

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 16, 2026

[Orthology.eg.db](#)

RRID:SCR_024740

Type: Tool

Proper Citation

Orthology.eg.db (RRID:SCR_024740)

Resource Information

URL: <https://bioconductor.org/packages/Orthology.eg.db/>

Proper Citation: Orthology.eg.db (RRID:SCR_024740)

Description: Software R package to provide orthology mappings between species, based on NCBI Gene IDs and NCBI orthology mappings.

Resource Type: software resource, software toolkit

Keywords: orthology mappings between species, NCBI Gene IDs, NCBI orthology mappings,

Availability: Free, Available for download, Freely available

Resource Name: Orthology.eg.db

Resource ID: SCR_024740

License: Artistic v2.0

Record Creation Time: 20231204T190628+0000

Record Last Update: 20260912T200430+0000

Ratings and Alerts

No rating or validation information has been found for Orthology.eg.db.

No alerts have been found for Orthology.eg.db.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Jessa S, et al. (2024) FOXR2 targets LHX6+/DLX+ neural lineages to drive CNS neuroblastoma. Cancer research.