

# FlyBase 2.0 - What's new?

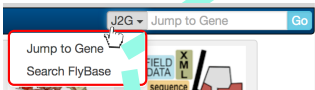
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## FlyBase on any device!



FlyBase 2.0 provides good usability on all devices, including smart phones and tablets

## J2G/Simple Search Box



There is a Jump-to-gene (J2G)/Search FlyBase box at the top right of every page. J2G goes straight to specific gene or allele report pages; Search FlyBase retrieves hits from all data classes matching the query entry

## HitLists - fresh from the oven!

### List view

All hits of all data classes

**HitList filters**

- Species filter \*coming soon\*
- Data class filter

### Table view

For single data classes

**Table sorting feature**

Symbol	Name	Annotation ID	Orthology	Scaffold	# Alleles	# Stocks	# Refs	Genome View
Egfr	Epidermal growth factor receptor	CG10079	57E9-37F1	2R	169	53	1597	
Egfr<sup>C22</sup>	Enhancer of Egfr C22	6303-93F14			1	1	1	
Egfr<sup>C12</sup>	Enhancer of Egfr C12				1	0	1	
Egfr<sup>B56</sup>	Enhancer of Egfr B56	49D1-49D4			1	1	4	
Egfr<sup>Hsp1</sup>/EGFR					1	1	1	
Egfr<sup>tor</sup>	Egfr<sup>tor</sup>				3	0	28	
tblt:Egfr	tblt:Egfr				17	1	111	
Egfr<sup>36</sup>	Egfr<sup>36</sup>				1	0	1	

## Report pages - loads of new features!

### Gene Snapshots

Some protein-coding genes have short summaries solicited from expert researchers, curated and standardized by FlyBase, which provide a quick overview of the function of a gene product.

### "Alliance" Linkouts

Gene reports linkout to the Alliance of Genome resources, a consortium that provides an integrated view of data and orthology for genes across the model organisms

### GO Summary Ribbons

Gene reports now include a visual summary of the associated Gene Ontology (GO) annotations

### Gene Groups

Gene groups are a new data class, bringing together genes/gene products that form a biological group (members of a gene family, subunits of a protein complex, or some other functional grouping)

### Protein Domains

The Gene Model and Products section now includes a section showing the protein domains

### Functional complementation

This section lists all orthologs for which there is experimental complementation (rescue) data in files

### New 'Dataset' data class

This section lists all associated large datasets (e.g. genetic screen, highthroughput study)

### Reference type filter

Allows selecting specific types of associated publications (e.g. Research papers, Reviews, Personal communications)

### RSS updates button

Automated updates on your favorite gene groups, genes, alleles

### Navigation panel

For easy navigation through all sections of report pages

## Sequence Downloader

## Expression tools

Allele	Insertions / Constructs	Assoc. gene	Common terms	Major tissue	Description	# Stocks	# Refs
ScarGAL4<sup>GMRP</sup>	PI(CAP-GAL4-P)	CCAP		larval CCAP neuron	Drives expression in CCAP neurons.	2	42
ScarGAL4<sup>C72</sup>	PI(SawB)C72 PI(SawB)			follicle cell	Drives expression in follicle cells from oogenesis stage 5B.	0	64
ScarGAL4<sup>P6</sup>	PI(Cp-GAL4-A)	Coat1		hemocyte	Drives expression in hemocytes and fat bodies at all developmental stages.	2	140
ScarGAL4<sup>GMR1.4</sup>	PI(CHAT-GAL4.7.4)	CHAT		cholinergic neuron	Drives expression in cholinergic neurons in larvae and adults.	4	118

\*The **FlyBase Consortium** includes: Nicholas H. Brown, Steven J. Marygold, Gillian Millburn, Giulia Antonazzo, Helen Attrill, Silvie Fexova, Phani Garapati, Alex Holmes, Aoife Larkin, Vitor Trovisco, Jose M. Urbano (FlyBase-Cambridge), Norbert Perrimon, Julie Agapite, Kris Broll, Lynn Crosby, Gil dos Santos, David Emmert, L. Sian Gramates, Kathleen Falls, Beverley Matthews, Susan Russo, Christopher Tabone, Pinglei Zhou, Mark Zytovicz (FlyBase-Harvard), Thomas Kaufman, Bryon Czoch, Josh Goodman, Victor Strelets, Jim Thurmond (FlyBase-Indiana), Richard Cripps (FlyBase-New Mexico). FlyBase is supported by a grant from the National Human Genome Research Institute at the **U.S. National Institutes of Health** (U41HG000739). Support is also provided by the **British Medical Research Council** (MR/N030117)