

# Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

## Interactive Fly

RRID:SCR\_012784

Type: Tool

### Proper Citation

Interactive Fly (RRID:SCR\_012784)

### Resource Information

**URL:** <http://www.sdbonline.org/fly/aimain/1aahome.htm>

**Proper Citation:** Interactive Fly (RRID:SCR\_012784)

**Description:** The InterActive Fly is an online database of Drosophila development and metazoan evolution. It contains information on biochemical pathways, organs, images, a cis-decoder, and EvoPrinter, a machine that allows users to identify Evolutionarily Resilient DNA Sequences.

**Abbreviations:** Interactive Fly

**Synonyms:** Interactive Fly: A cyberspace guide to Drosophila development and metazoan evolution

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

**Resource Name:** Interactive Fly

**Resource ID:** SCR\_012784

**Alternate IDs:** nif-0000-03030

**Record Creation Time:** 20220129T080312+0000

**Record Last Update:** 20260905T133203+0000

### Ratings and Alerts

No rating or validation information has been found for Interactive Fly.

No alerts have been found for Interactive Fly.

### Data and Source Information

## Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Thurmond J, et al. (2019) FlyBase 2.0: the next generation. Nucleic acids research, 47(D1), D759.

Marques E, et al. (2016) Par6G suppresses cell proliferation and is targeted by loss-of-function mutations in multiple cancers. Oncogene, 35(11), 1386.

Behura SK, et al. (2011) Comparative genomic analysis of *Drosophila melanogaster* and vector mosquito developmental genes. PloS one, 6(7), e21504.

Zeng V, et al. (2011) De novo assembly and characterization of a maternal and developmental transcriptome for the emerging model crustacean *Parhyale hawaiiensis*. BMC genomics, 12, 581.

Zhan S, et al. (2011) The monarch butterfly genome yields insights into long-distance migration. Cell, 147(5), 1171.

Yan H, et al. (2010) A genome-wide gene function prediction resource for *Drosophila melanogaster*. PloS one, 5(8), e12139.

Hooper SD, et al. (2007) Identification of tightly regulated groups of genes during *Drosophila melanogaster* embryogenesis. Molecular systems biology, 3, 72.