Resource Summary Report

Generated by RRID on Aug 4, 2026

Database Commons

RRID:SCR_023661
Type: Tool

Proper Citation

Database Commons (RRID:SCR_023661)

Resource Information

URL: https://ngdc.cncb.ac.cn/databasecommons/

Proper Citation: Database Commons (RRID:SCR_023661)

Description: Curated catalog of worldwide biological databases to provide landscape of biological databases throughout the world and enable easy retrieval and access to specific collection of databases of interest. Catalog of worldwide biological databases as well as their curated meta information and derived statistics.

Resource Type: catalog, data or information resource, database

Defining Citation: PMID:36572336, PMID:34718731

Keywords: Curated catalog, worldwide biological databases, curated meta information,

Funding: Strategic Priority Research Program of the Chinese Academy of Sciences ;
National Natural Science Foundation of China ;
Professional Association of the Alliance of International Science Organizations ;
Youth Innovation Promotion Association of Chinese Academy of Sciences ;
International Partnership Program of the Chinese Academy of Sciences

Availability: Free, Freely available

Resource Name: Database Commons

Resource ID: SCR_023661

Record Creation Time: 20230607T050210+0000

Record Last Update: 20260804T043806+0000

Ratings and Alerts

No rating or validation information has been found for Database Commons.

No alerts have been found for Database Commons.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 13 mentions in open access literature.

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

Yang LP, et al. (2026) Targeted methylation of cg24067911 suppresses colorectal cancer metastasis through BCL6-ATXN1-CDH1 axis. Oncogene, 45(2), 295.

Qu X, et al. (2025) Recent Advances of Processing and Detection Techniques on Crustacean Allergens: A Review. Foods (Basel, Switzerland), 14(2).

, et al. (2025) Database Resources of the National Genomics Data Center, China National Center for Bioinformation in 2025. Nucleic acids research, 53(D1), D30.

Chen W, et al. (2025) Qing-Re-Hua-Shi Decoction ameliorates DSS-induced colitis by modulating multiple signaling pathways and remodeling the gut microbiota and metabolite profile. Frontiers in cellular and infection microbiology, 15, 1541289.

Mu L, et al. (2025) Machine learning-based transcriptomic analysis identifies candidate genes in sepsis-induced coagulopathy and explores the immunomodulatory potential of baicalein. Human genomics, 19(1), 102.

Yang J, et al. (2024) Identification of genes related to fatty acid metabolism in type 2 diabetes mellitus. Biochemistry and biophysics reports, 40, 101849.

, et al. (2024) Database Resources of the National Genomics Data Center, China National Center for Bioinformation in 2024. Nucleic acids research, 52(D1), D18.

Reed CJ, et al. (2024) Beyond blast: enabling microbiologists to better extract literature, taxonomic distributions and gene neighbourhood information for protein families. Microbial genomics, 10(2).

Teng X, et al. (2024) Three bioactive compounds from Huangqin decoction ameliorate Irinotecan-induced diarrhea via dual-targeting of Escherichia coli and bacterial ??-glucuronidase. Cell biology and toxicology, 40(1), 88.

Li H, et al. (2024) Identifying key inflammatory genes in psoriasis via weighted gene co-expression network analysis: Potential targets for therapy. Biomolecules & biomedicine, 24(5), 1133.

Liu T, et al. (2024) Genome-Wide Identification and Expression Analysis of Sucrose Nonfermenting 1-Related Protein Kinase (SnRK) Genes in Salvia miltiorrhiza in Response to Hormone. Plants (Basel, Switzerland), 13(7).

Wei F, et al. (2024) Lymph node metastasis determined miRNAs in esophageal squamous cell carcinoma. Aging, 16(20), 13104.

Yang LP, et al. (2023) Comprehensive genomic characterization of sporadic synchronous colorectal cancer: Implications for treatment optimization and clinical outcome. Cell reports. Medicine, 4(10), 101222.