

# GMOD @ Plant and Animal Genome XIX

## GMOD Users

Gramene  
 RAP-DB  
 ConiferGDB  
 TAIR  
 MaizeGDB  
 SGN  
 TreeGenes  
 SoyBase  
 CottonDB  
 MOCCAdb  
 GDR  
 GnpIS  
 + hundreds more

## GMOD Tools

Chado  
 Apollo  
 GBrowse  
 CMap  
 Tripal  
 GBrowse\_syn  
 Ergatis  
 Textpresso  
 BioMart  
 JBrowse  
 MAKER  
 Galaxy  
 InterMine  
 ...



<p>Database Resources at the EBI: Ensembl (including BioMart)          Sunday, 8:00-9:00am, California Room, Bert Overduin</p> <p>The Generic Genome Browser: A Hands on Workshop          Sunday, 1:30-3:40pm, California Room, Scott Cain</p> <p>Using Gramene (including CMap)          Tuesday, 3:50-6:00pm, California Room, Ken Youens-Clark</p>	<p><b>Workshops</b></p>	
<p><b>MAKER: An easy to use genome annotation pipeline</b>          Wednesday, 10:30-11:30am, Pacific Salon 3, Carson Holt</p> <p><b>Galaxy: Analyze, Visualize, Communicate</b>          Wednesday, 11:30am-12:30pm, Pacific Salon 3, Dan Blankenberg</p> <p><b>JBrowse: A Next Generation Genome Browser</b>          Wednesday, 1:30-2:30pm, Pacific Salon 3, Mitch Skinner</p> <p><b>Tripal: A Construction Toolkit for Online Genomic Databases</b>          Wednesday, 2:30-3:30pm, Pacific Salon 3, Stephen Ficklin</p> <p><b>GMODviews: A Configurable Interface for GMOD Databases &amp; Tools</b>          Wednesday, 3:45-4:00pm, Pacific Salon 3, Michael Caudy</p> <p><b>GBrowse_syn: The Generic Synteny Browser</b>          Wednesday, 4:00-5:00pm, Pacific Salon 3, Robert Buels</p>		<p><b>GMOD:          Organize          Annotate          and          Visualize          Your Data</b></p> <p>Wednesday          10:20 - 5:00          Pacific Salon 3</p>
<p><b>Developing a Comprehensive Genomics, Genetics &amp; Breeding DB for Cacao</b>          Sunday, 3:45-4:00pm, Pacific Salon 6 &amp; 7, Ping Zheng</p> <p><b>How to Use MetaCyc and the Pathway Tools Software in Local Mode</b>          Sunday, 4:30-4:50pm, California Room, Lukas Mueller</p>		<p><b>Talks</b></p>
<p>P798: CottonDB: An Integrated Repository Resource for Cotton</p> <p>P800: Genome Annotation and Report System: An Integrated System for Structural &amp; Functional Genome Annotation</p> <p>P805: GnpIS, A Portal and Information System to Bridge Genetics and Genomics Plant &amp; Fungi Data</p> <p>P814: Informatics for Turbo-charged Comparative Biology: Highlights from the GMOD Evo Hackathon</p> <p>P818: A Complete Community Annotation System using GMOD Tools</p> <p>P828: GBrowse 2.2: Fast, Friendly, and Sociable</p> <p>P888: GDR (Genome Database for <i>Rosaceae</i>) Building Resources for Translational Research in <i>Rosaceae</i></p>		<p><b>Posters</b></p>
<p><i>and at least 8 other talks and 9 more posters by GMOD users (see <a href="http://gmod.org/wiki/PAG_2011">http://gmod.org/wiki/PAG_2011</a>)</i></p>		<p><b>+</b></p>

<http://gmod.org>

Got genome?