerland), 19(2).

Wang X, et al. (2026) Integrated multi-omics and phenotypic validations reveal biocontrol mechanisms of Bacillus velezensis XM18-5 against potato common scab. Frontiers in microbiology, 17, 1748645.

Geng X, et al. (2026) Positive effects of bile acids alleviating heat stress in laying hens by enhancing immunity via metabolome and transcriptome integration. Poultry science,

105(3), 106428.

Jin X, et al. (2026) Gut Microbiota-Derived Tyrosol Alleviates Radiation-Induced Intestinal Injury via Targeting SCD1-MUFA Axis to Suppress ER Stress. International journal of biological sciences, 22(5), 2469.

Fossati A, et al. (2026) Bioprospecting Italian biodiversity for identifying potential neuroprotective agents. Scientific reports, 16(1).

Huang Q, et al. (2026) Heat stress suppresses lactation through potential rumen-mammary communication mediated by extracellular vesicles: integrated analysis of microbiome, metabolome, and miRNA profiles. Microbiome, 14(1).

Kim JW, et al. (2026) Methionine metabolism is linked with phospholipid and glutamine metabolism to drive ferroptosis. Cell reports, 45(4), 117157.

Li Y, et al. (2026) Characteristics of gut microbiota and metabolites in patients with metabolic dysfunction-associated steatotic liver disease and colorectal adenoma. Scientific reports, 16(1).

Robinson SD, et al. (2026) Combined multi-omics and multi-spectral profiling of plasma extracellular vesicles reveals liquid biopsy biomarkers for glioma diagnosis. Cell reports. Medicine, 7(5), 102744.

Yao Y, et al. (2026) Preventive effects and mechanisms of yam exosome-like nanoparticles on acute liver injury. Journal of nanobiotechnology, 24(1).

Li G, et al. (2026) Global transcriptional and metabolic reprogramming underlies Fusarium wilt resistance in figleaf gourd (Cucurbita ficifolia). BMC genomics, 27(1).