

Resource Summary Report

Generated by [nidm-terms](#) on Aug 1, 2026

MetaP

RRID:SCR_014686

Type: Tool

Proper Citation

MetaP (RRID:SCR_014686)

Resource Information

URL: <http://metap.helmholtz-muenchen.de/metap2/>

Proper Citation: MetaP (RRID:SCR_014686)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on April 5,2023. Software tool for processing in metabolomics experiments., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Synonyms: metap, MetaP Server

Resource Type: computational hosting, service resource

Defining Citation: [DOI:10.1155/2011/839862](https://doi.org/10.1155/2011/839862)

Keywords: metabolomics, metabolomics tool, server, data analysis, processing, computational hosting, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: MetaP

Resource ID: SCR_014686

Alternate IDs: biotools:metap

Alternate URLs: <https://bio.tools/metap>

Record Creation Time: 20220129T080321+0000

Record Last Update: 20260731T042923+0000

Ratings and Alerts

No rating or validation information has been found for MetaP.

No alerts have been found for MetaP.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 14 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Okoye GD, et al. (2025) Single-cell map of innate-like lymphocyte response to Francisella tularensis infection reveals interleukin-17-dependent protection by MAIT cells. iScience, 28(3), 111810.

Scrivano L, et al. (2024) Active mobility and mental health: A scoping review towards a healthier world. Global mental health (Cambridge, England), 11, e1.

Davidson RK, et al. (2024) Loss of STAT3 in osteoblasts has detrimental and sexually dimorphic effects on skeletal development. PloS one, 19(12), e0315078.

Li J, et al. (2022) Determination of the nanoparticle- and cell-specific toxicological mechanisms in 3D liver spheroids using scRNAseq analysis. Nano today, 47.

Trzuppek D, et al. (2020) Discovery of CD80 and CD86 as recent activation markers on regulatory T cells by protein-RNA single-cell analysis. Genome medicine, 12(1), 55.

Capasso M, et al. (2017) The functional variant rs34330 of CDKN1B is associated with risk of neuroblastoma. Journal of cellular and molecular medicine, 21(12), 3224.

Madore AM, et al. (2016) Genes Involved in Interleukin-1 Receptor Type II Activities Are Associated With Asthmatic Phenotypes. Allergy, asthma & immunology research, 8(5), 466.

Hill CB, et al. (2015) Metabolomics, Standards, and Metabolic Modeling for Synthetic Biology in Plants. Frontiers in bioengineering and biotechnology, 3, 167.

Kim DS, et al. (2014) The relationship between diastolic blood pressure and coronary artery calcification is dependent on single nucleotide polymorphisms on chromosome 9p21.3. BMC medical genetics, 15, 89.

Liscovitch N, et al. (2014) Differential Co-Expression between α -Synuclein and IFN- γ Signaling Genes across Development and in Parkinson's Disease. PloS one, 9(12), e115029.

Firnberg E, et al. (2013) The genetic code constrains yet facilitates Darwinian evolution. Nucleic acids research, 41(15), 7420.

Chen Y, et al. (2012) Interaction between smoking and functional polymorphism in the TGFB1 gene is associated with ischaemic heart disease and myocardial infarction in patients with rheumatoid arthritis: a cross-sectional study. *Arthritis research & therapy*, 14(2), R81.

Mehrotra S, et al. (2011) Genetic and functional evaluation of the role of CXCR1 and CXCR2 in susceptibility to visceral leishmaniasis in north-east India. *BMC medical genetics*, 12, 162.

Sun L, et al. (2011) Bivariate genome-wide association analyses of femoral neck bone geometry and appendicular lean mass. *PloS one*, 6(11), e27325.