



Tools for Drosophila Research

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Guangzhou, China Nov 10-12, 2021

Overview

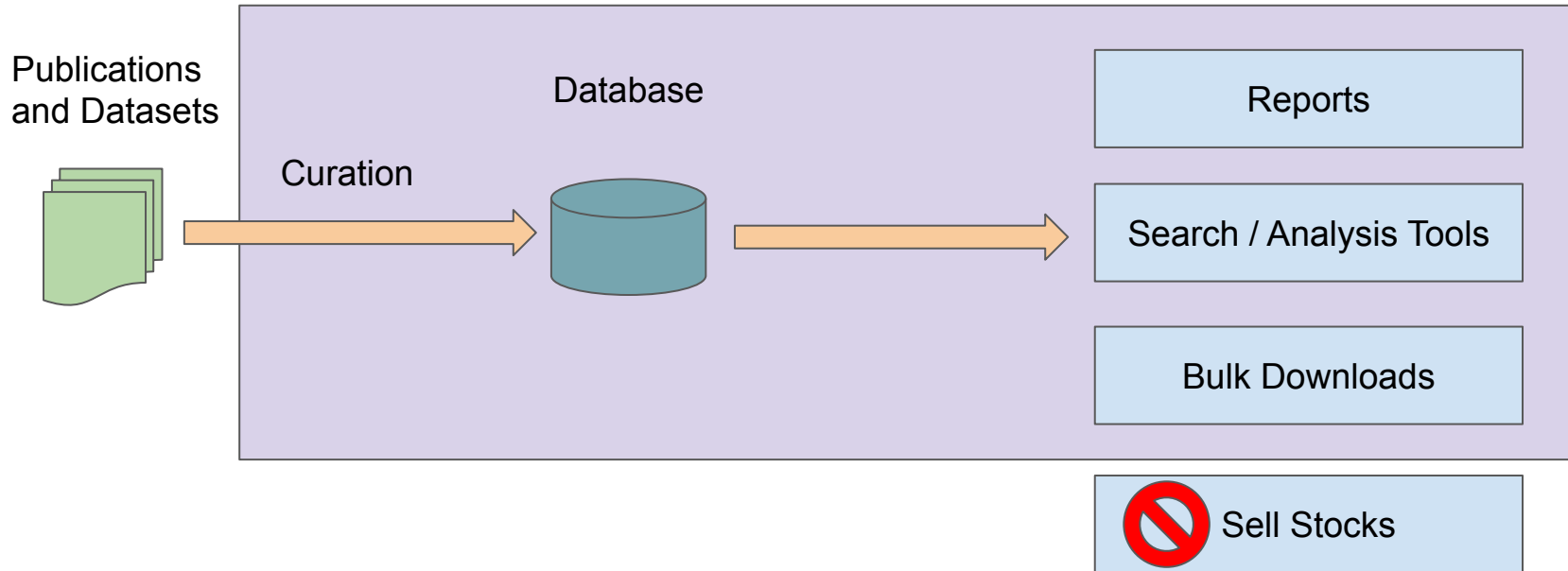
1. Introduction
2. ID Validator: Upload and validate IDs
3. HitLists
4. Fast-Track Your Paper (FTYP)
5. Current and upcoming publications
6. Staying in touch with FlyBase

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Introduction: What does FlyBase do?

Main data warehouse for 100+ years of *Drosophila* research.

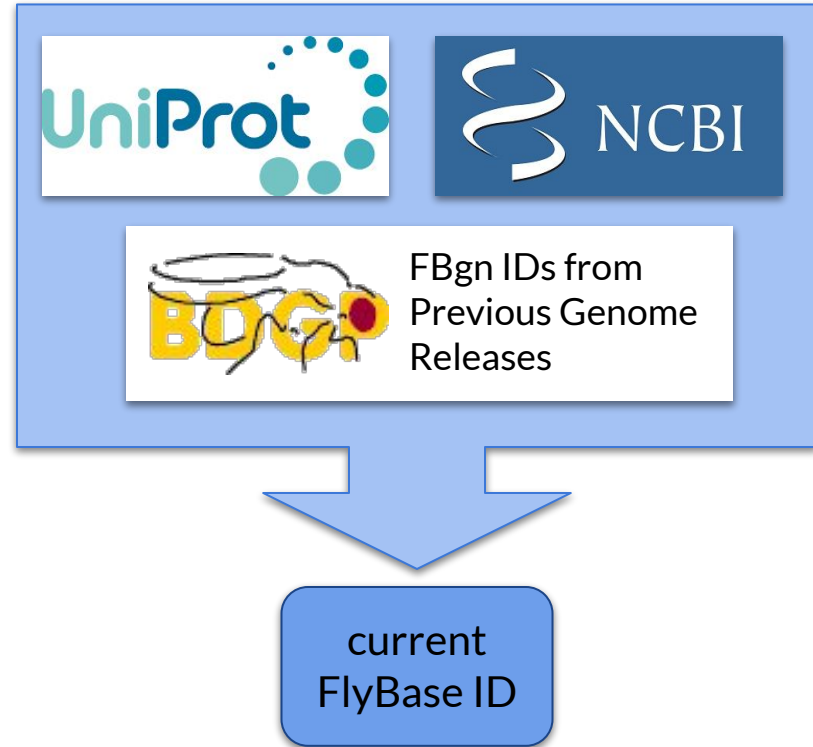
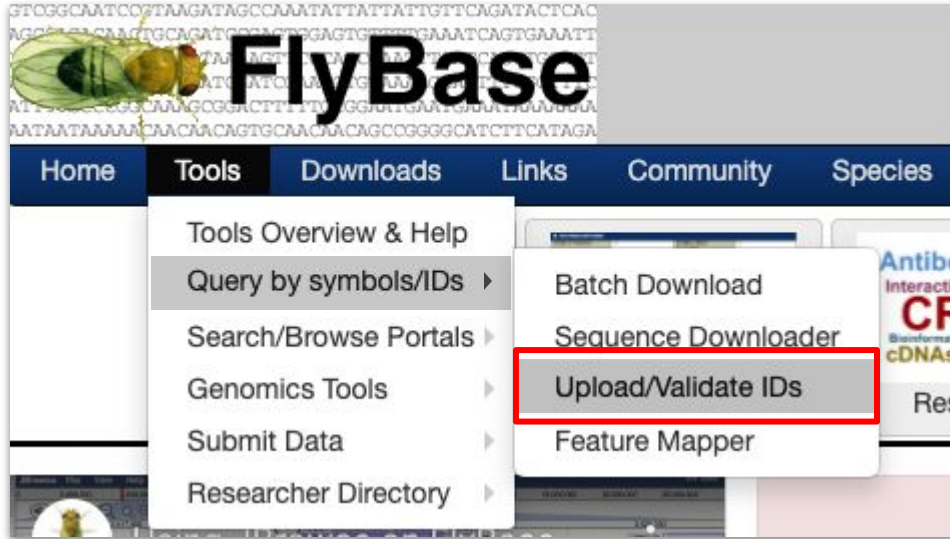


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ID validator: Upload and validate IDs

A helpful first step for batch queries and downloads.



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HitLists: Filter, sort, convert, analyze...

leucine zipper

View As

564 selected

← Previous **1** 2 3 ... 12 Next → Items
1-50 of 564

Filter by species

- D. melanogaster* (538)
- H. sapiens* (transgenes in files) (6)
- other *Drosophila* species (0)
- Other species (20)

Filter by data class

- Allele (72)
- Experimental Tool (14)
- FlyBase Ontology (1)
- Gene (65)
- Gene Group (9)
- Gene Ontology (3)
- Human Disease Model (5)
- Molecular Construct (39)
- Natural Transposon (1)
- Physical Interaction (8)
- Polypeptide (83)
- Reference (169)
- Transgenic Construct (95)

[show all](#)

Lztr1 Leucine zipper like transcription regulator 1 (CG3711, FBgn0040344) *D. melanogaster*

Feature type: protein coding gene
Sequence Location: X:922,889..927,582 [-]

Letm1 Leucine zipper and EF-hand containing transmembrane protein 1 (CG4589, FBgn0284252) *D. melanogaster*

Feature type: protein coding gene
Sequence Location: 2R:24,509,508..24,515,320 [-]

BASIC LEUCINE ZIPPER TRANSCRIPTION FACTORS (FBgg0000726)

21 genes in this group
Component groups: Parent group: [BASIC DOMAIN TRANSCRIPTION FACTORS](#)

Description >

leucine zipper domain binding (GO:0043522)

Binding to a leucine zipper domain, a protein secondary structure exhibiting a periodic repetition of leucine residues at every seventh position over a distance covering eight helical turns.

HitLists: Filter, sort, convert, analyze...

Filter by species [clear](#)

- D. melanogaster* (55)
- H. sapiens* (transgenes in flies) (5)
- > other *Drosophila* species (0)
- Other species (5)

Filter by data class [clear](#)

- Allele (72)
- Experimental Tool (14)
- FlyBase Ontology (1)
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[show all](#)

[Convert](#) [Export](#) [Analyze](#)

View As

[List](#) [Table](#)

← Previous **1** 2 Next →

selected items to:

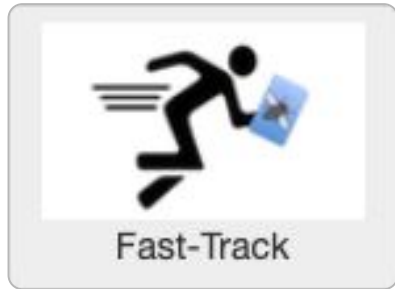
- Alleles
- Transcripts
- Polypeptides
- Insertions
- Clones
- References
- Stocks
- Sequence Features
- Human Disease Models

Gene Results			
Gene	Annotation ID	Cytology	Scaffo
leucine zipper like transcription regulator	CG3711	1D2-1D2	X
<input checked="" type="checkbox"/> <i>Letm1</i> Leucine zipper and EF-hand containing	CG4589	60D3-60D3	2R

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Fast-Track Your Paper



1 Publication — 2 Contact — 3 Data — 4 Genes — 5 Confirmation — 6 Finished

Choose a Publication to Annotate

enter key terms, author name, PubMed ID, etc.

Q Search ?

Fast-Track Your Paper guide at:

[https://wiki.flybase.org/wiki/FlyBase:Fast-Track Your Paper](https://wiki.flybase.org/wiki/FlyBase:Fast-Track_Your_Paper)

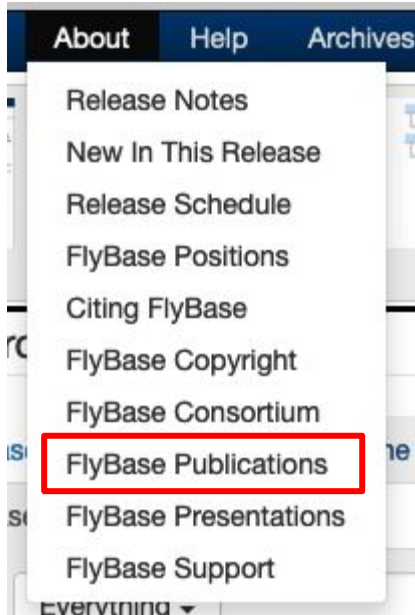
Links to papers where authors marked the “contains technical advance” box can be found at:

[https://wiki.flybase.org/wiki/FlyBase:Papers with technical advances](https://wiki.flybase.org/wiki/FlyBase:Papers_with_technical_advances)

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Current and in press publications



Current publications:

- Larkin A, Marygold SJ, Antonazzo G, Attrill H, dos Santos G, Garapati PV, Goodman JL, Gramates LS, Millburn G, Strelets VB, Tabone CJ, Thurmond J and the FlyBase Consortium (2021)
FlyBase: updates to the *Drosophila melanogaster* knowledge base.
Nucleic Acids Res. 49(D1) D899–D907 [DOI:10.1093/nar/gkaa1026](https://doi.org/10.1093/nar/gkaa1026)
- Marygold SJ, Crosby MA, Goodman JL and The FlyBase Consortium. (2016)
Using FlyBase, a Database of *Drosophila* Genes and Genomes.
Drosophila: Methods and Protocols, Second edition, vol. 1478 (C. Dahmann, ed.) pp. 1-31. Springer, New York [DOI:10.1007/978-1-4939-6371-3_1](https://doi.org/10.1007/978-1-4939-6371-3_1)

In press publications:

- Next edition of Drosophila: Methods and Protocols
- Genetics database issue.

Tutorials from @FlyBaseDotOrg

FlyBase @FlyBaseDotOrg · Aug 24, 2018
FlyBase has updated the 'Search FlyBase' QuickSearch tab. You can get the functionality you're used to, searching all of FlyBase by text string by choosing the (default) 'Everything' option, or narrow your search to only IDs, symbols & names.

QuickSearch

Human Disease GAL4 etc Expression Phenotype References
Search FlyBase Orthologs Protein Domains Gene Groups GO Data Class

Everything ▾ FBto*
ID, Symbol, Name
Everything

Commentary list

1 6 7

FlyBase @FlyBaseDotOrg · Aug 24, 2018
For example, we use the string 'FBto*', the identifier prefix for the new Experimental Tool reports. When 'Everything' is selected, the hitlist includes every FlyBase report with the string 'FBto' anywhere in it, while 'ID, Symbol, Name' includes only Experimental Tool reports.

FBto* 9475 selected
Filter by species
Filter by data class

FBto* 388 selected
Filter by species
Filter by data class

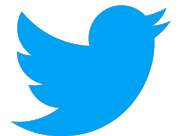
1 1 1

[#FlyBaseTweertorial](#)

FlyBase @FlyBaseDotOrg · Jan 7
Want to find a tweertorial? You can search Twitter using the hashtag [#FlyBaseTweertorial](#) + a hashtag for your feature of interest; this narrows your search to only tweets using both hashtags. The next few linked tweets alphabetically list all the FlyBaseDotOrg hashtags to date.

#FlyBaseTweertorial #paralogs

Search for "#FlyBaseTweertorial #paralogs"



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Staying in touch with FlyBase



FB2021_01, released Feb 18, 2021

A Database of *Drosophila* Genes & Genomes

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- FlySlack Community
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- FlyBase Presentations
- FlyBase Support

- Video Tutorials ↗
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- Report help
- Info for Authors ▶
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version FB2021_01, released Feb 18, 2021
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