

Resource Summary Report

Generated by [NIF](#) on Aug 7, 2026

AceDB

RRID:SCR_010671

Type: Tool

Proper Citation

AceDB (RRID:SCR_010671)

Resource Information

URL: <http://www.acedb.org/>

Proper Citation: AceDB (RRID:SCR_010671)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on April 6th,2023. Software genome database management system.

Synonyms: A C. elegans DataBase

Resource Type: service resource, database, data or information resource

Defining Citation: [DOI:10.1109/5992.764215](https://doi.org/10.1109/5992.764215)

Keywords: genome database management system, genome database, management system,

Availability: THIS RESOURCE IS NO LONGER IN SERVICE.

Resource Name: AceDB

Resource ID: SCR_010671

Alternate IDs: nlx_75315, OMICS_15828

Alternate URLs: <https://sources.debian.org/src/acedb-other/>

Record Creation Time: 20220129T080300+0000

Record Last Update: 20260807T042742+0000

Ratings and Alerts

No rating or validation information has been found for AceDB.

No alerts have been found for AceDB.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

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

Lockhead D, et al. (2016) The tubulin repertoire of *C. elegans* sensory neurons and its context-dependent role in process outgrowth. *Molecular biology of the cell*, 27(23), 3717.

Van Auken K, et al. (2012) Text mining in the biocuration workflow: applications for literature curation at WormBase, dictyBase and TAIR. *Database : the journal of biological databases and curation*, 2012, bas040.

Leschziner GD, et al. (2011) Q8IYL2 is a candidate gene for the familial epilepsy syndrome of Partial Epilepsy with Pericentral Spikes (PEPS). *Epilepsy research*, 96(1-2), 109.

Bentley SD, et al. (2007) Meningococcal genetic variation mechanisms viewed through comparative analysis of serogroup C strain FAM18. *PLoS genetics*, 3(2), e23.

N??gre V, et al. (2006) SPODOBASE: an EST database for the lepidopteran crop pest Spodoptera. *BMC bioinformatics*, 7, 322.

Jordan PW, et al. (2005) Strain-specific differences in *Neisseria gonorrhoeae* associated with the phase variable gene repertoire. *BMC microbiology*, 5, 21.

Lewis SE, et al. (2005) Gene Ontology: looking backwards and forwards. *Genome biology*, 6(1), 103.

Galperin MY, et al. (2005) The Molecular Biology Database Collection: 2005 update. *Nucleic acids research*, 33(Database issue), D5.

Sheps JA, et al. (2004) The ABC transporter gene family of *Caenorhabditis elegans* has implications for the evolutionary dynamics of multidrug resistance in eukaryotes. *Genome biology*, 5(3), R15.

Burren OS, et al. (2004) Development of an integrated genome informatics, data management and workflow infrastructure: a toolbox for the study of complex disease genetics. *Human genomics*, 1(2), 98.

Lewis SE, et al. (2002) Apollo: a sequence annotation editor. *Genome biology*, 3(12), RESEARCH0082.

Dowell RD, et al. (2001) The distributed annotation system. *BMC bioinformatics*, 2, 7.