

# 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><b>Thursday 7-9pm</b> <b>Community Contributions to Genome Annotation</b> Christine Elsik, Christopher Childers, Darren Hagen</p> <p><b>Friday 7:30-9:30pm</b> <b>Chado Databases and Integration with GMOD Tools</b> Scott Cain, Dave Clements</p>	<b>Workshops</b>
<p><b>Friday 11:25-11:45am</b> <b>Unlocated arthropod genes and ways to find them</b> Don Gilbert</p>	<b>Talks</b>
<p><b>Friday 5-6:30pm</b> A comparative annotation of Drosophilid dicistronic genes</p> <p><b>Saturday 10-11:30am</b> 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>	<b>Posters</b>

<http://gmod.org>

Got data?