

Resource Summary Report

Generated by [NIF](#) on Aug 26, 2026

Homophila

RRID:SCR_007717

Type: Tool

Proper Citation

Homophila (RRID:SCR_007717)

Resource Information

URL: <http://superfly.ucsd.edu/>

Proper Citation: Homophila (RRID:SCR_007717)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented on June 23, 2013. Homophila utilizes the sequence information of human disease genes from the NCBI OMIM (Online Mendelian Inheritance in Man) database in order to determine if sequence homologs of these genes exist in the current Drosophila sequence database (FlyBase). Sequences are compared using NCBI's BLAST program. The database is updated weekly and can be searched by human disease, gene name, OMIM number, title, subtitle and/or allelic variant descriptions.

Synonyms: Human disease to drosophila database

Resource Type: data or information resource, database

Defining Citation: [PMID:11752278](#), [PMID:11381037](#)

Keywords: homolog, human disease, human disease gene, human, gene, cognate

Funding: NCRR P 41 RR08605-06

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Homophila

Resource ID: SCR_007717

Alternate IDs: nif-0000-02976

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260821T193823+0000

Ratings and Alerts

No rating or validation information has been found for Homophila.

No alerts have been found for Homophila.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.