

# Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

## EGO - Eukaryotic Gene Orthologs

RRID:SCR\_000732

Type: Tool

### Proper Citation

EGO - Eukaryotic Gene Orthologs (RRID:SCR\_000732)

### Resource Information

**URL:** <http://compbio.dfci.harvard.edu/tgi/ego/>

**Proper Citation:** EGO - Eukaryotic Gene Orthologs (RRID:SCR\_000732)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23,2022. EGO is a database of eukaryotic gene orthologs generated by pairwise comparison between the Tentative Consensus (TC) sequences that comprise the Dana Farber Gene Indices from individual organisms.

**Synonyms:** EGO

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

**Keywords:** eukaryotic gene ortholog, dana farber gene indices, tentative consensus sequence

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** EGO - Eukaryotic Gene Orthologs

**Resource ID:** SCR\_000732

**Alternate IDs:** nif-0000-02790

**Record Creation Time:** 20220129T080203+0000

**Record Last Update:** 20260912T200120+0000

### Ratings and Alerts

No rating or validation information has been found for EGO - Eukaryotic Gene Orthologs.

No alerts have been found for EGO - Eukaryotic Gene Orthologs.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Hu P, et al. (2012) Microarray meta-analysis identifies acute lung injury biomarkers in donor lungs that predict development of primary graft failure in recipients. PloS one, 7(10), e45506.

Page GP, et al. (2008) Bioinformatic tools for inferring functional information from plant microarray data: tools for the first steps. International journal of plant genomics, 2008, 147563.