### Sage Bionetworks Mission

Sage Bionetworks is a non-profit organization with a vision to create a "commons" where integrative bionetworks are evolved by contributor scientists with a shared vision to accelerate the elimination of human disease



Michael Kellen Director of Technology

# Need: Community-based analysis is required to build accurate models of disease



#### Within the analytical community:

- Quality assessment of data, tools and models
- Reproducibility of models
- Reusability of data
- Validation of models across multiple disease models and patient subtypes

#### Across the field of biology:

- Validation of model predictions in experimental/clinical setting
- Accelerated pace towards improved therapeutic development

### Solution: Sage Bionetwork Synapse Project

#### Watch What I Do, Not What I Say

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Real-time tools to track data analysis projects

#### Most of the People You Need to Work with Don't Work with You



An online community to connect people and projects

#### Reduce, Reuse, Recycle

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A Repository of Data, Tools, and Models

#### My Other Computer is Amazon



Cloud-based data stores and compute resources 3

# Sage Metagenomics Project



- > 12K genomic and expression standardized datasets indexed in SCR
- Error detection, normalization in mG
- Access raw or processed data via download or API in downstream analysis
- Building towards open, continuous community curation

### Sage Metagenomics using Amazon Simple Workflow



Full case study at http://aws.amazon.com/swf/testimonials/swfsagebio/ 5

## Use of Ontologies in Synapse

- Synapse UI helps users annotate data / other objects in system with right concepts
- Automated data ingestion pipelines autogenerate appropriate annotations
- Ontologies drive faceted search
- Future development: consistency of clinical trail variables across studies
- Future development: integration with external indices

## **Ontologies in Synapse**

#### **Ontology Benefits**

- + Consistency of term use across many data sets facilitates large-scale meta analysis (e.g. finding gene expression signatures across all GEO prostate cancer studies)
- + Users can find public data to integrate with their own private studies.
- + Increase ability to integrate across systems (future)

#### **Ontology Difficulties**

- End users are the domain experts, but can't articulate everything they need up front
- End users can not understand complex ontological constructs e.g. OWL class restrictions / upper ontology schemes
- Many ontologies of dubious quality: incomplete term coverage; definition of concepts not mapping to terms as they are used in practice by people.

### Recommendations

- ⇒ Expect iterative refinement of ontology and system using it (we are still early in process)
- ⇒ Expose users to SKOS semantics; use more complicated constructs only on back end if necessary.
- ⇒ Service-oriented architectures allow separation of code and ontology updates.
- ⇒ Need to establish feedback mechanisms from end users to ontology designers directly from point of use.