Resource Summary Report

Generated by dkNET on Aug 27, 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

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Description: The InterActive Fly is an online database of Drosophilia 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: 20260821T193938+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

Source: SciCrunch Registry

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

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.

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

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

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.