

GMOD Help Desk

Dave Clements



GMOD Help Desk

- What I've been doing
- What I'm planning on doing
- What should I be doing?
- How am I doing?



What I've been doing

- Web site
 - Tags/Categories
 - Tagged every page
 - Made it easy to tag pages, lookup pages by tag
 - Pages for new users
 - Glossary, GMOD and Databases, ...
 - Community Portal
 - News, Calendar, Recent Changes
 - Need your help



What I've been doing

- Outreach
 - Posters:
 - Genome Informatics, IGERT EvoDevo, Arthropod Genomics, Genetics of Behavior, ISMB
 - Representation
 - Lepidoptera and ArthropodBase; U Oregon
 - Promotion
- Education
 - Chado Workshop with Scott @ Arthropod Genomics
 - ~35 people
 - GMOD Summer School with Scott, Ed, Ben,
 - 25 students for 2 1/2 days
- Components
 - MAKER, Phenote





@ 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

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

What I'm planning

- GMOD User Directory
 - How
 - UseTableEdit
 - Each user has a page
 - Each component lists users of that component
 - Comprehensive user list.
 - All driven by data on each user page.
 - Why
 - Community portal
 - Give new users an idea of who is doing something similar
 - I'll be in the neighborhood



Experience Logs

- Can't possibly describe or maintain HOWTOs for all OS/software combos.
- Create organization for including, encouraging and tagging user experience logs
 - A few already on the web site.
 - No maintenance involved with this.
 - Just catalog as we try them.



Documentation

- Chado doc reorganization
 - Including Chado Cookbook / Best Practices
 - Mine gmod-schema mailing list
 - Start user directory
- GBrowse Doc
 - GBrowse Glyphs
 - GBrowse Cookbook?
- Community Annotation Server Doc



Documentation 2

- Tutorials
 - Screencasts for sophisticated user interfaces
 - GBrowse, Apollo, CMAP
- Web site
 - Mediawiki upgrade
 - TableEdit
 - Better searching
 - New skin?



Components

- Galaxy, InterMine, BioMart
 - Seek tighter integration with rest of GMOD
- TableEdit
 - Documentation
 - Testing - will integrate with GMOD web site
 - Could be key to (non-sequence) community annotation
- Natural Diversity Chado Module
 - Get it integrated, documented and used
- Google 2009 Summer of Code?
- GMOD Logo Service?



Education & Outreach

- 2009 Summer School
- GMOD Course in Europe?
- Conferences
 - Tutorial at Plant and Animal Genomics?
 - Arthropod Genomics
 - IPlant?
 - More posters
- Particularly promote
 - Comparative Genomics
 - Community Annotation



Grants, Reviews & Funding

- I now offer to review grants that propose using GMOD
- Encourage grant writers to include funding for GMOD
 - Fund general project or specific components
 - Could increase stability of GMOD



What should I be doing?

?



How am I doing?

Don Gilbert is doing a review of my first year in August and September.

Please contact him if you any feedback.



Thank You!



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

<http://gmod.org>

Got data?