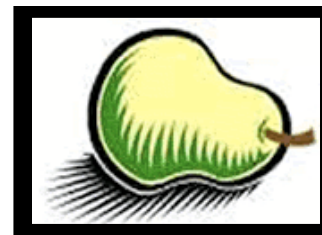




# GMOD @ ISMB & BOSC 2008

<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

GMOD: Managing genomic data from emerging model organisms

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

### 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