



Search

Search FlyMine. Enter **names, identifiers or keywords** for genes, proteins, pathways, ontology terms, authors, etc. (e.g. *eve*, HIPPO_DROME, glycolysis, *hb* allele).

e.g. zen, adh, CG2328, F8gn0000099

SEARCH

GENES

INTERACTIONS

The gene models and other genomic annotations in FlyMine are derived from Ensembl, and over 30 other data sources.

Query for genes:

- Gene → Proteins.
- Chromosome → All genes.
- Gene → Chromosomal location.
- Gene → Protein → Protein domains.
- Gene [D. melanogaster] → Regulatory elements.
- Protein → Gene.
- Organism → All intergenic regions.
- Gene → Transcripts → Exons → Introns

» [More queries](#)

Perl, Python, Ruby and Java API

Access our FlyMine data via our Application Programming Interface (API) too! We provide client libraries in the following languages:

[Perl](#) [Python](#) [Ruby](#) [Java](#)

Analysis & more queries

Gene

e.g. zen, adh, CG2328, F8gn0000099

Perl, Python, Ruby and Java API



Access our FlyMine data via our Application Programming Interface (API) too! We provide client libraries in the following languages:

[Perl](#)

[Python](#)

[Ruby](#)

[Java](#)

News & Updates

- 7 Aug [FlyMine 38.0](#)
We're pleased to announce a new version of FlyMine.
- 9 May [FlyMine 37.0](#)
We've released an updated version of FlyMine. All of the data sets have been updated.
- 11 Dec [FlyMine 36.0](#)
We've released version 36.0 of FlyMine. We've fixed a few bugs and updated some data.

» [More news](#)