

Data Discovery Index Ecosystem

Enabling the Big Data Commons through indexing of data and their interactions



bioCADDIE Overview

- 1. Help users find accessible data
- 2. Assist data producers on how to publish data for maximal discoverability
- 3. Build a prototype/platform to dock related products

PubMed of Data = DataMed



SCIENTIFIC DAT The FAIR Guiding Principles for scientific data

Mark D. Wilkinson, Michel Dumontier, IJsbrand Jan Aalbersberg, Gabrielle Appleton, Myles Axton, Arie Baak, Niklas Blomberg, Jan-Willem Boiten, Luiz Bonino da Silva Santos, Philip E Bourne, Jildau Bouwman, Anthony J Brookes, Tim Clark, Mercè Crosas, Ingrid Dillo, Olivier Dumon, Scott Edmunds, Chris T Evelo, Richard Finkers, Alejandra Gonzalez-Beltran, Alasdair J G Gray, Paul Groth, Carole Goble, Jeffrey S. Grethe, Jaap Heringa, Peter A.C. 't Hoen, Rob Hooft, Tobias Kuhn, Ruben Kok, Joost Kok, Scott J. Lusher, Marvann E. Martone, Albert Mons, Abel L. Packer, Bengt Persson, Philippe Rocca-Serra, Marco Roos, Rene van Schaik, Susanna-Assunta Sansone, Erik Schultes, Thierry Sengstag, Ted Slater, George Strawn, Morris A. Swertz, Mark Thompson, Johan van der Lei, Erik van Mulligen, Jan Velterop, Andra Waagmeester, Peter Wittenburg, Katherine Wolstencroft, Jun Zhao, and Barend Mons

management and stewardship



- Findable
- Accessible
- යා ඉදාගෙන් මෙය

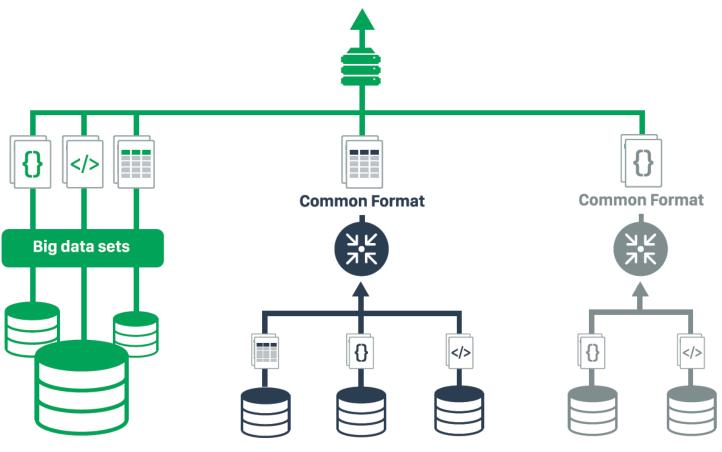
http://www.nature.com/sdata/

nature publishing group 1102





Data Discovery Index



Big data sets of particular interest to NIH and not covered by aggregators.

e.g. NIH Commons

Major aggregator services (i.e., indices or repositories that use a common metadata format)









Data Repositories...

































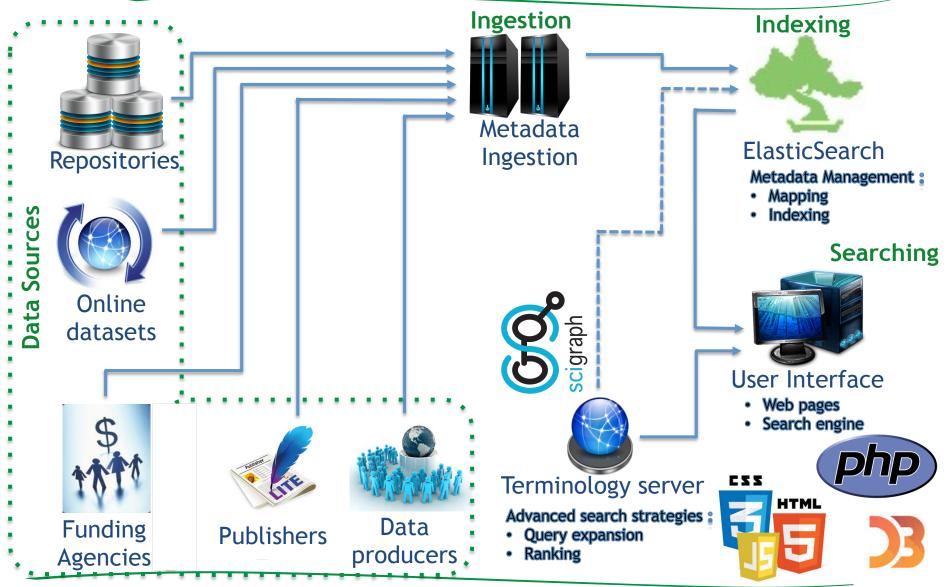






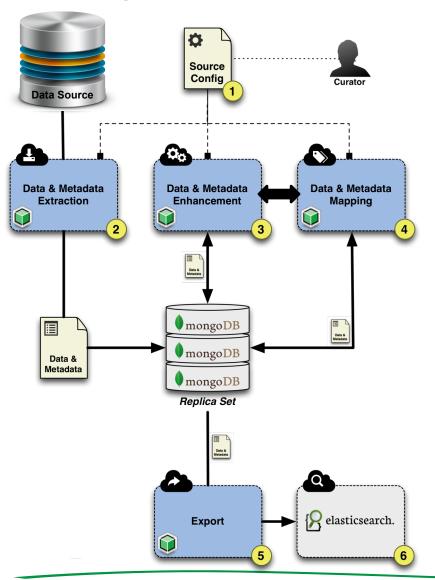


bioCADDIE Prototype Architecture





Data Indexing Pipeline



- Configuration file developed by curator
- Extraction of metadata/data from data resource or dataset via ingestion module
 - Cache information for further processing
- Process metadata/data via a set of processing modules
 - e.g. ID conversion, keyword extraction, data normalization
- Mapping of metadata/data to metadata model(s)
- Export to target endpoint(s)via export modules
- 6. Search via ElasticSearch APIs



WG3: Metadata Specifications

Metadata specification v1, future-proofed for progressive extensions, to support intended capability of the DDI prototype

PHASE 1 OUTPUT:

- NIH BD2K bioCADDIE Data Discovery Index WG3 Metadata Specification v1. 2015. Zenodo. 10.5281/zenodo.28019
 - The WG3-MetadataSpecifications-v1.zip contains a document, two Appendixes, JSON schema and examples.



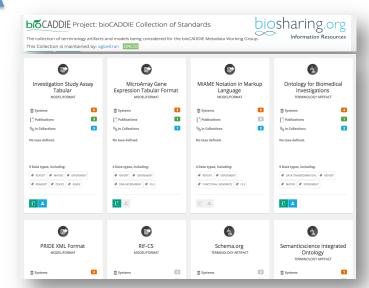


Created using 2 complementary approaches

Competency question ological process y (apoptosis) at scale z with an Search for organism x in 5 estimate of the reliability of the annotations Search for new drug x to predict and track x (cardiotoxicity) Search for data type x ('omics correlates) of to drug x Search for data types a, b, and c (EHR data, self-report, sensor) to determine natural history of patients given drugs similar to drug x Track responses to treatment to ensure detection of blological process x Find patient data "like these" with similar treatments, responses to treatment, genetics Search for studies a-z with patient data with measured by BMI) and interventions a-z. Then filter on demographic characteristics.

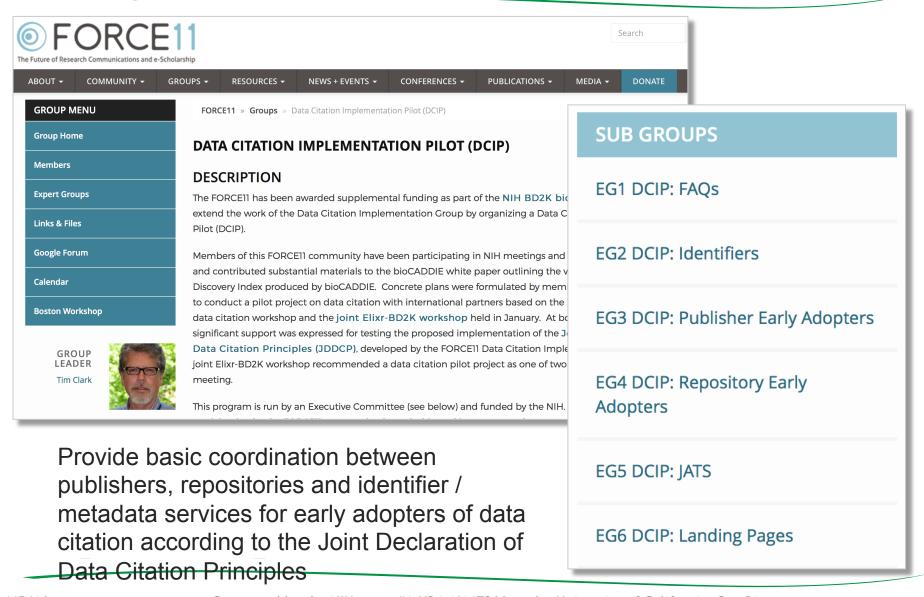
top-down: analyzing use cases

bottom-up: mapping existing standards/schemas



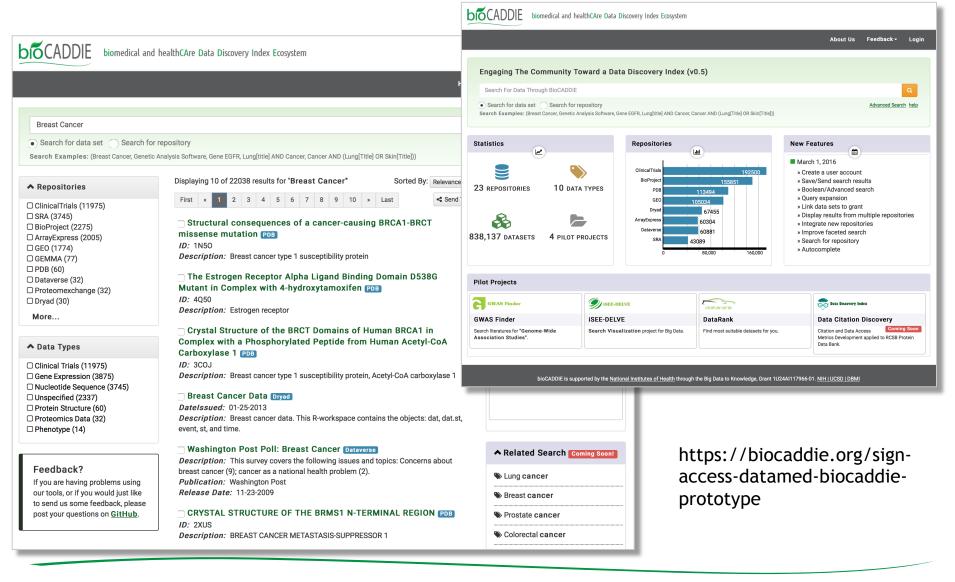


Data Citation Implementation Pilot





bioCADDIE Prototype





bioCADDIE Acknowledgements

- 93 working group members
- 12 steering committee members
- 8 pilot application reviewers
- staff and trainees
- collaborators

