

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 1, 2026

SQLite

RRID:SCR_017672

Type: Tool

Proper Citation

SQLite (RRID:SCR_017672)

Resource Information

URL: <https://www.sqlite.org/index.html>

Proper Citation: SQLite (RRID:SCR_017672)

Description: Relational database management system contained in C library. SQLite is not client server database engine but it is embedded into the end program.

Resource Type: software resource, software application

Keywords: Database, management, system, C library, engine, embedded, the end program

Availability: Free, Freely available

Resource Name: SQLite

Resource ID: SCR_017672

License: Public domain

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260731T043012+0000

Ratings and Alerts

No rating or validation information has been found for SQLite.

No alerts have been found for SQLite.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 65 mentions in open access literature.

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

Chorbadjiev L, et al. (2025) Analyzing the large and complex SFARI autism cohort data using the Genotypes and Phenotypes in Families (GPF) platform. *Genome research*, 35(10), 2352.

Miyata K, et al. (2025) Pharmacovigilance study for the identification of mogamulizumab-induced immune-related adverse events using a real-world database. *The oncologist*, 30(7).

Zuo Z, et al. (2025) AlgicideDB: a comprehensive database enhanced by large language models for algicide management and discovery. *Frontiers in microbiology*, 16, 1611403.

Mendel HC, et al. (2025) PeptideMiner-neuropeptide discovery across the animal kingdom. *GigaScience*, 14.

Yang F, et al. (2025) Deep learning reconstruction for accelerating synthetic magnetic resonance imaging of the breast: a comparative analysis of accelerated and standard protocols. *Quantitative imaging in medicine and surgery*, 15(10), 8953.

Rajesh MK, et al. (2024) Chromosome scale genome assembly and annotation of coconut cultivar Chowghat Green Dwarf. *Scientific reports*, 14(1), 28778.

Furuta T, et al. (2024) MCPtaggR: R package for accurate genotype calling in reduced representation sequencing data by eliminating error-prone markers based on genome comparison. *DNA research : an international journal for rapid publication of reports on genes and genomes*, 31(1).

Mok BA, et al. (2024) Web-based psychoacoustics: Hearing screening, infrastructure, and validation. *Behavior research methods*, 56(3), 1433.

Söderholm JJ, et al. (2024) How changes in depression severity and borderline personality disorder intensity are linked - a cohort study of depressed patients with and without borderline personality disorder. *Borderline personality disorder and emotion dysregulation*, 11(1), 3.

Navandar M, et al. (2024) Long non-coding RNAs expression and regulation across different brain regions in primates. *Scientific data*, 11(1), 545.

Maia Cordeiro C, et al. (2024) BioPipeline Creator-a user-friendly Java-based GUI for managing and customizing biological data pipelines. *Scientific reports*, 14(1), 16572.

Chen C, et al. (2024) PancanQTLv2.0: a comprehensive resource for expression quantitative trait loci across human cancers. *Nucleic acids research*, 52(D1), D1400.

Stawiski M, et al. (2024) Optimizing neuroscience data management by combining REDCap, BIDS and SQLite: a case study in Deep Brain Stimulation. *Frontiers in neuroinformatics*, 18, 1435971.

Strayer EC, et al. (2024) NaP-TRAP reveals the regulatory grammar in 5'UTR-mediated translation regulation during zebrafish development. *Nature communications*, 15(1), 10898.

Yamauchi K, et al. (2024) Combined substituent number utilized machine learning for the development of antimicrobial agent. *Scientific reports*, 14(1), 4106.

Chen Z, et al. (2024) PungentDB: Bridging traditional Chinese medicine of medicine food homology and modern food flavor chemistry. *Food chemistry: X*, 23, 101742.

Sun L, et al. (2024) GEPREP: A comprehensive data atlas of RNA-seq-based gene expression profiles of exercise responses. *Journal of sport and health science*, 14, 100992.

Zheng M, et al. (2023) ChemPert: mapping between chemical perturbation and transcriptional response for non-cancer cells. *Nucleic acids research*, 51(D1), D877.

Michell-Robinson MA, et al. (2023) Hypomyelination, hypodontia and craniofacial abnormalities in a Polr3b mouse model of leukodystrophy. *Brain : a journal of neurology*, 146(12), 5070.

Lipman D, et al. (2023) Integrative multi-omics approach for identifying molecular signatures and pathways and deriving and validating molecular scores for COVID-19 severity and status. *BMC genomics*, 24(1), 319.