Resource Summary Report

Generated by NIF on Apr 22, 2025

ProteinPilot

RRID:SCR_024414

Type: Tool

Proper Citation

ProteinPilot (RRID:SCR_024414)

Resource Information

URL: https://sciex.com/products/software/proteinpilot-software

Proper Citation: ProteinPilot (RRID:SCR_024414)

Description: Software tool for protein identification and protein expression analysis. Used to identify proteins and search large numbers of post translational modifications, without increasing search time or false positives. Compatible with all proteomics MS/MS systems.

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

Keywords: protein identification, protein expression analysis, post translational modification search.

Funding:

Availability: Restricted

Resource Name: ProteinPilot

Resource ID: SCR_024414

Alternate URLs: https://sciex.com/content/dam/SCIEX/pdf/posters/ProteinPilot-Software-

Overview.pdf

Record Creation Time: 20230913T050234+0000

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Ratings and Alerts

No rating or validation information has been found for ProteinPilot.

No alerts have been found for ProteinPilot.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 153 mentions in open access literature.

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

Blanco-Pintos T, et al. (2025) Diagnostic Accuracy of Novel Protein Biomarkers in Saliva to Detect Periodontitis Using Untargeted 'SWATH' Mass Spectrometry. Journal of clinical periodontology, 52(2), 199.

Adim H, et al. (2025) iTRAQ-based quantitative proteomic analysis of herbicide stress in Avena ludoviciana Durieu. Scientific reports, 15(1), 577.

Freitas-Ribeiro S, et al. (2025) Cryogenic, but not hypothermic, preservation disrupts the extracellular matrix of cell sheets. Bioactive materials, 46, 301.

Bastola S, et al. (2025) Endothelial-secreted Endocan activates PDGFRA and regulates vascularity and spatial phenotype in glioblastoma. Nature communications, 16(1), 471.

Serrat J, et al. (2025) Binding and cleavage of pro-urokinase by a tegument extract of Fasciola hepatica newly excysted juveniles activate the host fibrinolytic system. Veterinary research, 56(1), 20.

Hussain M, et al. (2025) Interaction of the Wolbachia surface protein with a novel pro-viral protein from Aedes aegypti. mBio, 16(1), e0148624.

Wu F, et al. (2025) A multiprotein regulatory module, MED16-MBR1&2, controls MED25 homeostasis during jasmonate signaling. Nature communications, 16(1), 772.

Wang Y, et al. (2025) Transcriptomic and proteomic analysis reveals the mechanism of chicken cecum response to Salmonella enterica serovar Enteritidis inoculation. iScience, 28(1), 111571.

Qiu R, et al. (2025) AAV2-PDE6B restores retinal structure and function in the retinal degeneration 10 mouse model of retinitis pigmentosa by promoting phototransduction and inhibiting apoptosis. Neural regeneration research, 20(8), 2408.

Schneeweis A, et al. (2025) Mass spectrometry identifies tau C-terminal phosphorylation cluster during neuronal hyperexcitation. Journal of neurochemistry, 169(1), e16221.

Luna-Nophal A, et al. (2025) Preclinical Efficacy and Proteomic Prediction of Molecular Targets for s-cal14.1b and s-cal14.2b Conotoxins with Antitumor Capacity in Xenografts of Malignant Pleural Mesothelioma. Marine drugs, 23(1).

Ding Y, et al. (2024) Oral supplementation of gut microbial metabolite indole-3-acetate alleviates diet-induced steatosis and inflammation in mice. eLife, 12.

Chen X, et al. (2024) Small extracellular vesicles from young plasma reverse age-related functional declines by improving mitochondrial energy metabolism. Nature aging, 4(6), 814.

Domínguez-Lobo MT, et al. (2024) Double blocking of carbon metabolism causes a large increase of Calvin-Benson cycle compounds in cyanobacteria. Plant physiology, 195(2), 1491.

Tuomivaara ST, et al. (2024) Fluoride-related changes in the fetal cord blood proteome; a pilot study. Environmental health: a global access science source, 23(1), 66.

Netherton JK, et al. (2024) The role of HnrnpF/H as a driver of oligoteratozoospermia. iScience, 27(7), 110198.

Kadonosawa Y, et al. (2024) Overexpression of carbonyl reductase 1 in ovarian cancer cells suppresses proliferation and activates the eIF2 signaling pathway. Oncology letters, 28(2), 359.

Chen W, et al. (2024) Exosome-derived tRNA fragments tRF-GluCTC-0005 promotes pancreatic cancer liver metastasis by activating hepatic stellate cells. Cell death & disease, 15(1), 102.

Lv L, et al. (2024) Inhibition of ABI2 ubiquitination-dependent degradation suppresses TNBC cell growth via down-regulating PI3K/Akt signaling pathway. Cancer cell international, 24(1), 222.

Shender VO, et al. (2024) Therapy-induced secretion of spliceosomal components mediates pro-survival crosstalk between ovarian cancer cells. Nature communications, 15(1), 5237.