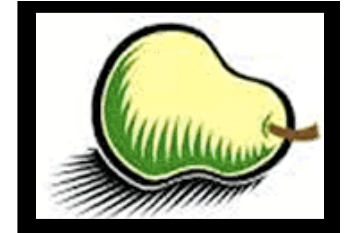




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<http://gmod.org>



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

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

BOSC Talks

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