Biodiversity science: A new, unique cyberinfrastructure challenge: Or familiar, generic problem space?

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SEAIP Biodiversity

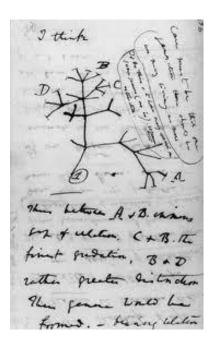
Biological Diversity

 Biodiversity: the variety of all forms of life, from genes to species, through to the broad scale of ecosystems.









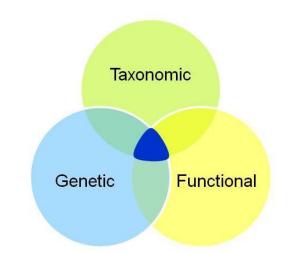
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Elements ->> Process

- "atomistic" bias of western culture towards objects.
- Has biodiversity science been overly focussed on "inventory" of species, genes, ecosystems and has neglected processes that create and maintain natural values.
- Cf. reductionist focus on model species (Arabidobsis, mouse, human, etc.)

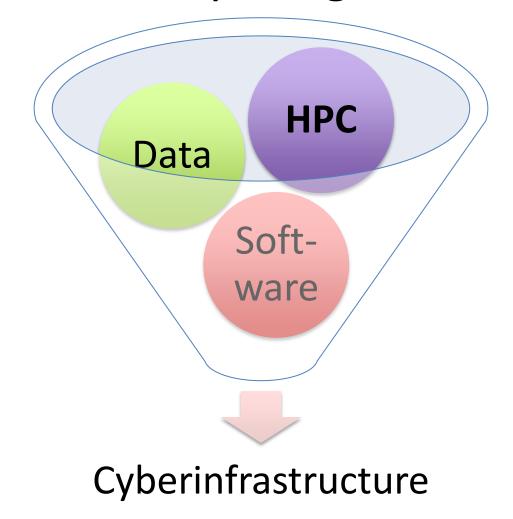
Biodiversity: the broader view

- NSF Dimensions of Biodiversity Program (cross-directorate)
- How many dimensions are there?



Systematics, taxonomy, evolution, biogeography, ecology, population genetics, genomics, metagenomics....

What does this have to do with computing?



Phylogenetic cyberinfrastructure

Sequencing facility

NCBI

GenBank

- CIPRES
- AToL (NSF program)
- iPlant/iPTOL,
- AToL, AvaToL



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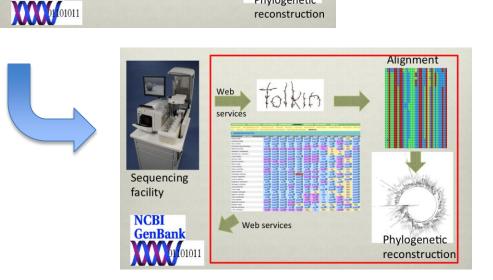
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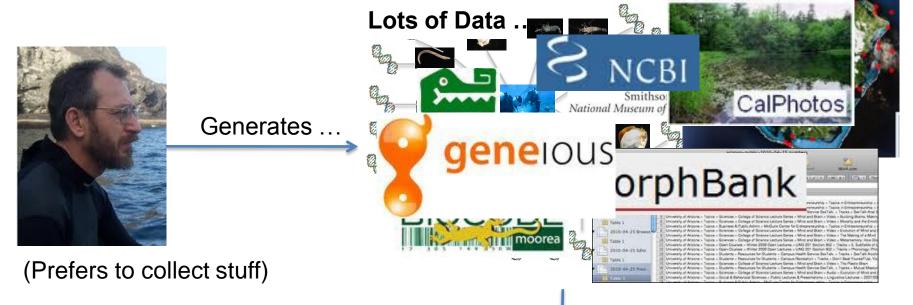
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Phylogenetic

- High Throughput sequencing
- Genes to Genomes

Data? Gustav's Problem....

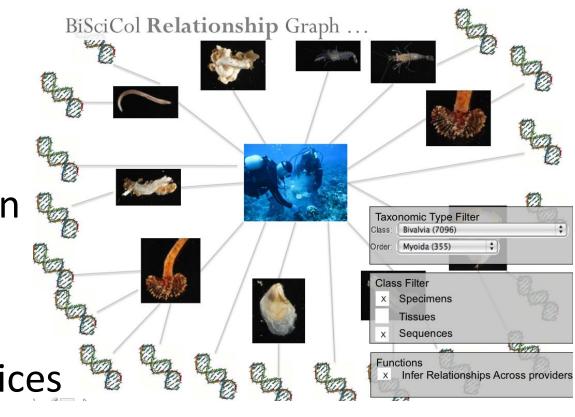




Due to project requirements and integration needs, Gustav is left navigating a plethora of redundant and disconnected distributed Databases. Lots of effort to track objects And their derivatives.

Knowledge networking challenge

- Ontologies and semantics
- Data curation
- Genotype to phenotype
- Gene expression and function
- Ecotypes
- Ecosystem services



250 years of Biological Ontologies



Chapter 1: Philosophy and Biomedical Information Systems Barry Smith and Bert Klagges

1. The New Applied Ontology





Each partial category system will divide its domain into classes, types, groupings, or kinds, in a manner analogous to the way in which Linnaeus's taxonomies divided the domain of organisms into various upper-level categories (kingdom, phylum, class, species, and so forth), now codified in works such as the *International Code of Zoological Nomenclature and the International Code of Nomenclature of Bacteria*.

Storage and Archiving: 250 years of documenting biodiversity

- Scientific collections (Musuems and Herbaria)
- ca. 2.5 billion collections objects (specimens) world-wide
- Documents 2 10 million species
- NSF ADBC program (Advancing Digitization of Scientific Collections): National HUB and Thematic Networks







National Resource:



- Making collections accessible
- Enabling answers to scientific questions
- Information integration across collections and across domains
- Adding value to collection data
- Digital data capture and
- management





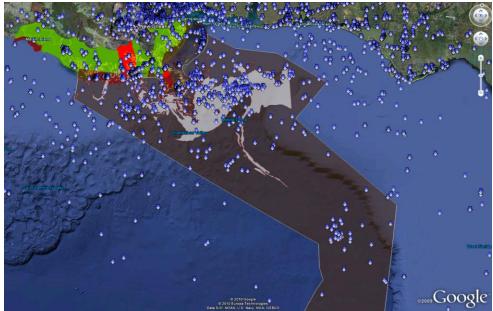


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Societal Challenge

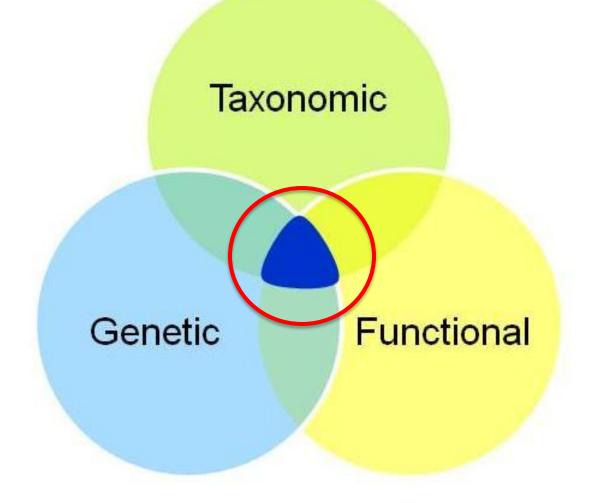
Oil Spill response

- How are we doing in terms of measuring effects on biodiversity?
- What information infrastructure is available for comparison to baselines, modeling, impact, and assessment?
- Data problem: Sensor networks, crowdsourcing, specimens/observations

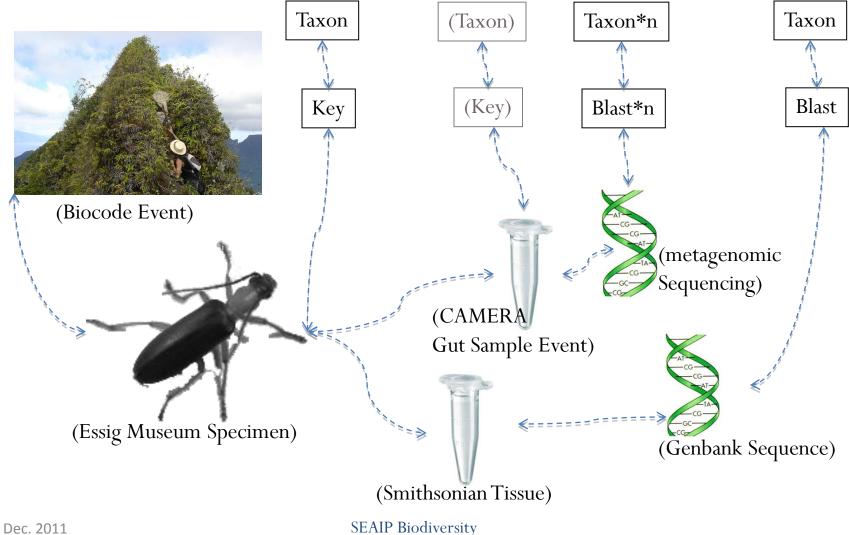


Courtesy Hank Bart, Nelson Rios, Tulane Univ.

Integration challenge



Moorea Biocode Example: Tracking biological material from field collection through analysis, across multiple systems



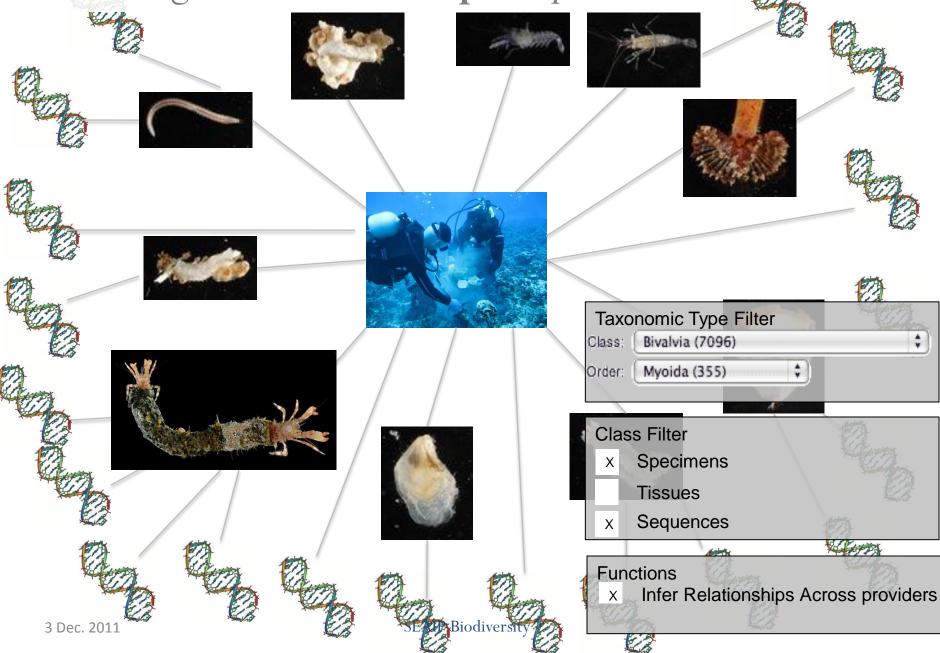
Linked queries

- List all participating institutions that house sequenced DNA samples for marine organism species Y
- For plant species X, indicate if an image and/or a tissue sample is available.
- List crab taxonomy updates by species and acceptance status by participating institution.
- For sponge species N, find updated determinations and list institutions holding the reference specimens
- List all species in publication Y that have determinations and locality updated in date range *d1-d2*.
- Find all plants of Species N collected within date range *d1-d2* and have digital images.

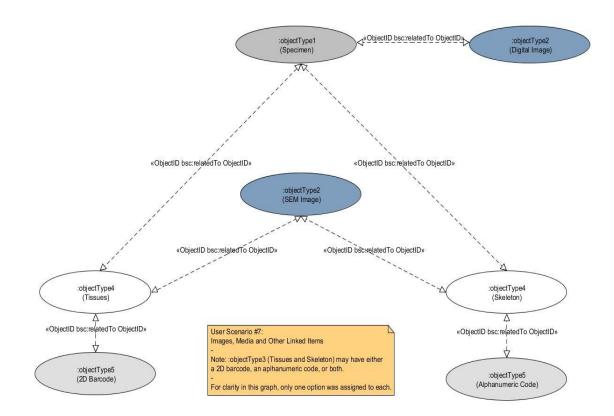
What relationships exist that haven't been explicitly expressed?

What can you discover with linked relationships that you can't do easily with distributed data (e.g., DiGIR over Darwin Core) or standard relational database queries?

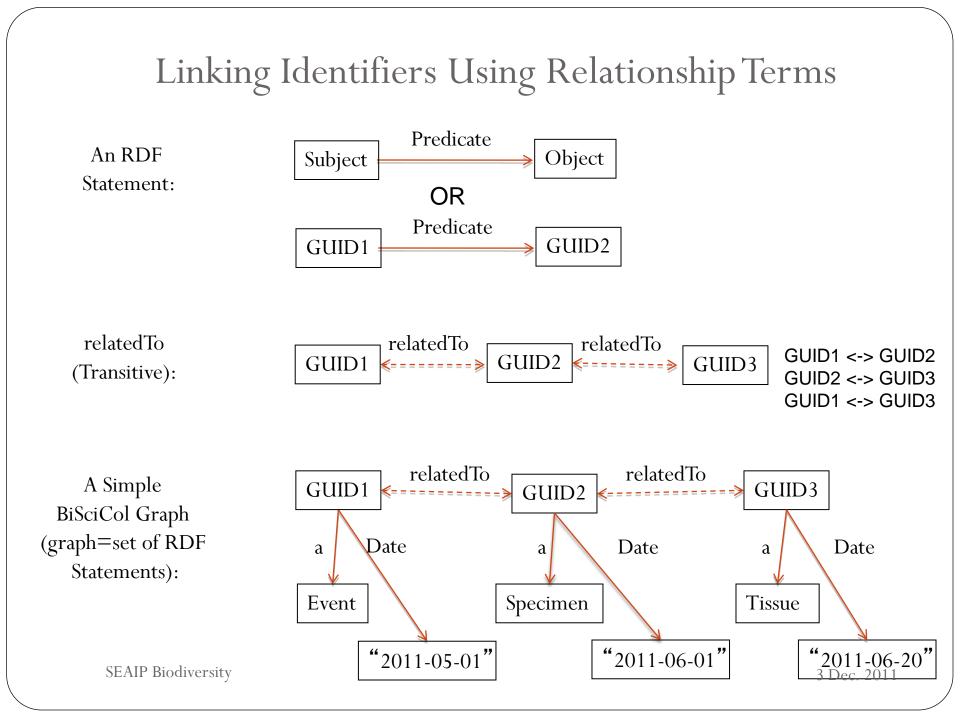
A Biological **Relationship** Graph ...



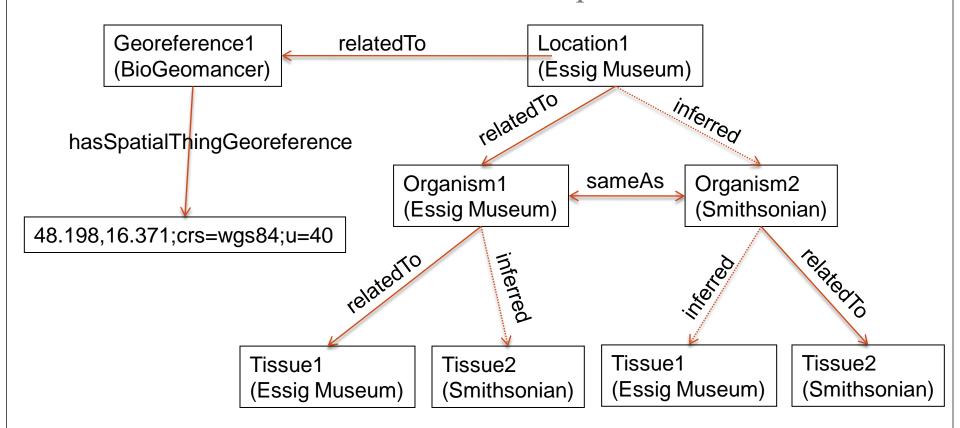
How do we Track Biological Objects and their **Relations** Across Distributed, Heterogeneous systems?



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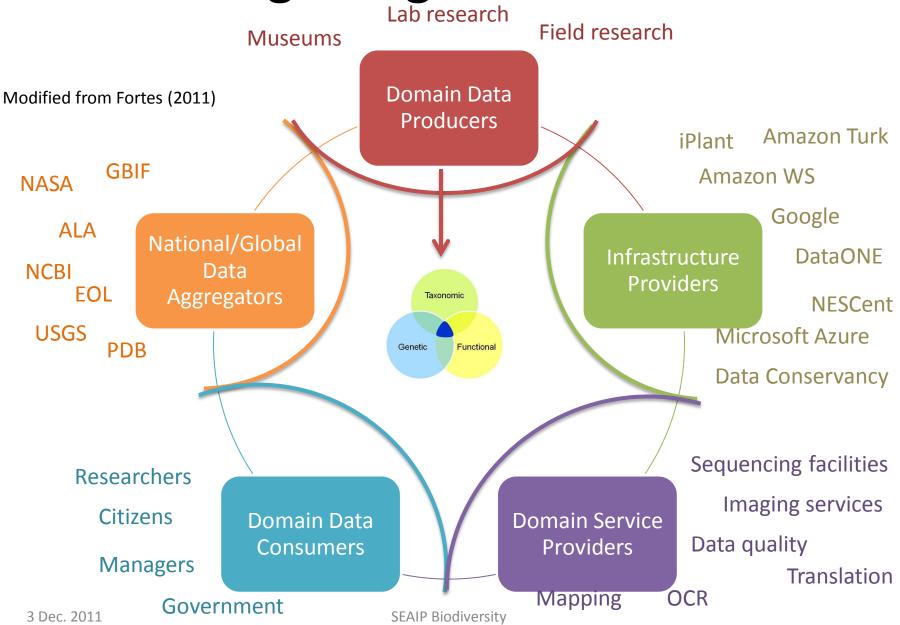


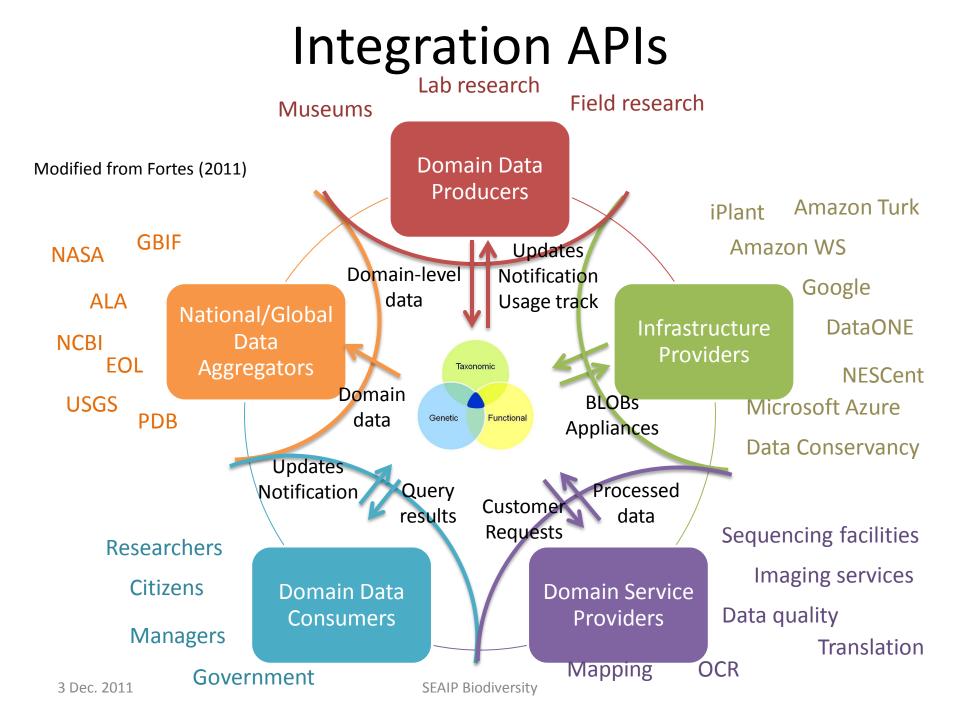
Inferred Relationship Chains



Even though Tissue #2 is not directly related to Location1, we can still infer its relationship through Organism1 and Organism2 being the same as each other. SEAIP Biodiversity 3 Dec. 2011

Integrating Dimensions



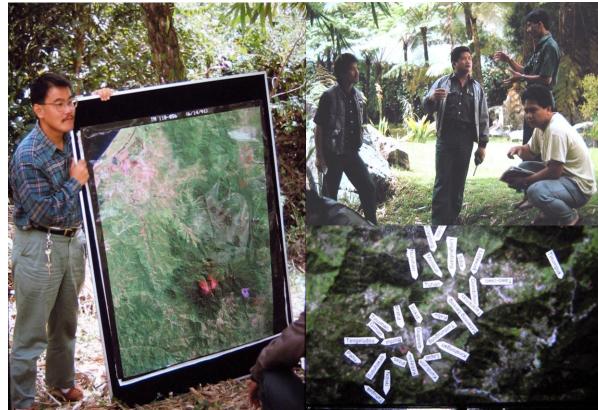


Geospatial knowledge from primary sources

Not all is legacy or retrospective but based on imagery, GPS, and sensor nets Pro

- NEON/LTER
- Planetary biodiversity surveys (PBI)
- new plot and census data

Projek Etnobotani Kinabalu (PEK)



SEAIP Biodiversity

Biodiversity valuation

Standing forests have value other than just timber

$$PV_{B} = \frac{\int_{0}^{T} P_{b}B(t) \cdot e^{-rt} dt}{1 - e^{-rT}}$$
Hartman (1976) modi

Hartman (1976) modified by Caparros (2003)

- PV_B is the present value for biodiversity values;
- B the biodiversity function; and
- P_b is the biodiversity shadow price

P_b and B are not easy to value in current practice

Making collections relevant to economics

- Biological collections are the ultimate documentation of what, where, and why for vouchering biodiversity research
- Communicate value of collections based data through education and outreach
- Economic modelers can only use information readily available. Biodiversity and informatics collaborators must
 - Provide raw and synthesized data
 - Provide [more] complete data sets
 - Provide a basis for trust

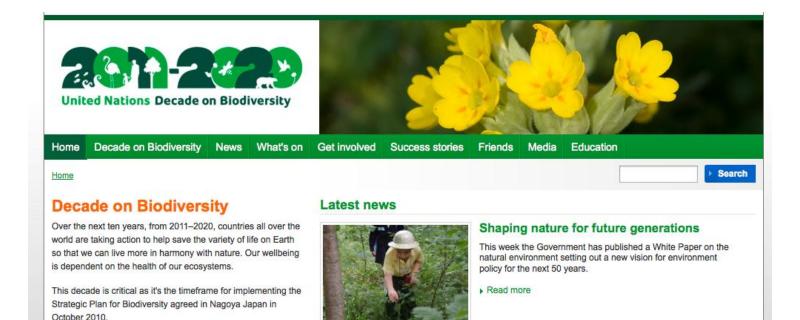
Relevant biological diversity metrics

based on vouchered biological collections

- Identifying hotspots
- Organism occurrence and distribution modeling
- Geochronological change
- Endangered/threatened/i nvasive species mapping
- Phylogenetic and genetic diversity
- Diversity of economically useful organisms

$$PV_{B} = \frac{\int_{0}^{T} P_{b}B(t) \cdot e^{-rt} dt}{1 - e^{-rT}}$$

Decade on Biodiversity: 2011-2020



 Projects mentioned were funded by NSF and the Moore Foundation. Material in slides provided by Jose Fortes and John Deck. Thanks!