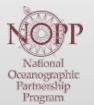


MBON All-Hands Meeting Silver Spring, MD May 03, 2016









Monterey Bay, Florida Keys, and Flower Garden Banks National Marine Sanctuaries

Principal Investigators: Frank Muller-Karger (USF) Francisco Chávez (MBARI)















The Sanctuaries MBON pilot

<u>Outline</u>

- Goals
- Data collection and management
- Outreach
- International dimensions
- Project successes and challenges

The Sanctuaries MBON pilot: goals

- Develop a <u>practical and sustainable</u> pilot MBON with the *National Marine Sanctuaries Program*
- Integrate, augment, synthesize information from ongoing programs
- Develop technologies for biodiversity assessments:
 - environmental DNA (eDNA)
 - Remote sensing time-series: Seascapes
- Integrate biological data into national/international databases
- Plan to transition to an operational system
- Help develop international connections for a global network

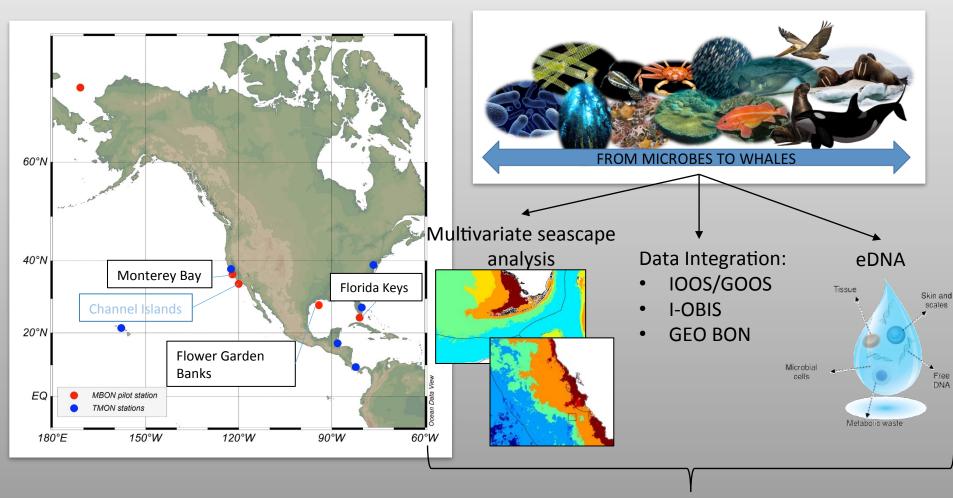
Why we do this: The importance of biodiversity

Diversity defines the function and health of ecosystems

Ultimately the right mix of species on Earth helps ensure our own survival

Understanding habitat and biological diversity and function is fundamental to define conservation strategies

The Sanctuaries MBON pilot concept



- Sanctuary Condition Reports
- Resource managers and policy makers
- Scientists and educators

Supports Web-based information system

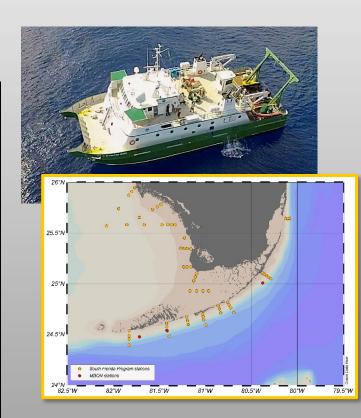


Field programs

Complement Multiple Data Collection Efforts in Each Sanctuary



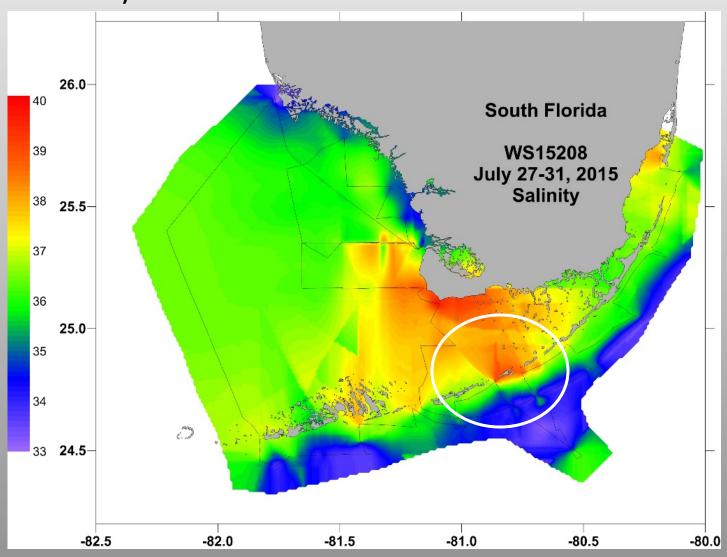




South Florida Program run by AOML partners:

- Chris Kelble–bio oc./prim prod
- Lindsey Visser
- Libby Johns phys oc. analyses

High salinity event during July 27-31, 2015, in Florida Bay and FKNMS



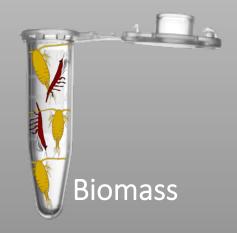


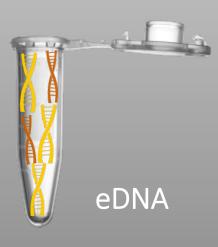
eDNA - A New Frontier

Sanctuaries MBON eDNA: vertebrates, zooplankton, and microbes

- Testing of eDNA extraction methods completed manuscript in prep.
- Primers: selection completed for some organisms
- eDNA validation in progress (i.e. zooplankton and whales)







MBON Genomics Team

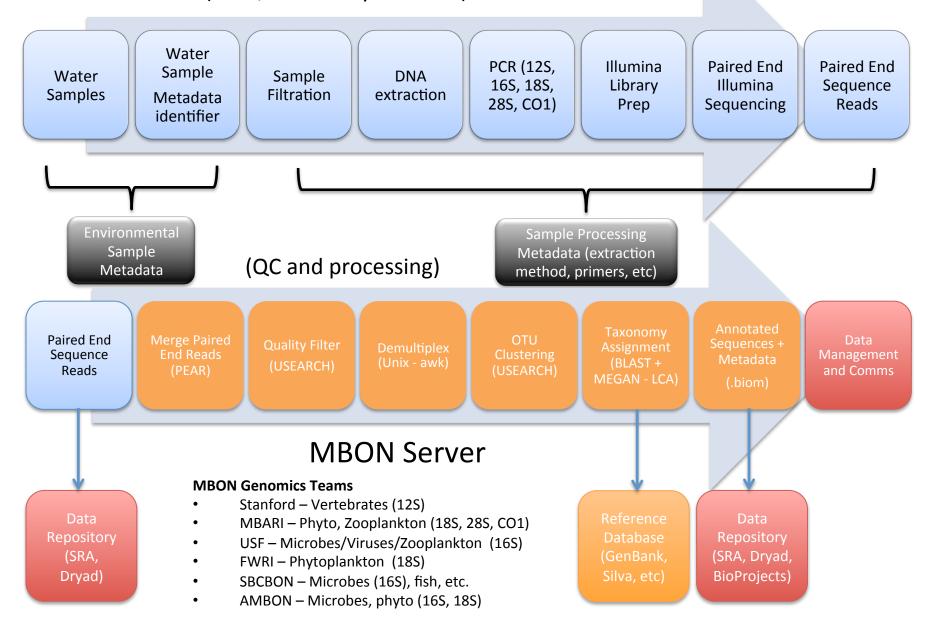
- Stanford Vertebrates (12S)
- MBARI Phyto, Zooplankton (18S, 28S, CO1)
- USF Microbes/Viruses/Zooplankton (16S)
- FWRI Phytoplankton (18S)
- SBCBON Microbes (16S), fish, etc.
- AMBON Microbes, phyto (16S, 18S)

Collaborations

- X-MBON: Santa Barbara, Alaska MBON
- NOAA Omics Research: (K. Goodwin, J. Hendee)
 - Atlantic Oceanographic & Meteorological Lab
 - Southwest Fisheries Science Center (SWFSC)

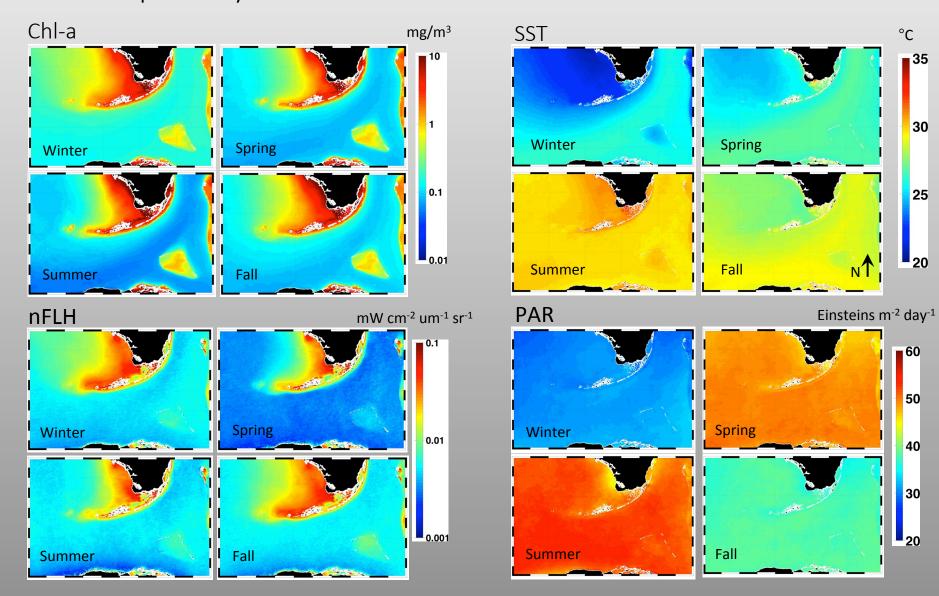
MBON Sequencing Work and Data Flow

(Field, laboratory and NGS)



Remote Sensing of Seascapes

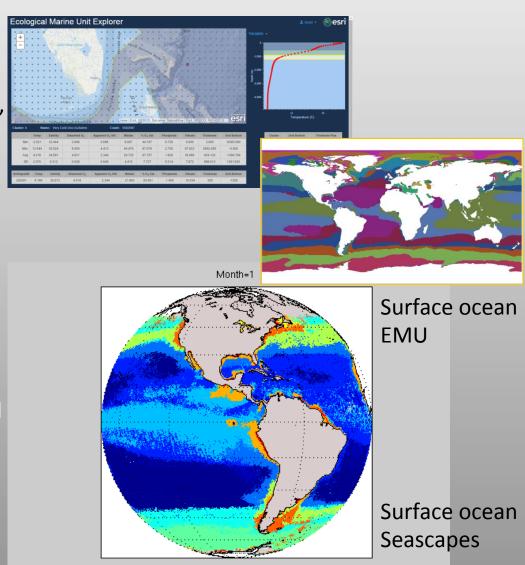
MODIS-Aqua 1-km seasonal climatologies (2002 -2015) for seascape analyses - FKNMS



Regional to Global Seascapes

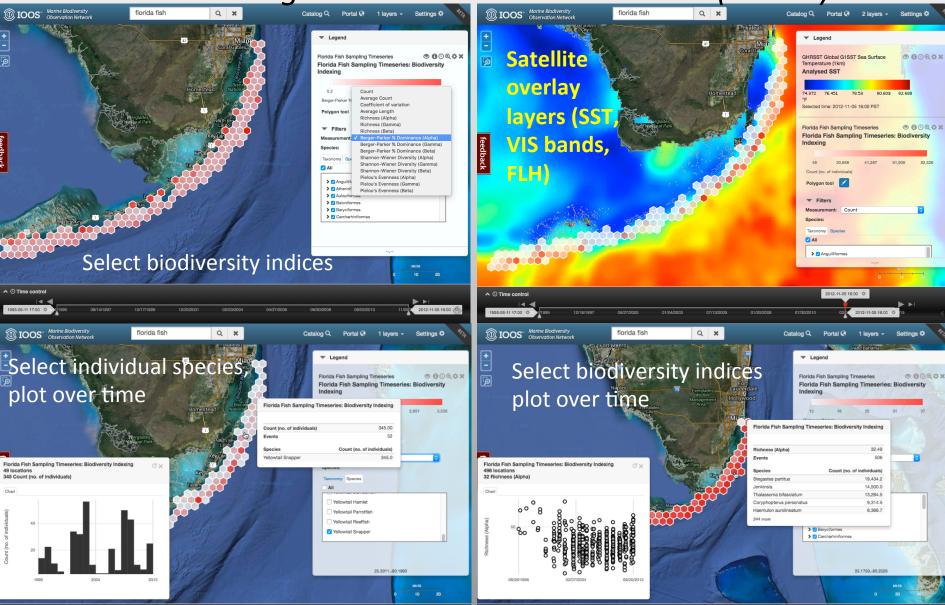
In progress/Next steps:

- In situ validation
- Automate data flow between NASA, USF, WHOI, IOOS DMAC / Axiom
- Improving algorithms: Harmful Algal Blooms (HAB), carbonate system
- Expand high-resolution coverage:
 - Flower Garden Banks, AK-BON, Santa Barbara, other NMS
- Evaluate user metrics
- GEOBON and broader links:
- ➤ Work with Roger Sayre (USGS) to link remote sensing Seascapes and Ecological Marine Units/EMU (USGS/esri)
 - Pole to Pole Biodiversity and Biogeogeography
- ➤ Other partners/internationally



Integrated Ocean Observing System (IOOS)

Data Management and Communications (DMAC)





MBON Communications & Outreach

CJ Reynolds, Jennifer Brown

• Newsletter (distributed via all_mbon@marine.usf.edu)

SANCTUARY MBON

Updates from the Florida Keys and Monterey Bay Projects

DMAC UPDATE

The DMAC team is revising the data management plan under the leadership of Dave Anderson at CeNCOOS. The Florida and Monterey Bay teams are working on enrollment of historical data sets for ingestion in close collaboration with Philip Goldstein with OBIS-USA. Axiom contin-ues to make progress on data ingestion of Florida and California biological data sets and indested a large number of Florida Keys National Marine Sanctuary (FKNMS) ecological data GIS layers and several time series data sets.

Eighteen years (1995-2012) of the Reef Visual Census (RVC) data have been transformed from their original forms into the standards-based forms used by OBIS-USA, such as Darwin Core and the Marine Bio-Geography (MGB) common terms definitions. Guidance for this effort was derived from the "MBG 2.1 Enrollment Journal Florida Marine Sanctuaries Fish 2004 Draft 20140521" produced by Philip Goldstein. The transformed data and metadata were written into NetCDF files and in-stalled in an ERDDAP server making them available through the network via human or URL-based interfaces.

Axiom and a team of Monterey Bay and Florida scientists are working on the development of a generalized biodiversity indices tool. Axiom is working with the RVC data set to prepare it for real-time biodiversity in-dices calculations. Software engineers have started to incorporate biodi-versity indices calculations into backend server side analytical code. The prototype biodiversity indices tool is expected to be available for use with the AXIOM tool in early March. Axiom (along with IOOS and OBIS staff) met with Barbara Block and team at Hopkins Marine station and discussed best paths forward to accessing ATN data relevant to

The RVC data is available through a GCOOS <u>ERDDAP server</u>. The ERDDAP interface allows users to filter the data, produce maps, and deliver data and metadata in many different formats. Click <u>here:</u> The "Make a Graph" page allows users to plot selected data and export the plots. You can enter constraints by hand or generate the plot directly from here.

Another impressive dataset, the Historical NOAA CTD data for the Florida Kevs (30 raw CTD data sets collected from the R/V Walton Smith from 2006-2015) have been incorporated into the MBON data manage ment system. Data collected include temperature, salinity, beam atten ation, beam transmission, dissolved oxygen, surface and *in situ* irradiance (PAR), Underway systems collect near-surface temperature, salinitv. meteorological and navigation data. Some cruises include current meter data. In the coming month, we will have products from these three-dimensional time series datasets mapped. Additional water quality data will be indested from http://data.gcoos.org/nutrients/_and.other sources to visualize and analyze with the Axiom workspace

A DMAC X-MBON team has been formed with participants from Alaska Florida, Monterey Bay, Santa Barbara and the Smithsonian MarineGEO

Welcome

focuses on Florida pro-



NOAA SEFSC

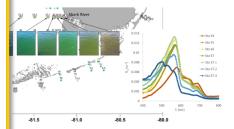
MBON project with John Lamkin (NOAA SEFSC). Shulzitski's exp

habitats throughout the FKNMS. Ana

lecting specific phytoplankton func-PFT) in the Florida Keys. Specifically, ific absorption spectra of phyto-

July 27-31, 2015, indicate high blue-green algae (very likely *Tricho*shown as absorption peaks at 545 Time series of (a*phy) and comple

practical tool for studying changes in biodiversity of lower trophic levels in this region and aid in the improvement of satellite ocean color algorithms for from samples collected at the three detecting shifts in phytoplankton community com-



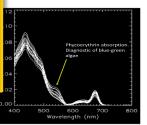


Figure 8. Chlorophyll-specific phytoplankton absorption (a*phy) spectra of surface and bottom samples from the three MBON sites during July 27-31, 2015. Absorption peaks at 545 nm are indicative of the presence of the pigment phycoerythrin, and therefore of cyanobacteria in the

Marine Biodiversity Observation Network Update Feb 2016 7

Roughly quarterly distribution

RI and USF have been working on methods. We are testing whether we ze the choice of filter type and DNA ethod to capture eDNA from multiple simultaneously (i.e. bacteria, phyto-pplankton, vertebrates).

amples collected on an MBARI CANfall 2015 to assess the spatial distriebrate eDNA across

time-series at station rev Bay (6 cruises.10 led per cruise from 0vielded very low/absent for 12S mtDNA DNA); we have providof this time-series to has successfully ampli-

ter volume is related to DNA concentration in ers, our team collected 10-L and 100-L mples at station M1 us-I flow filtration (TFF)

inkton and zooplankton

across these volumes and to convenmay repeat this experiment in anothfore finalizing our results.

eam is also comparing two library s for NGS to determine whether this standardized across the three ic teams. Results should be obtained

sting MBARI with setting up a NA studies. USF and FWRI are test-

gene markers used by the MBON genetic teams. The USF/FKNMS genomics team has collected

monthly samples in the Florida Keys near some of the major coral reefs. Samples are collected with help of the USF remote sensing group and scienists at the FKNMS during cruises on the R/V Walton Smith and on small boats.

Currently, USF/FKNMS collect three samples from surface and bottom at each of the three key MBON stations - resulting in 18 samples per cruise. We work are complete and data analysis already have 198 samples from the three stations annuscript are underway. Stanford is and an additional 286 samples from the surrounding ocean!



Figure 9. Fall 2015 CANON cruise sampling

The USF group has also collected samples for zooplankton morphological identifications, eDNA genomics, and for sequencing tissue of whole zoo-plankton communities. The goal is to ground truth the eDNA method for zooplankton and move for-ward to eventually use only eDNA to assess zooplankton diversity and seasonal oscillations in the Florida keys.

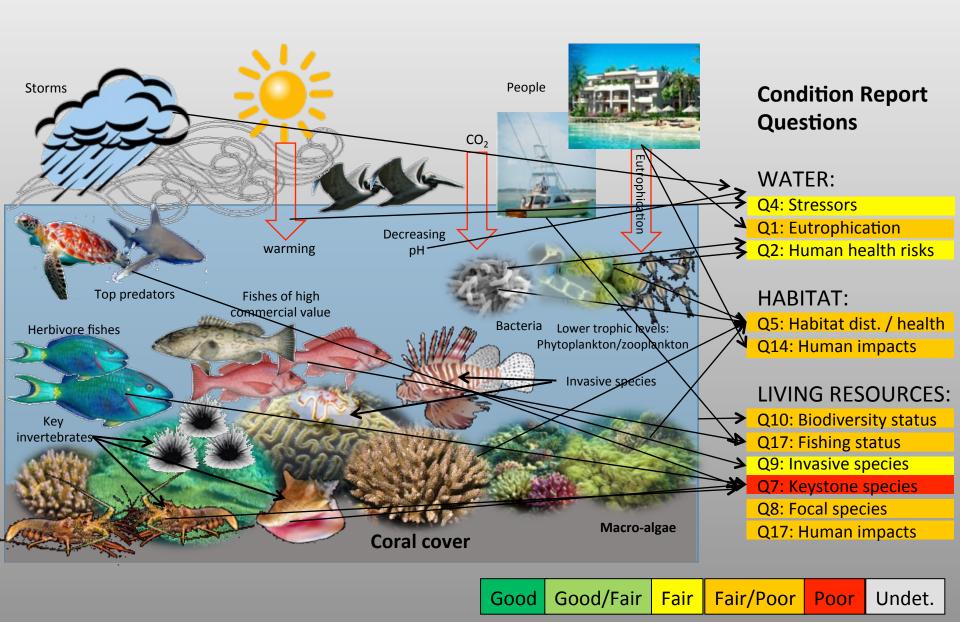
Morphological zooplankton identification is almost done and we plan to have the sequencing ready for the Genomics Team (USF, FWC, Stanford, and pipeline to process and analyze NGS MBARI) meeting at MBARI on Feb. 10-12. This will enable comparison of zooplankton data between his system. This pipeline will provide a bridged data analysis framework for different ing, the genomics teams will decide on a final the Florida Kevs and Monterey Bay. At the meetmethod for sample collection and data analysis



Next steps: Putting it all together

 Build the decision-support tool to answer Sanctuary Condition Report questions

Dynamic support for Condition Reports

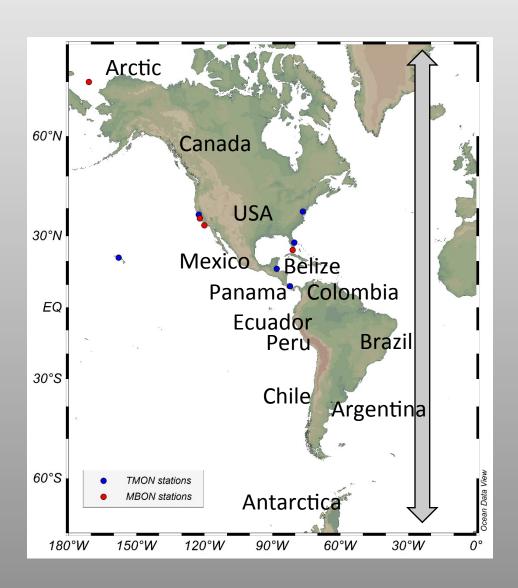


MBON

Marine Biodiversity Observation Network



MBON beyond the US: GEO, GOOS, CBD



GOALS:

- Pole-to-Pole MBON pilot
 - the Americas
- BON in a Box

Outreach and planning

- MBON presentations at GEO Plenary, Mexico (2015)
- MBON presentation at the Convention of Biological Diversity (Montreal, Apr 24, 2016)
- GEO BON Open Science Meeting (Leipzig, Jul 4-6,2016)
- GEO MBON All-Hands Workshop (Leipzig, Jul 6-7,2016)
- Pole-to-Pole in the Americas Workshop (Puerto Morelos, Mexico, Sep 26-30, 2016)
- MBON presentation at the GEO-XIII Plenary (St Petersburg, Russia, Nov 9-10, 2016)

Successes and challenges within the Sanctuaries MBON

Successes:

- Conceptual framework for collaborative MBON:
- eDNA: collection and extraction methods tested
- Satellite-based, dynamic seascape products: automated pipeline
- Biodiversity field monitoring program implemented in both Sanctuaries
- MBON data used to update 2015 Monterey Bay NMS Condition Report
- Making links: NOAA ocean acidification program, NOAA Omics, State and Federal fisheries & environmental monitoring, NSF LTER (Everglades), IOOS and other observation programs
 - IOOS GCOOS has funded a Doctoral fellowship at USF to work MBON-LTER-SFP

Challenges:

- Development and maintenance of data system / visualization tool:
 - The amount of work is staggering many details
 - How to transition to a sustainable model?
- eDNA validation at different trophic levels
 - For ex. vertebrates in different regions
- Curation and permanent archive of biological datasets from various sources:
 - Identifying and understanding datasets is an ongoing effort
 - How do these data transition into a permanent archive? How do we maintain access needed for operational & research utility?
 - How do we engage monitoring programs to enroll data?
- Communications flow on news and outreach
- Integrating the MBON observations into the FKNMS Condition Report-2017(?)
- Coordination of myriad moving parts with partners and X-MBON projects
- Building critical international partners and linkages for Pole-to-Pole
- Operational MBON
 - path to sustainability not clear

Marine Biodiversity Observation Network