Towards a Web of Distributed Registries: JBEI Registry of Biological Parts

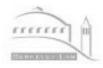
Timothy Ham
29 April 2009

Advances in Synthetic Biology















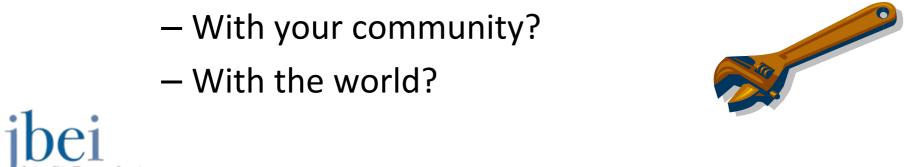




Synthetic Biology Software: **Current Status**

- MS Excel
- MS Word
- VectorNTI / APE / DNA Star

- How to share your data?
 - With your group?
 - With your community?

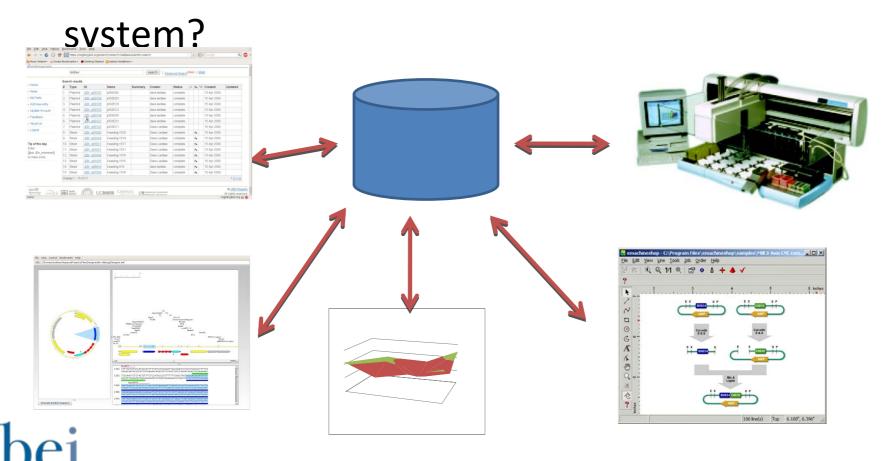




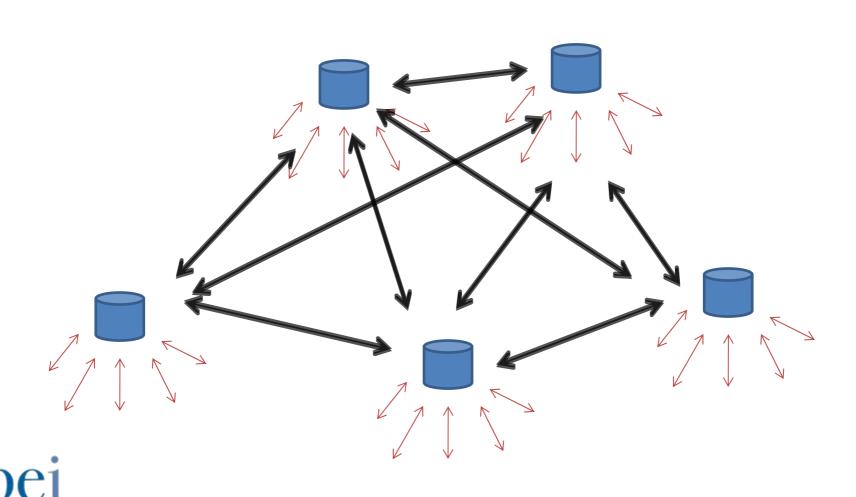
Synthetic Biology Software: Goals

Blank Canvas. What would be an ideal

Joint BioEnergy Institute



Synthetic Biology Software Goals



MIT Registry

- Centralized
 - One Registry to Rule Them All
 - All parts are public
- Wiki based
 - Easy to add pictures and descriptions
 - Hard to automatically organize information
- Rapid Growth
 - iGEM contributions



MIT Registry

- Too Brick-Centric
 - No concept of plasmids or strains, outside of brick context
 - Brick as both a basis of unit as well as "kind" of a part
- Software is not available



JBEIR: Goals

- "Backward" compatible
 - Support for strains, plasmids, as well as parts
- Distributed
 - Software is Open Source (BSD-style license)
 - Allows individual labs to run their own registry
- Encourages sharing of data
 - Core Data Model will be published with Clotho (UC Berkeley BioCAD) group.



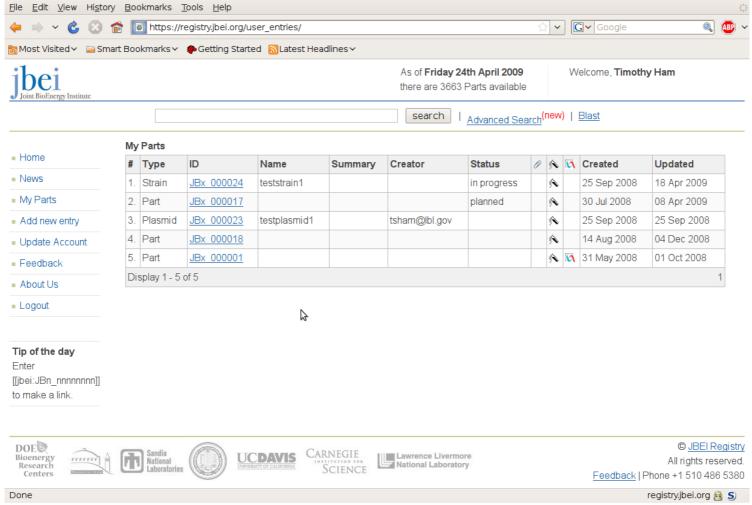


JBEIR: Status

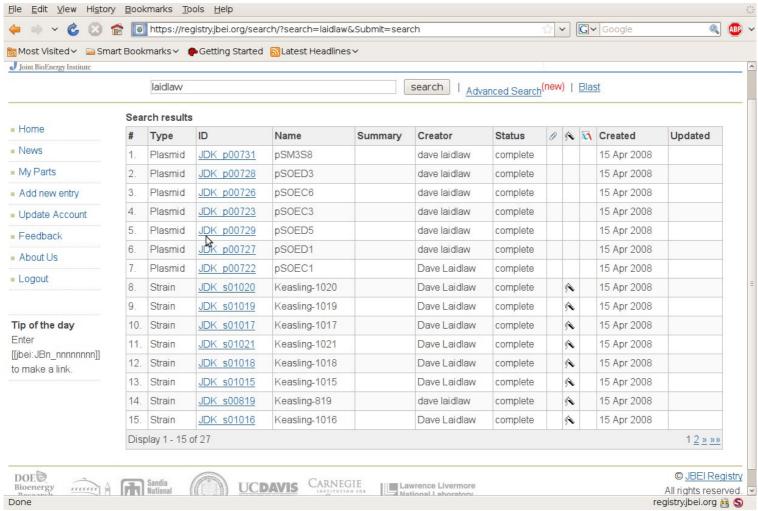
- 3500+ parts
- 200+ new parts since last year
- Advanced searches
 - Google-like full text search
 - Field based search
 - BLAST search



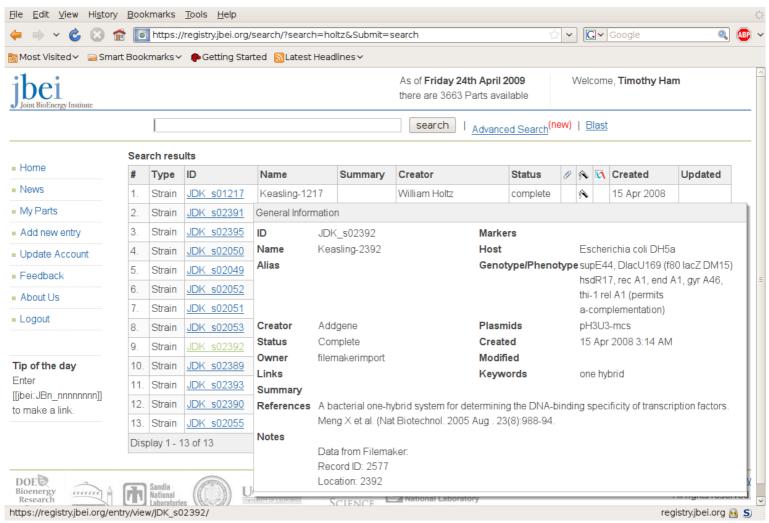




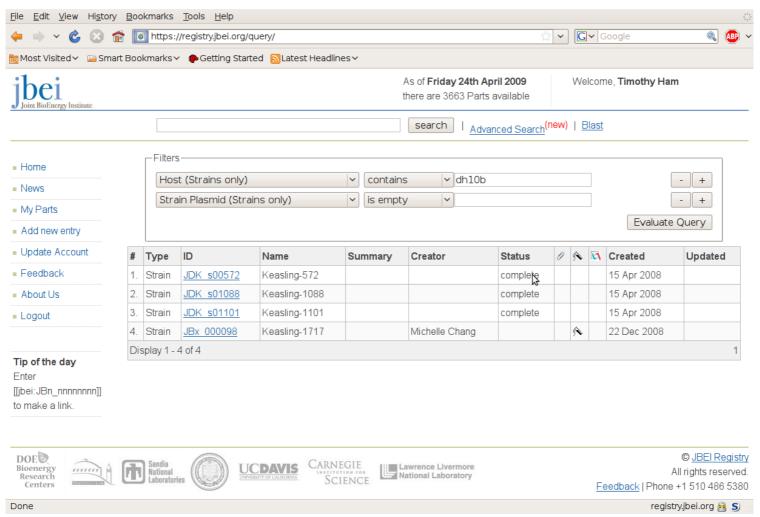




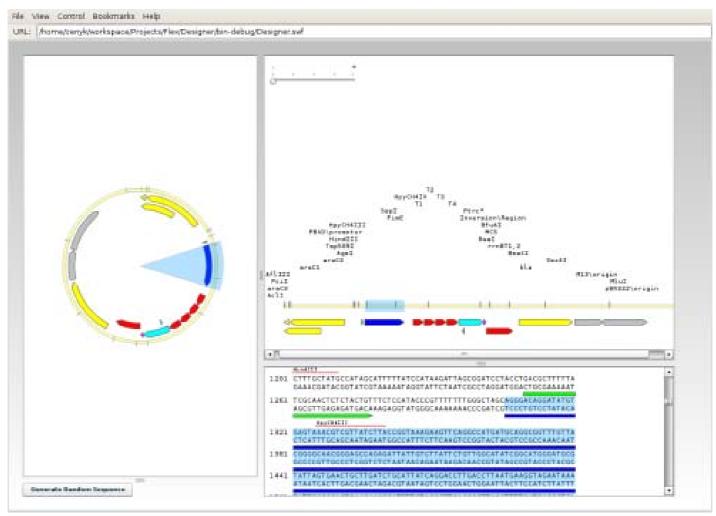






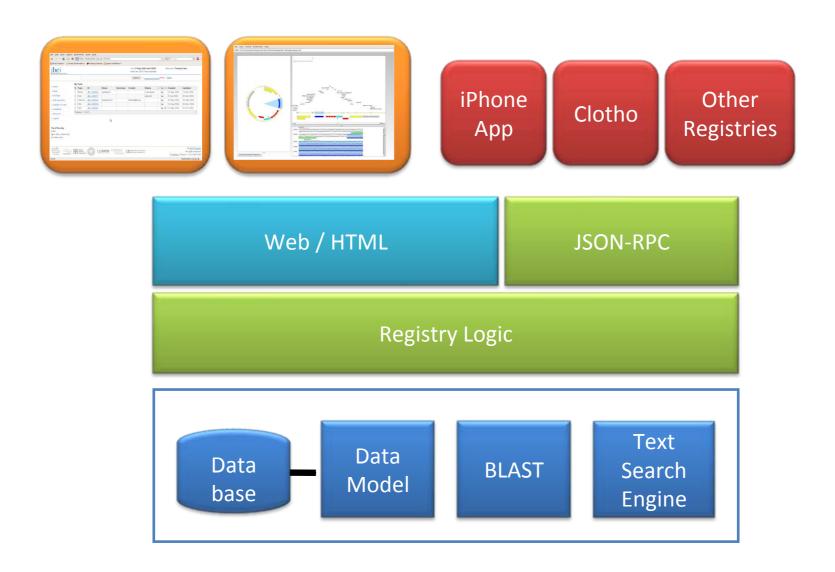








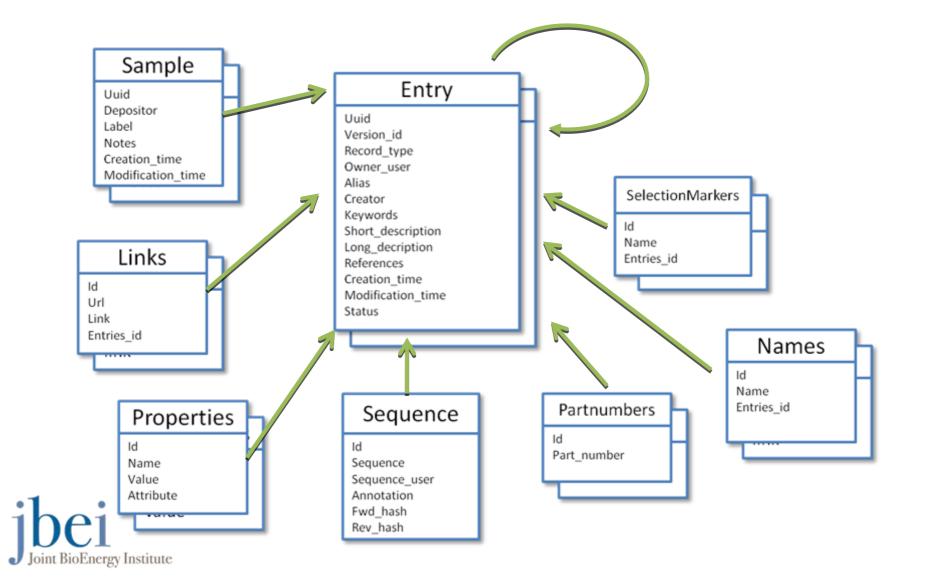
JBEIR: Architecture



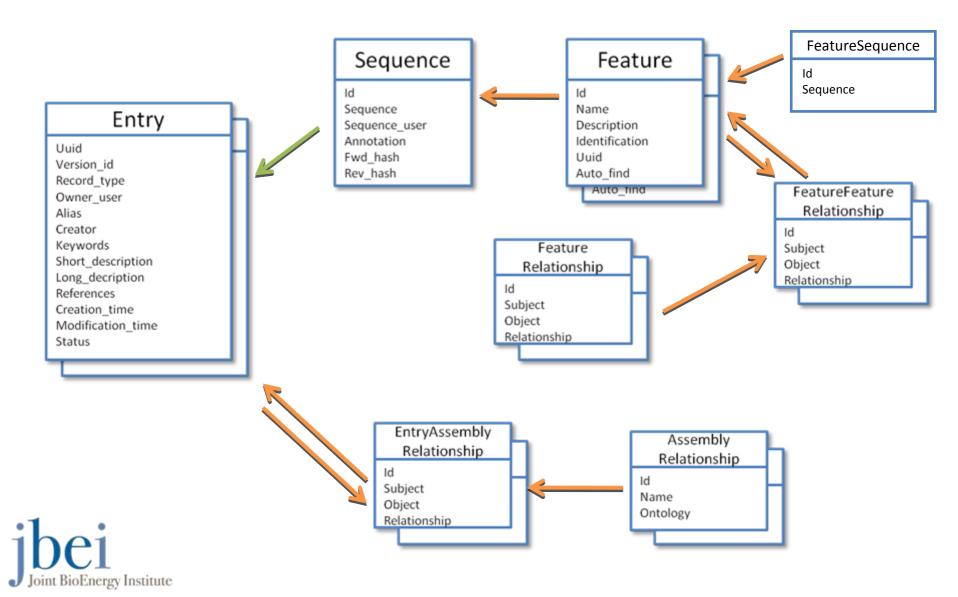
JBEIR: Architecture

- Capabilities
 - Modular brings flexibility
 - Web accessible, both by browser, and through Javascript, RPC, and REST.

JBEIR: Data Model



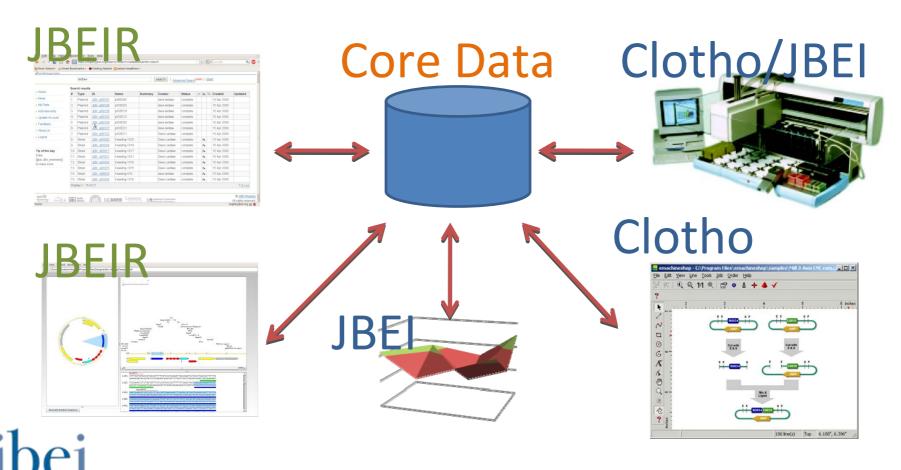
JBEIR: Ontology Model



JBEIR: Data Model

- Advanced Capabilities
 - Strains can have multiple plasmids
 - Plasmid can be in multiple strains
 - Can determine all strains with a given plasmid
 - Determine construction lineage: derived parts, sibling parts (mutations), etc.
 - Determine Families/Categories:
 - Lac Promoter -> Promoter -> Transcriptional Regulator
- Draft "Core Data" RFC in progress

Synthetic Biology Software: Goals



Joint BioEnergy Institute

JBEIR: Current Work

- Getting ready for a public release. Coming this summer.
 - Set up your own instance of the registry.
 - Contact me for a preview release.
 - Public Registry to deposit your parts.
- Share data with other software
 - Clotho, Synthetic Biology Core Data Model
- Integration of graphical annotation tools
 - This Summer

Thank you

Thanks goes out to:

- Zinovii Dmitriv
- Blake Simmons
 - Paul Adams
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 - Will Holtz
 - Swap Chhabra
 - Jay Keasling
 - JBEI