Oceans and Ecosystems Research

Molecular & Environmental Microbiology

to explore biodiversity, understand ecological function, and protect health

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Microbial Dynamics Underpin Ecosystem Function

(they control...pretty much everything)

Good:

- Nutrient Cycles (including photosynthesis)
- Climate-controlling gases
- Ozone-depleting gases
- Waste Treatment
- Degrade toxic compounds
- Life-saving medicines



Bad:

- Impair Water Quality (including oxygen depletion)
- Diseases & Infections of fish, protected species, corals, humans



Main Science Questions

- What are the sources of degraded water quality in coastal ecosystems*?
- How can we improve microbial water quality assessments (faster, easier, more accurate) and transfer that knowledge to stakeholders?
- How can we best harness the power of 'omic technologies to characterize a main driver of the Earth System – microbial diversity & function.

For what purpose?

- Better protect health, ecosystems, and economies from sewage pollution, pathogens, and harmful algae.
- Improve understanding of ecosystem diversity and function to improve assessment and stewardship of ecosystem services



*geographically specified system of organisms (including humans), the environment, and the processes that control its dynamics. – NOAA Five Year Research and Development Plan 2013-2017



AOML Program Review

Microbial Water Quality Assessments

Applications include recreational water & seafood safety, regulatory compliance, diseases of protected species & commercial fish

Recreational Water Focus –

Beaches Are Bucks:





- 85% of US tourism dollars are from coastal recreation with the creation of 1.6 million jobs in 2000. CA beaches: >100 million visitors annual. FL beaches: \$19.1 billion from out-of-state tourists in 2003.
- **Repercussions of degraded water quality:**
 - Microbial contamination makes people sick.
 - Contamination (or fear of) adversely affects consumer confidence.
 - Local governments forced to spend millions \$\$ annually on monitoring.
 - Mitigation is mandated costs can be 100's of millions \$\$\$\$.



• Links to NOAA Strategic Plan

• Holistic Understanding of the Earth System through Research

- Multiple links, including specific Implementation Plan Inclusion (transformative research & innovation/Research to Operations)
- Engagement Enterprise
- Resilient Coastal Communities and Economies
 - Improved coastal water quality supporting human health and coastal ecosystem services
- Healthy Oceans
 - Improved understanding of ecosystems to inform resource management decisions; Recovered and healthy marine and coastal species
- Other NOAA Links: Microbe and Ecosystem Health Working Group, One Health Working Group, Ecoforecasting Pathogen Team, OAR Ecosystem Portfolio, 2013 National Ocean Policy Implementation Plan



Partnerships and Stakeholders

Public Health Agencies, Fisheries, Tourism Industry, Environmental/Conservation Organizations, Concerned Citizens, Beach Managers, Water Quality Managers, Academic Institutions, Environmental and Public Health Scientists



AOML Molecular and Environmental Microbiology



- Development, Validation, and Transfer of Molecular Tools
- Application of Microbial Water Quality Assessments for:
 - coral reef and coastal ecosystem health research
 - investigation of ocean and human/animal health interactions
- 'Omic tools to characterize Earth System drivers:
 - microbial diversity and function controlling, e.g., global biogeochemistry, food webs, biological response to change
- Research Mentorship and International Engagement



Development, Validation and Transfer of Molecular Tools



Why are molecular tools needed for microbial water quality assessments?



Traditional Methods are too Slow



Consent decree lead to new EPA guidelines (2012) and spurred research on molecular and automated methods



Development, Validation and Transfer of Molecular Tools

Microbial Source Tracking (MST)

MST identifies contamination sources (e.g., humans, birds, dogs) to decrease risks to health and to preserve coastal economies. MST helps managers devise mitigation strategies and assess whether investments are working.

AOML MST assays for dog and gull contamination:

- validated -- large state-funded project
- transferred -- academia, industry, state & local governments. Protocols provided in the 2013 "*California Microbial Source Identification Manual*"





AOML gull assay matched the number of gulls measured by automated cameras

AOML inter-laboratory validation efforts contribute to state and national guidelines



Assay	N	Sensitivity		Specificity		
	Human n ^a	DNQ+	DNQ-	Non-human nª	DNQ+	DNQ-
BacH	12	100%	75%	26	77%	85%
BacHum-UCD	72	97%	97%	156	37%	67%
BsteriF1	48	100%	96%	96% 104	44%	61%
BtH	12	100%	92%	26	54%	96%
gyrB	12	92%	50%	26	58%	96%
HF183						
endpoint	84	75%	NA	182	96%	NA
HF183 SYBR	48	100%	92%	104	78%	89%
HF183				UNACTORIES - 1111		
Taqman	60	100%	95%	130	46%	92%
HumM2	72	93%	67%	156	75%	94%
Mnif	60	78%	60%	130	68%	76%

AOML testing of multiple human-associated fecal markers for the SIPP study. HF183 Taqman emerged as one of the recommended assays for human MST. Examples:

- EPA National Exposure Research Laboratory, Multi-Laboratory Validation Study for enterococci and Bacteroidales qPCR.
- EPA Office of Water, Inter-Laboratory Validation Study of EPA fecal indicator qPCR assays in support of revised 2012 recreational water quality criteria.
- State of California, Source Identification Protocol Project (SIPP) for 41 MST methods tested by 27 laboratories.



Development, Validation and Transfer of Molecular Tools

Since 2000, AOML has developed and evaluated a variety of platforms for molecular assays. Goals: faster, cheaper, more accurate, high throughput, multiple target detection













colorimetric microplate suspension array

electrochemical

QPCR

SCODA

whole cell dielectrophoresis

get cells

collect

sample

Recent work includes sample preparation methods for in situ, automated biosensing platforms, such as the Environmental Sample Processor (ESP)



Application of Microbial Water Quality Assessments



Land-Based Pollution



Epidemiology



Protected Species



BMP Evaluation



Natural Source Exclusion



Coral Pathogens

Example Studies:

- FL Keys National Marine Sanctuary Little Venice Canal
- F.A.C.E. Program
- Tijuana Watershed
- Hobie Beach Baseline
- High Intensity Diurnal
- Epidemiology FL (B.E.A.C.H.E.S)
- Epidemiology CA (CA Water Study)
- Bather Shedding
- Coastal MSSA & MRSA
- Sand Pathogens
- Pathogen/Indicator Environmental Persistence
- DOH/DOE Event Response
- Zoonotic Pathogens
- Seaweed Wrack & Sand
- Poche Beach Watershed
- Cabrillo Beach TMDL
- Coral Pathogen, S. marcescens

AOML MST efforts guide and evaluate remediation efforts





AOML MST assay shows widespread contamination from gulls. Fecal indicator bacteria concentrations drop after completion of a bird-exclusion structure.

AOML testing of residential canals in the Florida Keys indicate that switching from septic to sewer improves water quality.



Application of Microbial Water Quality Assessments: for investigation of ocean and human/animal health interactions

Pathogen detection is used to characterize mechanisms of contamination and risk assessment for humans & animals



Pathogen research leads to recommendations for human-animal interactions during rehabilitation.





Pathogen load on dolphin skin tracks to distance of habitat from shore, suggesting impacts from land-based sources of pollution.

Staphylococcus aureus (MSSA and MRSA) in seawater correlated with the number of human bathers, suggesting that humans are a source of beach contamination.

SA (In CFU/L) = -4.6 + 0.58 *SA (In CFU/100 dryg) + 0.22*watertemp(C)+0.22ENT(In CFU/L) + 0.018xswimmers

Multivariate linear regression describes *S. aureus* concentrations.

The 'omic revolution: transforming medical, food, energy, and environmental sciences

	Metagenomics & microbes: circumvents the obstacle of laboratory cultivation
from this	
The pace of technology: 100 DNA sequences to 100 Million DNA sequences per run in 10 years	AGGGGGGGCGGCAGCTTACACATGCA AGTCGAGCGGCAGCTTACACATGCA AGTCGAGCGGTAGCACAAGGAGA GCTTGCTCTCTGGGGTGACGAGCGG CGGACGGGGTGAGTAATGTCTGGG AAACTGCCTGATGGAGGGGGGATA ACTACTGGAAACGGTAGCTAATA CGCATAACGTCGCAAGACCAAAG AGGGGGACCTTCGGGCCTCTTG

Transformational Research:

"...Revolutionizing entire disciplines, creating entirely new fields, or disrupting accepted theories and perspectives."

-National Science Foundation

'Omic tools to characterize Earth System drivers:

Metagenomics to uncover microbial diversity & function, set ecosystem baselines, and support ocean exploration



Samples collected pre-DWH showed metabolic pathways consistent with the ecosystem response to the spill. Methanotrophic pathway coverage was similar in shallow (200m) and deep (650m) samples; BTEX degradation and methanogenesis coverage was enhanced in deep samples.





Bacterial Order	%	Archaeal Order	%
Campylobacterales	27	Methanosarcinales	55
Alteromonadales	12	Methanomicrobiales	18
Desulfobacterales	8	Methanobacteriales	12

Gut samples from the rare methane ice worm -- dominated by organic, sulfur, and methane degraders.



Methane hydrate covered in methane ice worms



Bioinformatic extraction of methane ice worm genome from a mix of bacterial and archaeal sequences.



Research Mentorship

Science Education and Research Mentorship at NOAA AOML: Providing Opportunities for the Next Generation

Examples:

- NOAA Hollings Scholars
- •NOAA-Smith College Program Interns
- Environmental Protection Agency Interns
- University Research Assistants
- •NSF Research Experience for Undergraduates
- High School Research Interns















Next Steps: International Engagement & Coral Genomic Observatories

EU-US Commission Joint Consultative Group 2013 Meeting Opportunities for Collaboration in Marine Microbial Ecology

• Short-term tangible identified: NOAA participation in Ocean Sampling Day (OSD), a worldwide synchronized sampling of seawater for metagenomic analysis using internationally-accepted protocols for sampling, shipping, analyzing, storing and data archiving. *Summer Solstice June 21-22, 2014.*



• New funding:

Establish Coral Reef Genomic Observatories to characterize microbial community structure, biodiversity, and land-based pollution source impacts in Gulfstream Reef tracts by next-generation sequencing & community genomic characterization.



Questions?







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