

Resource Summary Report

Generated by [NIF](#) on Aug 28, 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: 20260821T194006+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 15 mentions in open access literature.

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

Zhang Y, et al. (2026) Co-occurrence of transcriptionally distinct persister cell states underpins neoadjuvant therapy resistance in triple-negative breast cancer. *Genome medicine*, 18(1).

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.

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

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.