

Resource Summary Report

Generated by [NIF](#) on Aug 12, 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 application, software resource

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: 20260810T040725+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 83 mentions in open access literature.

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

Hwang J, et al. (2026) Development of K-CORE: a web-based platform for integrated clinico-genomic analysis. Life science alliance, 9(8).

Lorenzo MCC, et al. (2026) Single-Cell Morphometrics From Hematoxylin and Eosin-Stained Images Reveals Subtype-Specific Features in Chronic Lymphocytic Leukemia. Cytometry. Part A : the journal of the International Society for Analytical Cytology, 109(1), 62.

Rudl Kulhava L, et al. (2025) Circadian ontogenetic metabolomics atlas: an interactive resource with insights from rat plasma, tissues, and feces. Cellular and molecular life sciences : CMLS, 82(1), 264.

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).

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.

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.

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

Zhai J, et al. (2025) GenMasterTable: a user-friendly desktop application for filtering, summarising, and visualising large-scale annotated genetic variants. BMC bioinformatics, 26(1), 211.

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

Berikashvili LB, et al. (2025) Postoperative outcomes in patients with COPD undergoing elective non-cardiac surgery: a propensity score-matched cohort study. Frontiers in medicine, 12, 1641613.

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

S??derholm 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.

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.

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

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

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).

Du Q, et al. (2024) CBGDA: a manually curated resource for gene-disease associations based on genome-wide CRISPR. *Database : the journal of biological databases and curation*, 2024.

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.

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.

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