Taming an Ocean of Data at AOOS

End to end data lifecycle management





Background - Axiom

- Cyberinfrastructure technology development:
 - environmental, biological and geoscientific data
- AK Headquarters, OR and RI satellite offices
- Support federal, university and NGO groups
- Mission driven







Background - Axiom

- Shared cyberinfrastructure approach
- Community developed software, standards and protocols
- Scalable compute and storage infrastructure (HPC)
 - o 5 petabytes storage; 3,000 processing cores







Background - Alaska Ocean Observing System

- Officially established in 2005
- Regional member of the Integrated
 Ocean Observing System (IOOS)
- Network of critical ocean and coastal observations, data and information products
- "Eye on Alaska's coasts and oceans"

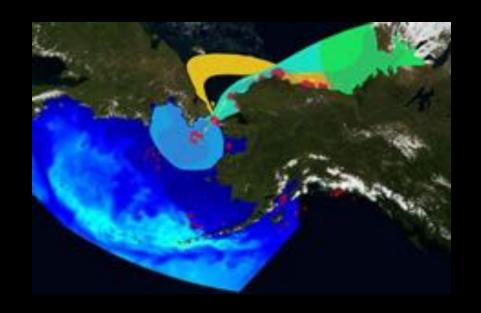






Goals - Alaska Ocean Observing System

- Increase access to existing coastal and ocean data
- Package information and data in useful ways to meet the needs of stakeholders
- Increase observing and forecasting capacity in all regions of the state, with a priority on the Arctic and Gulf of Alaska

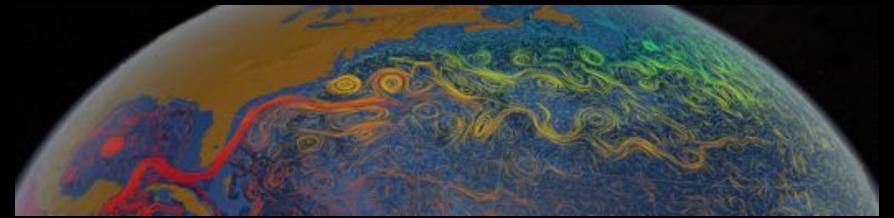






Challenges - Alaska Ocean Observing System

- Lots of existing data and research efforts in the region
 - Varying data/metadata quality
 - Limited sharing beyond specific research effort
 - Often not ready for public consumption
 - Complex data formats
 - Lack of context/metadata
 - Hard to find (not discoverable)
 - Isolated (not interoperable/comparable/synthesizable)





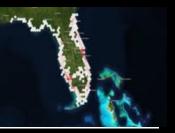


Data Types

Biodiversity

count, richness, diversity indices





Platforms

moorings, shore stations







Products

skill assessment, shoreline change, etc.

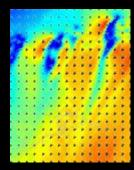


Grids

models, satellite, radar

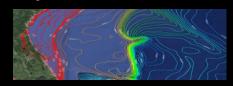




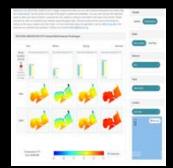


GIS

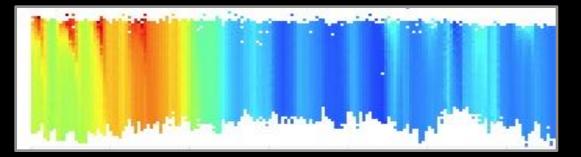
Habitat types, bathymetry, fishing zones, etc.

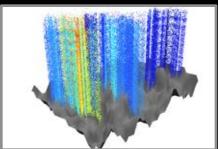






Gliders









Data Pipeline





Geospatial Database



Tabular Data



Web Service (ERDDAP, WMS, etc.)



Research Workspace



Other







Storage





GeoServer



Documentation



Metadata







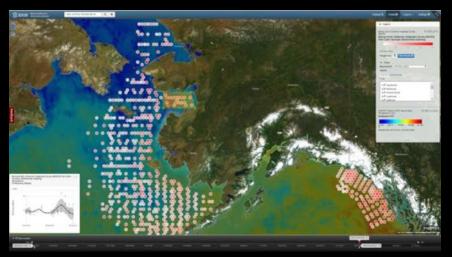
Data Cleaning/Upgrades

- Data
 - Structural/syntax problems
 - Quality Control/Quality Assurance (QA/QC)
 - Clean up or flag invalid/suspect data
 - Monitor data stream sources for outages
- Metadata
 - Make sure observable properties are clearly defined
 - Make sure units are clearly defined
 - Make sure spatial and temporal axes are defined
 - o Use community standards when possible
 - CF conventions/standard names
 - ACDD metadata attribute conventions
 - Dataset should be self describing!!
 - Even to people outside of the domain

Data Portal - Public Data Exploration and Access

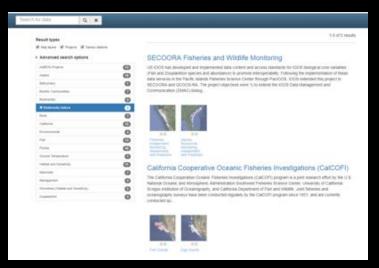
Map

Integrate & visualize data from many sources



Catalog

Search, metadata, & data download

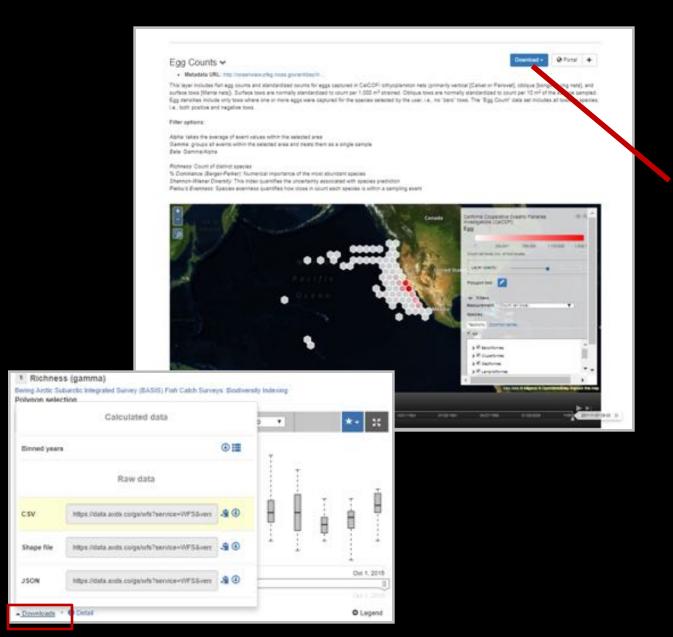


Data Views

Rapidly assimilate & compare different data streams



Downloads Using Interoperability Services



ncWMS
Shapefile
CSV
THREDDS
netCDF
OPenDAP
ERDDAP

ERDDAP



Dataset Title: Granite Crk SS 0000

ERDDAP > tabledap > Make A Graph ..

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That's great for structured, predictable data. What about research programs, synthesis projects, citizen science, etc.?

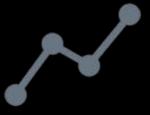
RESEARCHAWORKSPACE

~web-based platform for collaboratively managing science projects through the entire data lifecycle~

Share Analyze Preserve

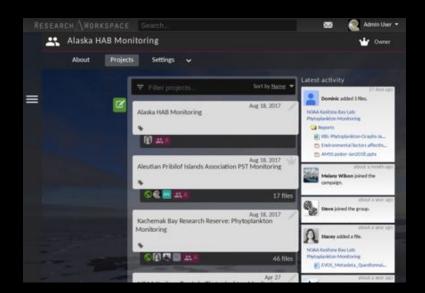


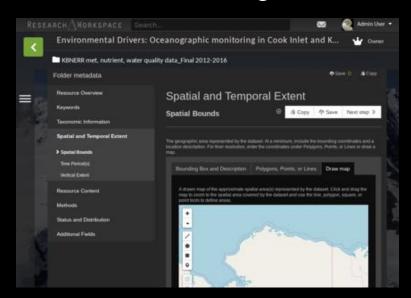




RESEARCH MORKSPACE

- Organize into projects, research campaigns, and organizations
- Manage sharing through advanced security permissions
- Coordinate data exchange across networks, groups, programs
- ISO 19115/19110 standards metadata editor
- Execute server side Python and R numeric workflows (Jupyter) on uploaded data AND any data in Axiom CI stack
- Archive pathway to DataONE & Datacite DOI minting





DATA COLLECTION & QUALITY CONTROL

Scientists or Ingestion

REUSE & TRANSFORMATION

Jupyter Notebook & data analyses

Supporting the entire data lifecycle

ACCESS & DISCOVERY

Data portals & search catalogs

STORAGE

Research Workspace

DESCRIPTION

Metadata Editor

ARCHIVE & PRESERVATION

Repository submission pathway

Research Workspace - Metadata

- We help build large collections of diverse datasets.
- Unorganized, undocumented data collections benefit no one.
- Metadata tells the story (who, what, when, where, why, and how) of the data.
- Metadata makes data discoverable.



Research Workspace - Metadata

- The best metadata is written by the people closest to the data.
- Many researchers aren't familiar with authoring it.
- Metadata can be difficult to write well (existing standards are complex and confusing).
- Researchers are also very busy people.



Resource Overview

> Basic Overview

Contacts

Category and Form

Keywords

Taxonomic Information

Spatial and Temporal Extent

Resource Content

Methods

Status and Distribution

Additional Fields

Resource Overview

Basic Overview



This section provides an overview about the dataset and any associated file(s).

Resource Title @

A descriptive title for the data that includes the subject mater, where data was collected, and when it was collected.

Assessing abundance of beluga whales in Bristol Bay using genetic mark-recapture methods, 2002-2011

Abstract ©

A surressry of the key aspects of the dataset that excludes when, where, why, and have it was collected, as well as a brief description of its usuables and file formats.

This project estimated the abundance of beluga whales within the Bristol Bay stock using genetic markrecapture methods and combined genetic data with aerial survey data to develop an unbiased correction factor for use in future aerial surveys. The project was started in 2004 by the Alaska Beluga Whale Committee, which funded sample collection from 2004 until 2012 and genotyping from 2004 until 2011, and continued through funding from the North Pacific Research Board (NPRB 1516) from 2015 through 2017. The data for this project were generated using genetic markers from skin biopsies of beluga whales Bristol Bay from 2002 to 2011 using mark-recapture methods.

Data from this project consists of 2 .csv data files archived here (NPRB_1516_Bristol_Bay_beluga_whale_abundance_data_sample_list.csv and NPRB_1516_Bristol_Bay_beluga_whale_abundance_data_matching_file.csv).

Purpose ©

The intention of the dataset and why it was collected or developed, as well as a statement about the dataset's relevance to any target project or effort.

The Bristol Bay beluga whole stock is genetically distinct from other stocks and tagging studies show it is restricted to Bristol Bay year-round. Quantifying the abundance of belugas in the Bristol Bay stock is important for their management and is critical information for upcoming stock status reviews. This is the first estimate of abundance of belugas in Bristol Bay with appropriate confidence limits.

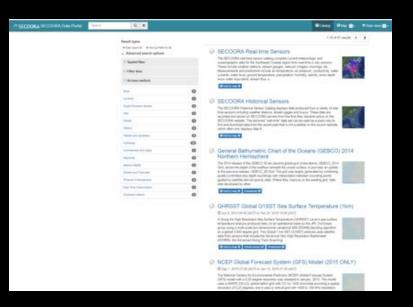
Research Workspace - Publish & Archive

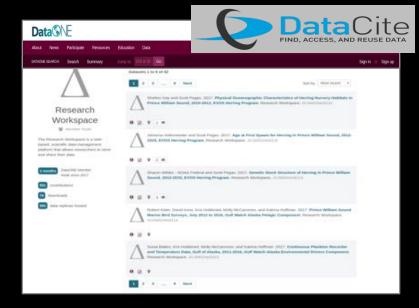


Data Portal











Mission:

"Enable new science and knowledge creation through universal access to data about life on earth and the environment that sustains it."

How:

Cyberinfrastructure + Community

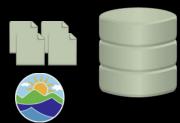


DataONE - Networked Repositories

Member Nodes







Coordinating Nodes





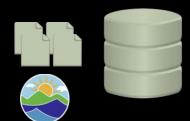
DataONE - Member Nodes

Member Nodes

- Defined policies
- Persistent IDs
- Immutable content
- Standardized metadata
- Resource maps (bagit, etc)
- Implement DataONE API





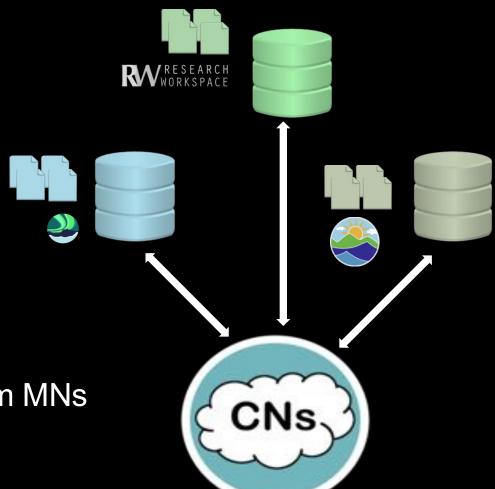






DataONE - Coordinating Nodes

Member Nodes (Preservation Repo + DataONE API)



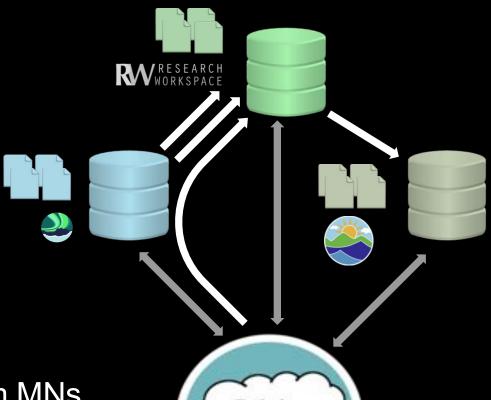
Coordinating Nodes

Sync metadata from MNs



DataONE - Coordinating Nodes

Member Nodes (Preservation Repo + DataONE API)

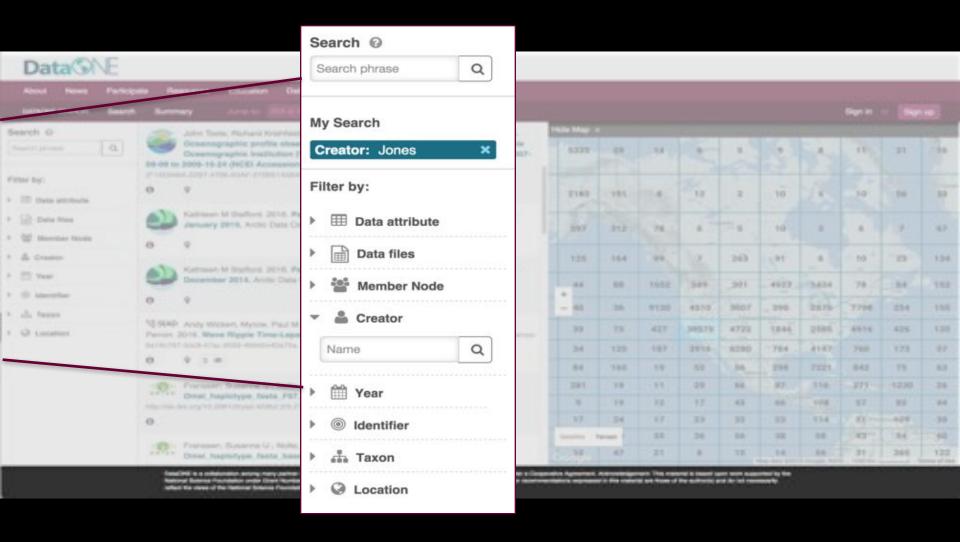


Coordinating Nodes

- Sync metadata from MNs
- Control MN to MN replication

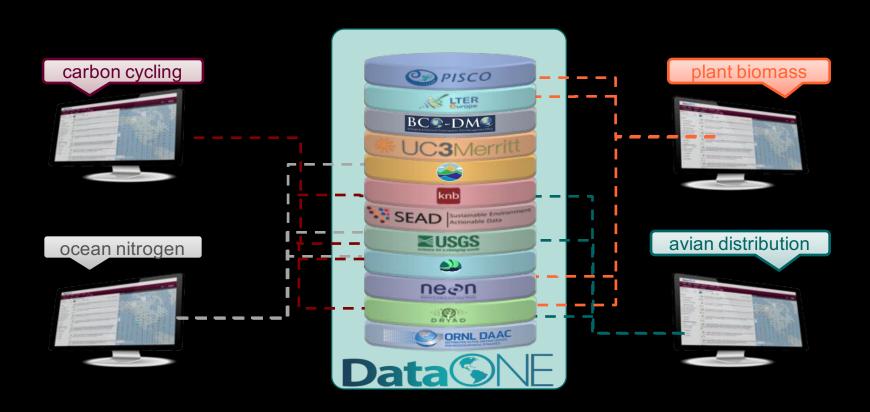


DataONE - Federated Search





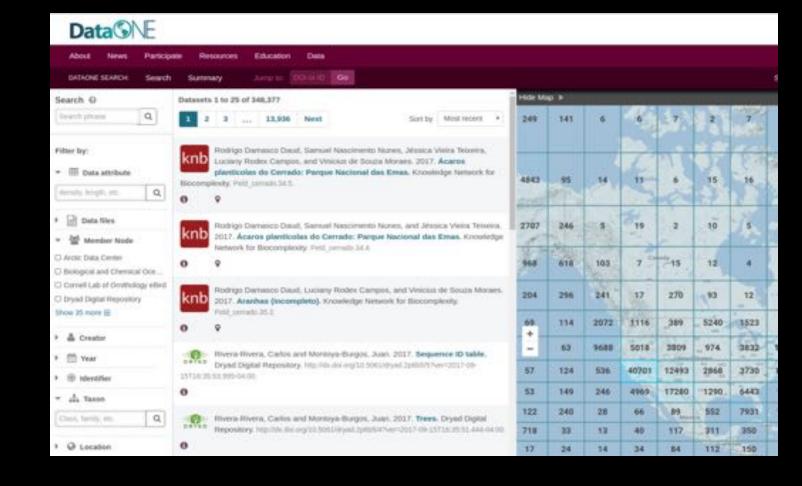
DataONE - Discoverability







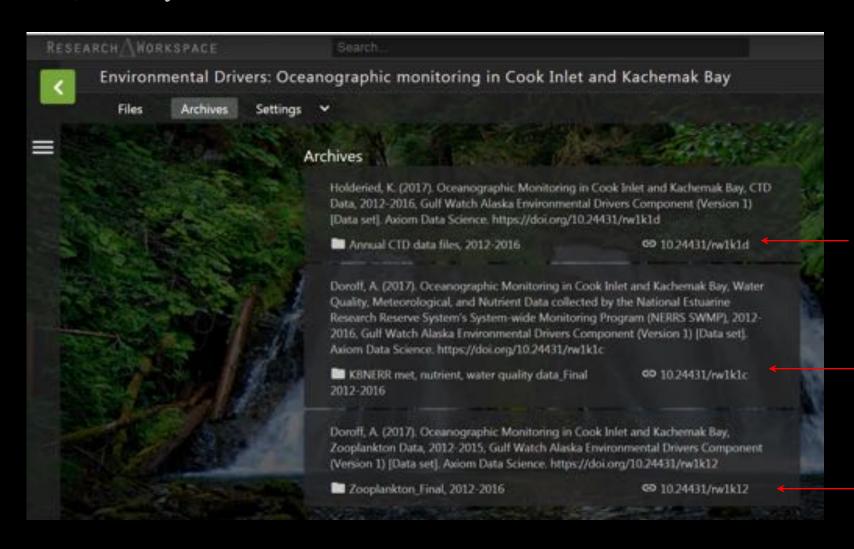




RESEARCH \(\frac{1}{2}\) WORKSPACE



Locate, identify and cite research data



Research Workspace - Analysis/Synthesis

- Challenges
 - Data Availability
 - Compute Resources
 - Barriers to Entry
- Solution: Jupyter Notebooks
 - Local availability of large datasets
 - TB+ model/satellite data
 - Real time sensor system
 - No need to download data to analyze
 - Powerful compute resources
 - High bandwidth/throughput
 - Powerful CPUs / large RAM
 - Hardware optimized for numerical computation
 - No software management burden!





RESEARCH \(\frac{1}{2}\) WORKSPACE



- Create and share documents that contain code, equations, and visualizations
- Reproducible numerical simulations and statistical modeling
- Access uploaded data stored in the Workspace or data portal



the number of distinct species found in a sample % Dominance (Berger-Parker) the ratio between the number of individuals belonging to the most abundant species and the toal number of individuals in the sampe $Dominance = max(p_i)$ Shannon-Wiener Diversity index quantifies the uncertainty associated with species prediction Pielou's Evenness species evenness quantifies how close in count each species is within a sampling event In [17]: # create dominance and shannon-weaver diversity indices p_i_stats = aggregated_df.groupby(['location_id', dateField])['p_i'].agg diversity = aggregated df.groupby(['location_id', dateField]).agg({ species tsn' : np.count nonzero, 'lat station' : np.mean, # why are decimals truncated? 'lon station' : np.mean, # why are decimals truncated? }).rename(columns={'species tsn': 'richness'}) diversity = diversity.merge(p i stats, left index=True, right index=True # add Pielou's Evenness Index diversity['evenness'] = diversity['sw diversity']/np.log(diversity['rich diversity = diversity.reset index(level=[dateField, 'location id'])

diversity



Richness

the number of distinct species found in a sample

$$S = \sum (p_i > 0)$$

% Dominance (Berger-Parker)

the ratio between the number of individuals belonging to the most abundant species and the toal number of individuals in the sampe

$$Dominance = max(p_i)$$

Shannon-Wiener Diversity

index quantifies the uncertainty associated with species prediction

$$H' = -\sum_{i=1}^{S} p_i ln(p_i)$$

Pielou's Evenness

species evenness quantifies how close in count each species is within a sampling event.

$$J' = \frac{H'}{ln(S)}$$

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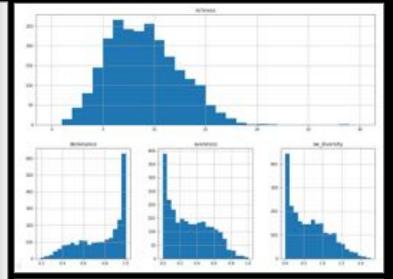
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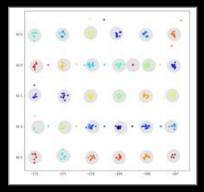
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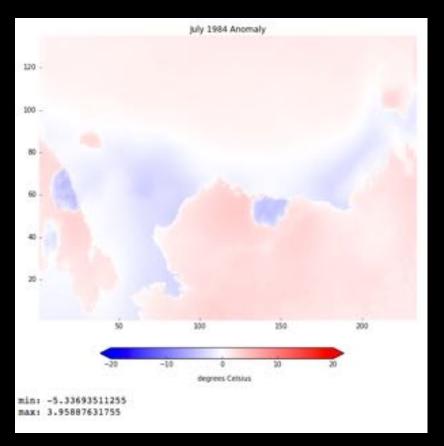
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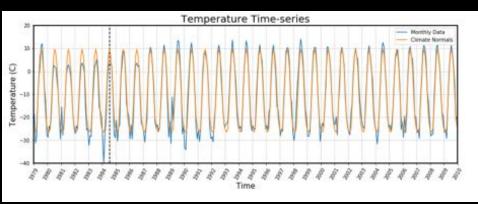


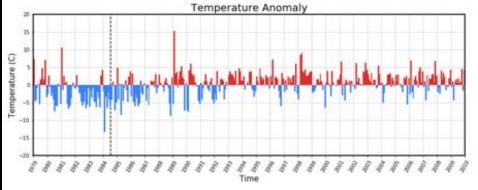




Research Workspace - Time-series Anomalies







- Calculate climate normals on a 31-year long, multi-terabyte dataset
- Then plot temperature anomalies over a region



