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

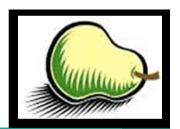








@ ISMB & BOSC 2008



http://gmod.org

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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

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

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

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

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?

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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!



Dave Clements GMOD Help Desk

315 Pacific Hall National Evolutionary Synthesis Center clements@nescent.org help@gmod.org

http://gmod.org/GMOD_Help_Desk http://nescent.org



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

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InterMine

Community Contributions to Genome Annotation

Christine Elsik, Christopher Childers, Darren Hagen

Friday 7:30-9:30pm

Chado Databases and Integration with GMOD Tools

Scott Cain, Dave Clements

Friday 11:25-11:45am

Unlocated arthropod genes and ways to find them

Don Gilbert

Friday 5-6:30pm

A comparative annotation of Drosophilid dicistronic genes

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

Talks

Workshop

Posters



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