Resource Summary Report

Generated by NIF on Apr 25, 2025

Metascape

RRID:SCR_016620

Type: Tool

Proper Citation

Metascape (RRID:SCR_016620)

Resource Information

URL: http://metascape.org/gp/index.html#/main/step1

Proper Citation: Metascape (RRID:SCR_016620)

Description: Web service to analyze gene or protein lists. Provides automated meta analysis tools to understand pathways within a group of orthogonal target-discovery studies.

Resource Type: analysis service resource, data analysis service, web service, production service resource, data access protocol, software resource, service resource

Keywords: analyze, gene, protein, list, data, annotate, resource, bio.tools

Funding:

Availability: Free, Freely available

Resource Name: Metascape

Resource ID: SCR_016620

Alternate IDs: biotools:Metascape

Alternate URLs: https://bio.tools/Metascape

Record Creation Time: 20220129T080331+0000

Record Last Update: 20250425T060155+0000

Ratings and Alerts

No rating or validation information has been found for Metascape.

No alerts have been found for Metascape.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2701 mentions in open access literature.

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

Zhang Y, et al. (2025) Exploring markers in nursing care of prostate cancer. Medicine, 104(4), e41357.

Zeng S, et al. (2025) Inhibition of triglyceride metabolism-associated enhancers alters lipid deposition during adipocyte differentiation. FASEB journal: official publication of the Federation of American Societies for Experimental Biology, 39(2), e70347.

Ishikawa H, et al. (2025) Effects of aging on diurnal transcriptome change in the mouse corpus callosum. iScience, 28(1), 111556.

Liu S, et al. (2025) Multi-Omics Analysis for Identifying Cell-Type-Specific Druggable Targets in Alzheimer's Disease. medRxiv: the preprint server for health sciences.

Zhou D, et al. (2025) Dysfunction of ATP7B Splicing Variant Caused by Enhanced Interaction With COMMD1 in Wilson Disease. Cellular and molecular gastroenterology and hepatology, 19(2), 101418.

Dos Santos FM, et al. (2025) Dynamic Cellular Proteome Remodeling during SARS-CoV-2 Infection. Identification of Plasma Protein Readouts. Journal of proteome research, 24(1), 171.

Zhang C, et al. (2025) Proteomic analysis across aged tissues reveals distinct signatures and the crucial involvement of midgut barrier function in the regulation of aging. Aging cell, 24(1), e14344.

Hu Z, et al. (2025) Analysis of the Clinicopathological Characteristics, Genetic Phenotypes, and Prognostics of Primary Pulmonary and Bronchial Adenoid Cystic Carcinoma. Thoracic cancer, 16(2), e15526.

Zhang Y, et al. (2025) Exploring similarities and differences in anti-atherosclerotic potential bioactives among Dendrobium species by UPLC-Q-Exactive Orbitrap MS. NPJ science of food, 9(1), 6.

Du W, et al. (2025) Integrated bioinformatics and experimental analysis of CHAF1B as a novel biomarker and immunotherapy target in LUAD. Discover oncology, 16(1), 43.

Yang X, et al. (2025) CHST3, PGBD5, and SLIT2 can be identified as potential genes for the diagnosis and treatment of osteoporosis and sarcopenia. Scientific reports, 15(1), 374.

Hayakawa T, et al. (2025) Genome-scale evolution in local populations of wild chimpanzees. Scientific reports, 15(1), 548.

Liang H, et al. (2025) LcProt: Proteomics-based identification of plasma biomarkers for lung cancer multievent, a multicentre study. Clinical and translational medicine, 15(1), e70160.

Wang J, et al. (2025) Comparing acute versus AIDS ART initiation on HIV-1 integration sites and clonal expansion. Signal transduction and targeted therapy, 10(1), 23.

Huang S, et al. (2025) Single-base m6A epitranscriptomics reveals novel HIV-1 host interaction targets in primary CD4+ T cells. bioRxiv: the preprint server for biology.

Bai W, et al. (2025) Defining ortholog-specific UHRF1 inhibition by STELLA for cancer therapy. Nature communications, 16(1), 474.

Huang Y, et al. (2025) Metabolic Atlas of Human Eyelid Infiltrative Basal Cell Carcinoma. Investigative ophthalmology & visual science, 66(1), 11.

Yu Q, et al. (2025) Causal genes identification of giant cell arteritis in CD4+?Memory t cells: an integration of multi-omics and expression quantitative trait locus analysis. Inflammation research: official journal of the European Histamine Research Society ... [et al.], 74(1), 3.

Huang F, et al. (2025) Comprehensive bioinformatics analysis of metabolism?related microRNAs in high myopia in young and old adults with age?related cataracts. Molecular medicine reports, 31(2).

Hiraku Y, et al. (2025) Microarray analysis of gene expression in lung tissues of indium-exposed rats: possible roles of S100 proteins in lung diseases. Archives of toxicology, 99(1), 245.