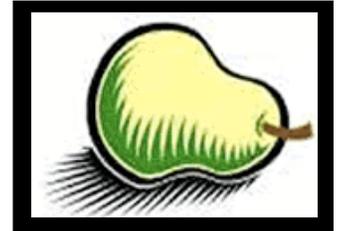




# @ ISMB & BOSC 2008

<http://gmod.org>



**Intermine - Open source Data Warehouse and Query Interface**  
 Richard Smith, Fri 11:10-11:35

**Ergatis: A Web-based Bioinformatics Pipeline Management and Collaborative Development System**  
 Joshua Orvis, Sat 9:30-9:55

**Apollo: a Sequence Annotation Editor**  
 Ed Lee, Sat 11:50-12:10

**EMBRACE - BioMart Developments and Future**  
 Syed Haider, Sat 1:50-2:10

**BOSC Talks**

**EMBRACE - Enabling Data Integration**  
 Syed Haider, Sun 2:15-2:40

**Intermine - Open source Data Warehouse and Query Interface**  
 Richard Smith, Mon 12:15-12:40

**Pathway Tools Software: Regulatory Networks and Recent Developments**  
 Peter Karp, Mon 3:15-3:40

**Galaxy Workflows: Enabling Complex Computational Analysis For Experimental Biologists**  
 James Taylor, Tue 3:15-3:40

**New GALAXY Tools**  
 Anton Nekrutenko, Tue 3:45-4:10

**ISMB Talks**

**Sun 5:45-8:30pm**  
 A Genome Grid for Finding new Bug Genes  
 A Complete System for Community Genome Annotation  
 Galaxy workflows for reproducible analyses of next-generation sequencing data, Anton Nekrutenko

**Mon 5:45-8:30pm**  
 Metagenomic Analysis with Galaxy:  
 Windshield Genomics and Beyond  
 EMBRACE – Enabling Data Integration  
 GMOD: Managing genomic data from emerging model organisms

**Posters about GMOD**

**Sun 5:45-8:30pm**  
 Multidimensional annotation of the Escherichia coli K-12 genome  
 Towards Data Analysis of Genome Wide Influenza RNAi Screen  
 Bioinformatics to support the sequencing of Brassica rapa chromosomes A2 and A10

**Mon 5:45-8:30pm**  
 Regional co-variation among rates of different types of mutations  
 Solutions for database interoperability: a report from the CASIMIR consortium

**Posters using GMOD**