

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 2, 2026

Evola

RRID:SCR_007651

Type: Tool

Proper Citation

Evola (RRID:SCR_007651)

Resource Information

URL: <http://jbirc.jbic.or.jp/hinv/evola/>

Proper Citation: Evola (RRID:SCR_007651)

Description: Evola is a sub-database of H-InvDB, providing ortholog data as evolutionary annotation. Representative transcripts (one transcript per one gene locus) were analyzed as genes. Orthologs were first detected by computational analysis. Then, more reliable orthologs were determined by manual curation inspecting the phylogenetic trees., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Synonyms: Evola

Resource Type: data or information resource, database

Keywords: ortholog

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Evola

Resource ID: SCR_007651

Alternate IDs: nif-0000-02824

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260801T190946+0000

Ratings and Alerts

No rating or validation information has been found for Evola.

No alerts have been found for Evola.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Kim Y, et al. (2019) A Mutation in ZNF143 as a Novel Candidate Gene for Endothelial Corneal Dystrophy. Journal of clinical medicine, 8(8).

Yang M, et al. (2017) Clinical, biochemical and molecular characterization of Korean patients with mucopolipidosis II/III and successful prenatal diagnosis. Orphanet journal of rare diseases, 12(1), 11.