TACACAATCAGTTAGTTTCCACCGACAGTCCGCAGAAACCATTCGACGGC GTCGGCAATCCGTAAGATAGCCAAATATTATTATTGTTCAGATACTCACT AGC ACACAGTGCAGATC AAATCAGTGAAATTC AATAATAAAAACAACAACAGTGCAACAACAGCCGGGGCATCTTCATAGAT

Human-health-related models in Drosophila melanogaster

Introduction

The number and proportion of Drosophila articles published each year that mention disease has risen steadily. Many of these papers describe Drosophila models of human diseases or investigate functional conservation between human-fly orthologs. To help researchers find genes and alleles that model human diseases in Drosophila FlyBase has started to collect and display this information from primary research articles.



Papers containing "disease" in the abstract or title

Human-health-related models in FlyBase

What disease models are captured by FlyBase?

In our initial efforts to assemble data on disease models on Drosophila we have limited the data curation to genetic models which result in a phenotype that recapitulates some aspect of human disease. The genetic models we have applied this to are at the allele level i.e. at-locus-mutations and transgenes.

What alleles are included?

•Disease-phenotype-causing- alleles with links to human disease •Alleles which modify the disease-model phenotype.

•How are alleles linked with disease?

·Alleles are associated with standardized human disease terms from the Disease Ontology (disease-ontology.org).

•An allele's relationship with a disease can be described as

• a model of

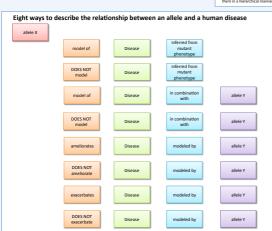
•a modifier: ameliorates or exacerbates a disease modeled by another allele

•When >1 allele models the disease, "in combination with" is used to describe their relationship.

•a DOES NOT prefix is used for cases when a particular allele fails to manifest the expected phenotype.

•Therefore using simple expressions there are multiple ways in which the relationship between an allele and disease can be described.





Finding disease-related papers

- •To identify potential papers for disease curation we have:
 •Searched the archives for papers with "disease" in the title or abstract,
- •Examined papers known to use human transgenes,
 •Sought author help: Flagging "Human Disease" on our "Fast Track Your Paper" tool.
 •From this we have been able to associate ~1000 alleles of over 500 genes, spanning ~100 diseases (DO terms).



Drosophila has proven an incredibly tractable model for neurodegenerative disease. The fly has a complex brain and its short life-span enables the study of these diseases which often have an agedependent pathology. It is therefore not surprising that the majority of Drosophila human-disease models relate to neurodegenerative diseases. This is illustrated in the word cloud above, generated from the Disease Ontology terms we have associated with alleles in FlyBase.

How can I find human disease models in FlyBase?

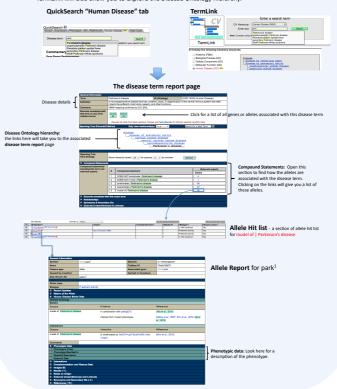
1. Is a particular gene associated with a disease model?

In the Gene Report page for each gene there is now a Human Disease Model Data section.



2. What genes/alleles are associated with a particular disease model?

There are two ways to directly search for a disease: QuickSearch and the dedicated Controlled Vocabulary search tool **TermLink.** Both have autocomplete functions, but TermLink will also allow you to explore the Disease Ontology hierarchy.



This is just the beginning

- •This newly compiled human disease data is in the FB2014 02 release of FlyBase.
- •Human disease curation is now an integral part of new paper curation at FlyBase.
- •We are still looking at past papers to identify potential disease models and text-mining will be employed to help trawl the archives.
- •A precomputed file will be available in the next release.
- •Please help us make this resource as comprehensive and useful as possible by providing feedback, suggestions and alerting us to models we have missed. Use the Contact FlyBase link (found at the bottom of each page in FlyBase).

