

GMOD @ Arthropod Genomics Symposium

GMOD Users

wFleaBase
BeetleBase
FlyBase
AphidBase
ButterflyBase
VectorBase
BeeBase
HeliconiusBase
GnpAnnot-Lep
+ hundreds more

GMOD Tools

Chado
Apollo
GBrowse
CMap
Table Editor
GMODWeb
Ergatis
Textpresso
Sybil
BioMart
LuceGene
SynBrowse
Galaxy
InterMine
...



<p>Thursday 7-9pm Community Contributions to Genome Annotation Christine Elsik, Christopher Childers, Darren Hagen</p> <p>Friday 7:30-9:30pm Chado Databases and Integration with GMOD Tools Scott Cain, Dave Clements</p>	Workshops
<p>Friday 11:25-11:45am Unlocated arthropod genes and ways to find them Don Gilbert</p>	Talks
<p>Friday 5-6:30pm A comparative annotation of Drosophilid dicistronic genes</p> <p>Saturday 10-11:30am A Complete System For Community Genome Annotation GMOD: Database Resources for Emerging Model Organisms Comparative genomics and database construction for Lepidoptera Unlocated Arthropod genes, and ways to find them VectorBase: A genome resource for arthropod vectors of human pathogens The VectorBase Manual and Community Annotation Submission pipeline ButterflyBase: A framework for comparative genomics in butterflies and moths</p>	Posters

<http://gmod.org>

Got data?