# Data Workshop Honolulu March 2014

Greg Riccardi
Florida State University
iDigBio
griccardi@fsu.edu



This material is based upon work supported by the National Science Foundation under Cooperative Agreement EF-1115210. Any opinions, findings, and conclusions or recommendations expressed in this material are those of the author(s) and do not necessarily reflect the views of the National Science Foundation.



# Goals of workshop

- Draft a requirements document for aggregators that describes information and services that are crucial to the success of biodiversity informatics.
- Writing group sessions
  - Discussion of issues for plenary
- Plenary session
  - Discussing issues from the perspective of
    - Providers
    - Users



#### Draft list of issues for discussion

- Full record-level information discovery and delivery
- Metadata harvesting protocols
- GUID per record with persistence
- Attribution metadata with all data records
- Media information ala Audubon Core
- Bi-directional portal
- Feedback from data users to providers (e.g. data quality)
- Usage analytics
- Attribution to providers from analysis
- Annotation management
- Active repository technology (incremental updates)



## **Plenary Discussion Summary**

- Providers need
  - Attribution for data use
  - Help with managing taxonomy
  - Feedback from determination and data cleaning
    - The effort required to process feedback will be considerable
    - Tools are needed to help providers with feedback
  - Details about determinations and geolocations
  - Help with identifiers
  - Registries for people and localities



## **Plenary Discussion Summary**

- Users need
  - Good global information discovery services
  - Assessments of data quality, per record or dataset
  - Data cleaning services
  - Feedback for data cleaning so that improvements are made by providers
  - Tools to find related data, e.g. sequences
  - Tools to aid in integration of data from multiple sources



# **Information Integrity and Attribution**

- Provenance tracking
  - Keep track of source of information
  - Ensure information is not changed
  - Control versions
- Attribution
  - Keep track of delivery of information
  - Make attribution information available to providers
  - Provide mechanism for users to report
    - Publications
    - Derived data
    - Evaluation and corrections



#### Identifier and identifier services

- Encourage identifiers for objects
  - Require stable identifiers
    - Provider must commit to consistent use of identifiers
  - Strongly suggest that providers maintain GUIDs
  - Add GUIDs as necessary
- Identifier services
  - Return metadata document upon request
  - Discover and maintain relationships among identifiers
  - E.g. If a provider changes the identifier, the aggregator must record that the old and new identifiers are equivalent



# **Search and Discovery**

- Search by common properties
- Discover across object types
  - E.g. Find image by scientific name or geography of specimen
- Provide for download in common formats
- Provide APIs for search and download



#### **Taxonomic services**

- Assumption
  - Provider sends scientific name and possibly higher classification
- Externalize taxonomic names and classifications
  - Participate in shared services
- Allow discovery beyond name string
  - Synonyms
  - Common names
  - Higher taxa



## Dealing with extended schema

- Assumption:
  - Providers will have important information that is not Darwin Core
- Properties
  - Keep track of properties and evaluate new data sets for new properties
  - Allow both literal- and resource-valued (relationship)
- Transformations and normal forms
  - Maintain information content when changing formats
  - Transform or coalesce properties according to community standards
- extending schemas
  - traits, measurements, interactions
- property similarity with respect to discovery



## Thanks to all participants

- Writing group
  - Greg Riccardi (iDigBio)
  - Reed Beaman (iDigBio)
  - Donald Hobern (GBIF)
  - Rich Pyle (GNA, Bishop)
  - Robert Whitton (GNA, Bishop)
  - Paul Flemons (Biodiversity Info Manager, Au)
  - James Macklin (Biodiversity Info Manager, Can)
- Plenary Group
  - Joanna McCaffrey (iDigBio)
  - Deb Paul (iDigBio)
  - Neil Evenhuis (GNA, Bishop)
  - Shelley James (Bishop, Macro Algae)
  - Michael Thomas (Biodiversity Informatics Manager, U. Hawaii)
  - Chris Neefus (Macro Algae)
  - Matt Goodale (data management/IT supervisor, NTBG)
  - Tom Schils (Biodiversity Informatics Manager, Botany, phycology, UOG)
  - Aubrey Moore (Biodiversity Informatics Manager, Entomology, UOG)
  - Ryan Caesar (IT Manager/programmer, Entomology, U. Hawaii)
- Others?

