## **Resource Summary Report**

Generated by dkNET on May 16, 2025

# **WorfDB**

RRID:SCR\_006028

Type: Tool

### **Proper Citation**

WorfDB (RRID:SCR\_006028)

### **Resource Information**

URL: http://worfdb.dfci.harvard.edu/

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**Description:** Database that integrates and disseminates the data from the cloning of complete set of predicted protein-encoding ORFs of Caenorhabditis elegans. It also allows the community to search for availability and quality of cloned ORFs. So far, ORF sequence tags (OSTs) obtained for all individual clones have allowed exon structure corrections for ORFs originally predicted by the C. elegans sequencing consortium. The database contains this OST information along with data pertinent to the cloning process.

Synonyms: WorfDB - Worm ORF Database, Worm ORFeome DataBase, Worm ORFeome

Resource Type: data or information resource, database

**Defining Citation:** PMID:12519990

**Keywords:** open reading frame, c elegans, orf sequence tag

Funding: NIGMS;

MGRI:

NHGRI 5R01HG01715-02; NCI 7 R33 CA81658-02

Resource Name: WorfDB

Resource ID: SCR\_006028

**Alternate IDs:** nif-0000-03644

**Record Creation Time: 20220129T080233+0000** 

**Record Last Update**: 20250507T060346+0000

## Ratings and Alerts

No rating or validation information has been found for WorfDB.

No alerts have been found for WorfDB.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 9 mentions in open access literature.

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

Leslie EJ, et al. (2016) A multi-ethnic genome-wide association study identifies novel loci for non-syndromic cleft lip with or without cleft palate on 2p24.2, 17q23 and 19q13. Human molecular genetics, 25(13), 2862.

Van Assche R, et al. (2015) Integrating -Omics: Systems Biology as Explored Through C. elegans Research. Journal of molecular biology, 427(21), 3441.

Ou CY, et al. (2010) Two cyclin-dependent kinase pathways are essential for polarized trafficking of presynaptic components. Cell, 141(5), 846.

Merritt C, et al. (2010) Transgenic solutions for the germline. WormBook : the online review of C. elegans biology, 1.

Warner GJ, et al. (2006) Interactome networks: the state of the science. Genome biology, 7(1), 301.

Farkas IJ, et al. (2006) Exploring transcriptional regulatory networks in the worm. Cell, 125(6), 1032.

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

Hare EE, et al. (2004) Function and evolution of the serotonin-synthetic bas-1 gene and other aromatic amino acid decarboxylase genes in Caenorhabditis. BMC evolutionary biology, 4, 24.

Davy A, et al. (2001) A protein-protein interaction map of the Caenorhabditis elegans 26S

proteasome. EMBO reports, 2(9), 821.