

## Digesting the Genome Glut Promoting the Use and Extension of GMOD

To Emerging Model Organisms



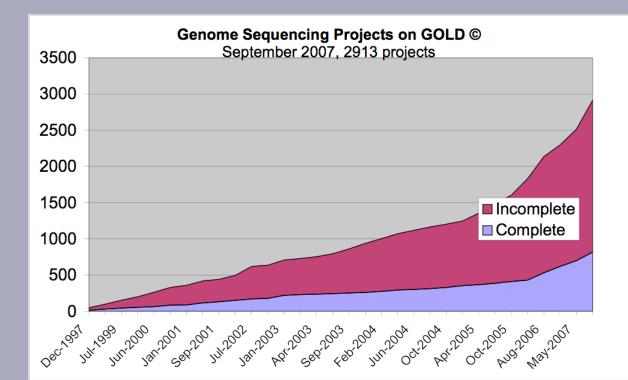
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## Abstract

An ever-increasing volume of genomic data is being generated from organisms well suited to comparative genomic, evolutionary and ecological studies. More data offers more potential for discovery, but it also makes it is harder to organize, visualize and annotate.

The Generic Model Organism Database (GMOD, http://gmod.org) is a toolkit of interconnected components that meet many of the data management, analysis, and visualization needs of these studies, and on a small informatics budget.



GMOD has recently added a helpdesk to promote use of GMOD more broadly in emerging model organism research communities. GMOD has also been extended to better support evolutionary research.



In the past year, the GMOD web site has been revamped and a GMOD Helpdesk (help@gmod.org) has been created.

## What Can GMOD Do?

GMOD is a collection of software tools for creating and managing genome-scale biological databases. You can use it to create a small laboratory database of genome annotations, or a large webaccessible community database. GMOD includes a modular database schema called Chado that supports many common needs.

If you have	& you need to	then GMOD can help with
Genomic Sequence	View	GBrowse
	Annotate	Apollo
	Organize	Chado Sequence Module
Physical, Genetic & Comparative Maps	View	CMap,SynView,SynBrowse,Sybil
	Annotate	Sybil
	Organize	Chado Map Module
Phylogenetic Trees	Organize	Chado Organism & Phylogeny modules
	View	GMODWeb
Phenotype,	Organize	Chado Genetics Module
Genotype	View	GMODWeb
Microarray and	View	Java TreeView, Caryoscope, GeneXplorer
Expression	Organize	Chado Mage and Expression Modules
Ontologies	Curate	Chado CV module
	View	GO Term Viewer, GMODWeb
Phylogenetic Trees	Organize	Chado Organism & Phylogeny modules
	View	GMODWeb
Publications	Curate, search	TextPresso, PubSearch
	Organize	Chado Publication Module

The National Evolutionary Synthesis Center (NESCent) Sponsors Synthetic, Interdisciplinary and Transformative Research in Evolutionary Biology.

The center hosts working groups, catalysis meetings, sabbaticals, postdoctoral fellowships, and a shortterm visitor program. NESCent's informatics branch undertakes initiatives in support of software interoperability, data sharing, and informatics training. The center also solicits whitepapers for projects that fall outside the scope of existing programs. More information on how to apply or become involved at:

http://www.nescent.org

at Chapel Hill.



NESCent is supported by the National Science Foundation (Grant # EF-0423641)

NESCent is located in Durham

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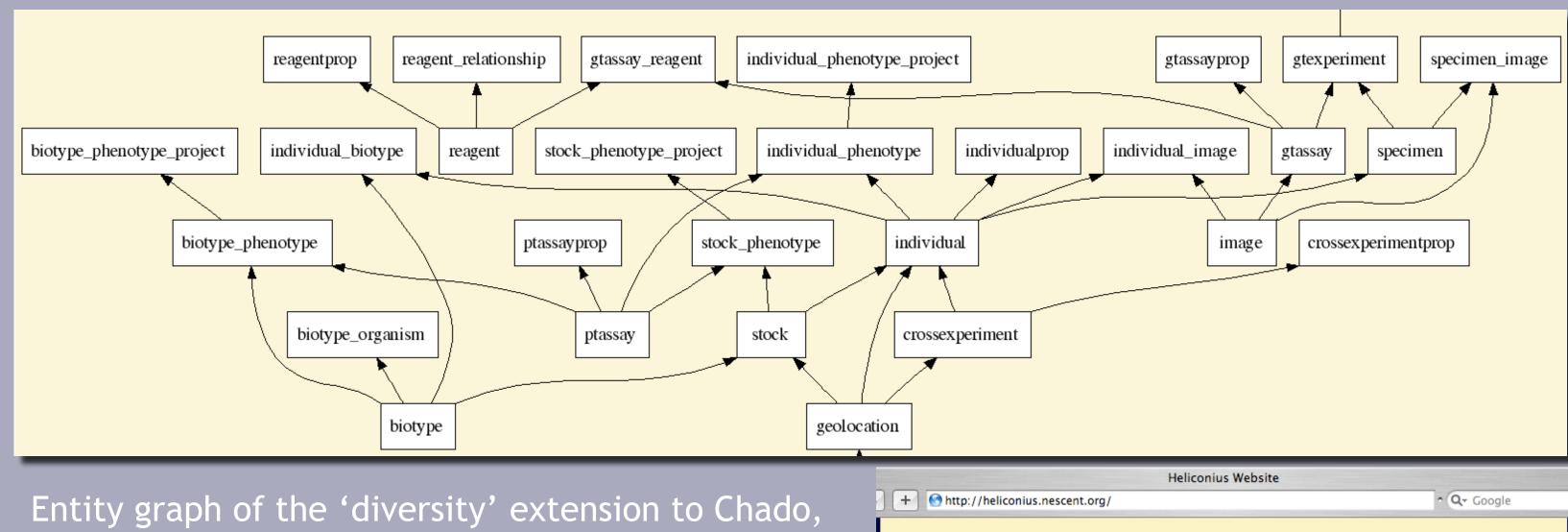
North Carolina State

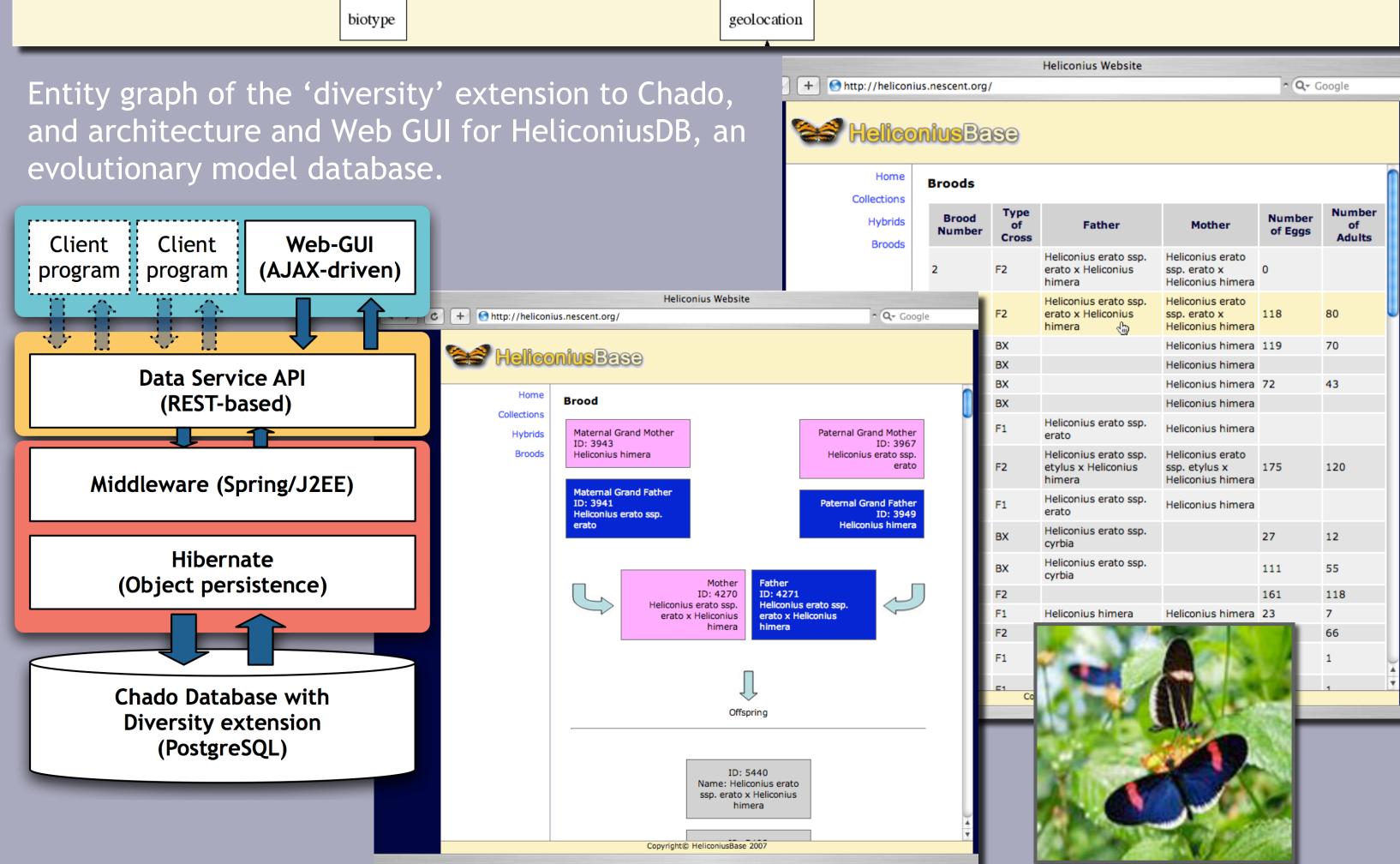
University, and the

## GMOD for Natural Diversity Data

Heliconius butterflies are a favored model for studying mimicry, because geographically overlapping populations of different species frequently converge on similar wing color patterns. Complementing the long history of evolutionary genetic studies in the genus, an increasing amount of molecular marker and sequence data are being collected.

To capture the diversity data in a GMOD-compatible way, Chado (GMOD's relational data model) has been extended with the 'diversity' module (see http://sf.net/projects/heliconiusdb). The module supports stocks, individuals, pedigrees, cross experiments, geolocations, and phenotype and genotype experiments. In addition, some of Chado's core modules have been revised to consolidate how organism relationships and phylogenetic trees are represented.







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