

Resource Summary Report

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MetaBase

RRID:SCR_001762

Type: Tool

Proper Citation

MetaBase (RRID:SCR_001762)

Resource Information

URL: <https://www.metabase.com/>

Proper Citation: MetaBase (RRID:SCR_001762)

Description: User-contributed list of biological databases available on the internet. Currently there are 1,801 entries, each describing a different database. The databases are described in a semi-structured way by using templates and entries can carry various user comments and annotations. Entries can be searched, listed or browsed by category. The site uses the same MediaWiki technology that powers Wikipedia, The Mediawiki system allows users to participate on many different levels, ranging from authors and editors to curators and designers. MetaBase aims to be a flexible, user-driven (user-created) resource for the biological database community. The main focuses of MetaBase are: * As a basic requirement, MB contains a list of databases, URLs and descriptions of the most commonly used biological databases currently available on the internet. * The system should be flexible, allowing users to contribute, update and maintain the data in different ways. * In the future we aim to generate more communication between the database developer and user communities.

Abbreviations: MB

Synonyms: MetaBase (MB)

Resource Type: narrative resource, database, data or information resource, wiki

Defining Citation: [PMID:22139927](#)

Keywords: biological, mediawiki, biology, bio.tools, FASEB list

Funding: MKE - Ministry of Knowledge Economy

Availability: Free, Freely available

Resource Name: MetaBase

Resource ID: SCR_001762

Alternate IDs: biotools:metabase, nif-0000-10293

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

Old URLs: http://biodatabase.org/index.php?title=Main_Page&oldid=8972

Record Creation Time: 20220129T080209+0000

Record Last Update: 20260805T044028+0000

Ratings and Alerts

No rating or validation information has been found for MetaBase.

No alerts have been found for MetaBase.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 81 mentions in open access literature.

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

Yang Z, et al. (2025) Large Language Model-Based Critical Care Big Data Deployment and Extraction: Descriptive Analysis. JMIR medical informatics, 13, e63216.

Puig-G??mez M, et al. (2025) Transcriptional signature of rapidly responding NK cells reveals S1P5 and CXCR4 as anti-tumor response disruptors. Scientific reports, 15(1), 10769.

Fomin V, et al. (2025) Machine learning identifies clinical tumor mutation landscape pathways of resistance to checkpoint inhibitor therapy in NSCLC. Journal for immunotherapy of cancer, 13(3).

Bachir ZR, et al. (2025) "Speed": A Dataset for Human Speed Estimation. Sensors (Basel, Switzerland), 25(20).

Klein D, et al. (2025) Building a Digital Health Research Platform to Enable Recruitment, Enrollment, Data Collection, and Follow-Up for a Highly Diverse Longitudinal US Cohort of 1 Million People in the All of Us Research Program: Design and Implementation Study. Journal of medical Internet research, 27, e60189.

Almeida JRG, et al. (2025) Comparison of joint position sense measured by inertial sensors embedded in portable digital devices with different masses. Frontiers in neuroscience, 19, 1561241.

Tran TA, et al. (2025) WEBA dataset as the Reflection of Work content effect on Workload perception in Real life Working conditions. *Scientific data*, 12(1), 11.

Piekniewska A, et al. (2025) Do organisms need an impact factor? Citations of key biological resources including model organisms reveal usage patterns and impact. *PloS one*, 20(8), e0327344.

Dal Col SVC, et al. (2025) Improving critical care through telemedicine: a comprehensive analysis of a Tele-ICU project in northern and northeastern regions of Brazil. *BMC health services research*, 25(1), 809.

Gao B, et al. (2025) xOmicsShiny: an R Shiny application for cross-omics data analysis and pathway mapping. *Bioinformatics advances*, 5(1), vbaf097.

Wesson T, et al. (2025) Feasibility of Using Inertial Measurement Units (IMUs) to Augment Cadaveric Temporal Training. *The Laryngoscope*, 135(4), 1465.

Janowski R, et al. (2025) Optimizing Engagement With a Smartphone App to Prevent Violence Against Adolescents: Results From a Cluster Randomized Factorial Trial in Tanzania. *Journal of medical Internet research*, 27, e60102.

Levinger P, et al. (2025) IMProving older people's health through physical ACTivity: the implementation of a community-based framework to promote park-based active ageing. *BMC public health*, 25(1), 4385.

Kaur P, et al. (2025) Scaling India Hypertension Control Initiative strategies to 15 states-treatment outcomes and risk factors for uncontrolled blood pressure, India: a cohort study, 2018-2021. *BMJ open*, 15(11), e106372.

Talay LA, et al. (2024) Why People Seek Obesity Care Through Digital Rather Than In-Person Services: A Quantitative Multinational Analysis of Patients From a Large Unsubsidized Digital Obesity Provider. *Cureus*, 16(12), e75603.

Talay L, et al. (2024) Patient Adherence to a Real-World Digital, Asynchronous Weight Loss Program in Australia That Combines Behavioural and GLP-1 RA Therapy: A Mixed Methods Study. *Behavioral sciences (Basel, Switzerland)*, 14(6).

Nuhamunada M, et al. (2024) BGCFLOW: systematic pangenome workflow for the analysis of biosynthetic gene clusters across large genomic datasets. *Nucleic acids research*, 52(10), 5478.

Dam D, et al. (2024) COVID-19 outcome trends by vaccination status in Canada, December 2020-January 2022. *Canada communicable disease report = Relevé des maladies transmissibles au Canada*, 50(1-2), 40.

Talay L, et al. (2024) The Role of Pharmacists in Minimizing the Risk Inherent in Unbundled Telehealth Services: A 12-Month Retrospective Case Study. *Pharmacy (Basel, Switzerland)*, 12(6).

Tay T, et al. (2024) Degradation of IKZF1 prevents epigenetic progression of T??cell exhaustion in an antigen-specific assay. *Cell reports. Medicine*, 5(11), 101804.