

The AOML Environmental Microbiology Program:

Collaborative Inter-Disciplinary Research for Molecular Assessment
of Microbial Water Quality In Coastal Ecosystems

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NOAA AOML: The Environmental Microbiology Program

What do we do?

Molecular Microbiology to make coastal water quality assessment faster, easier, cheaper, and more accurate

Why do we do it?

To better protect the public health, ecosystems, and economy from:
Sewage Pollution, Human Pathogens, Harmful Algae

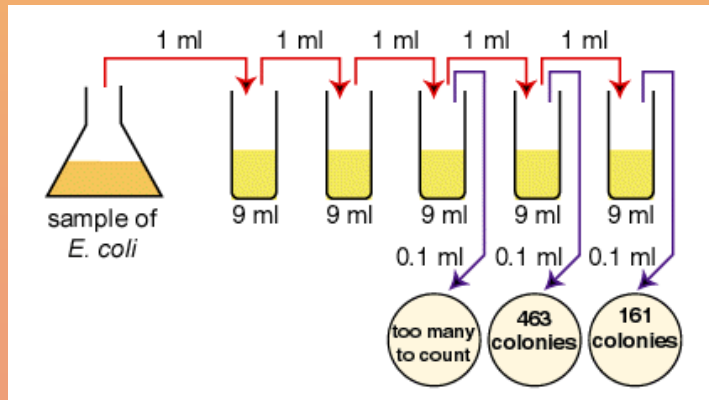
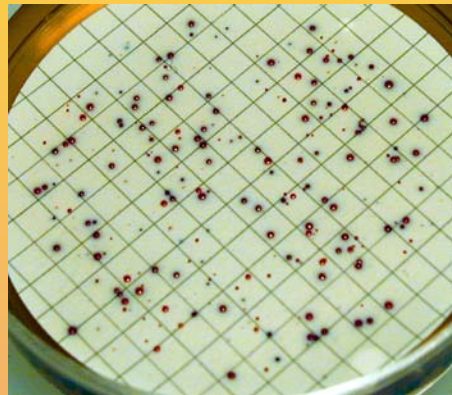
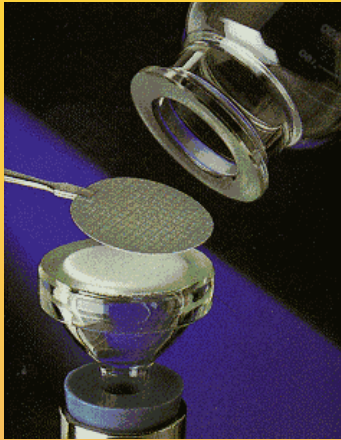
The Specifics?

Develop and/or adapt ***molecular assays and sensors*** to detect microbial contaminants in coastal waters by measuring DNA or RNA signatures.

Who are our Customers and Partners?

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

Traditional assays need improvement



Fecal Indicator Assays:

- too slow (>18 hr)
- it is unclear if the correct thing is being monitored (axioms of fecal indicator theory violated)
- no pathogen detection
- no ability to source track (needed to guide remediation)

The Potential of Biotechnology

- ✓ species-specific ID
- ✓ rapid
- ✓ sensitive
- ✓ high-throughput
- ✓ allows sample storage
- ✓ culture & microscope independent
- ✓ simultaneous information on multiple targets
- ✓ Source-tracking of fecal indicator hosts



Key Scientific Questions

- Can molecular analysis be used to quickly and reliably identify human pathogens in coastal waters?
- Can molecular microbial source tracking aid in the reliable discrimination of input sources of microbial contamination to coastal waters?
- What are the sources of nutrients, pathogens, and fecal indicators in coastal waters?
- Is there a linkage between sources of nutrients and microbial contamination in coastal waters?

NOAA AOML

Tools for Environmental Microbiology:

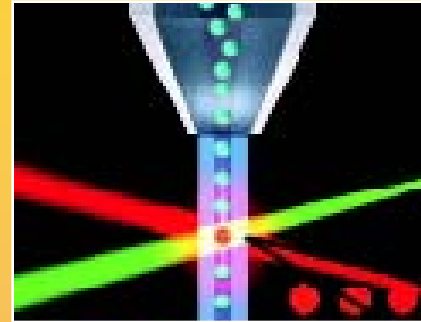
Molecular-Based



**Real-Time
quantitative
PCR**



**colorimetric
microplate
hybridization**



**Luminex
suspension
array**



**Handheld
Electrochemical
Biosensor**

Traditional Culture-Based



IDEXX EnteroLert™



**Membrane-Filtration
Plate Counts**

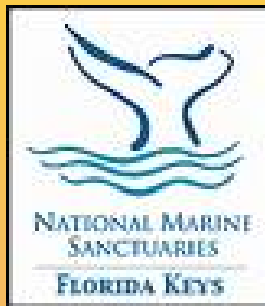


**Alternative Indicators
(Clostridium, etc.)**

Incorporation of Molecular Assays into Field Programs

Southeast Environmental Research Center
Working to Restore and Protect our Unique Environment

FLORIDA INTERNATIONAL UNIVERSITY
Miami's public research university



Atlantic Oceanographic & Meteorological Laboratory

National Oceanic & Atmospheric Administration

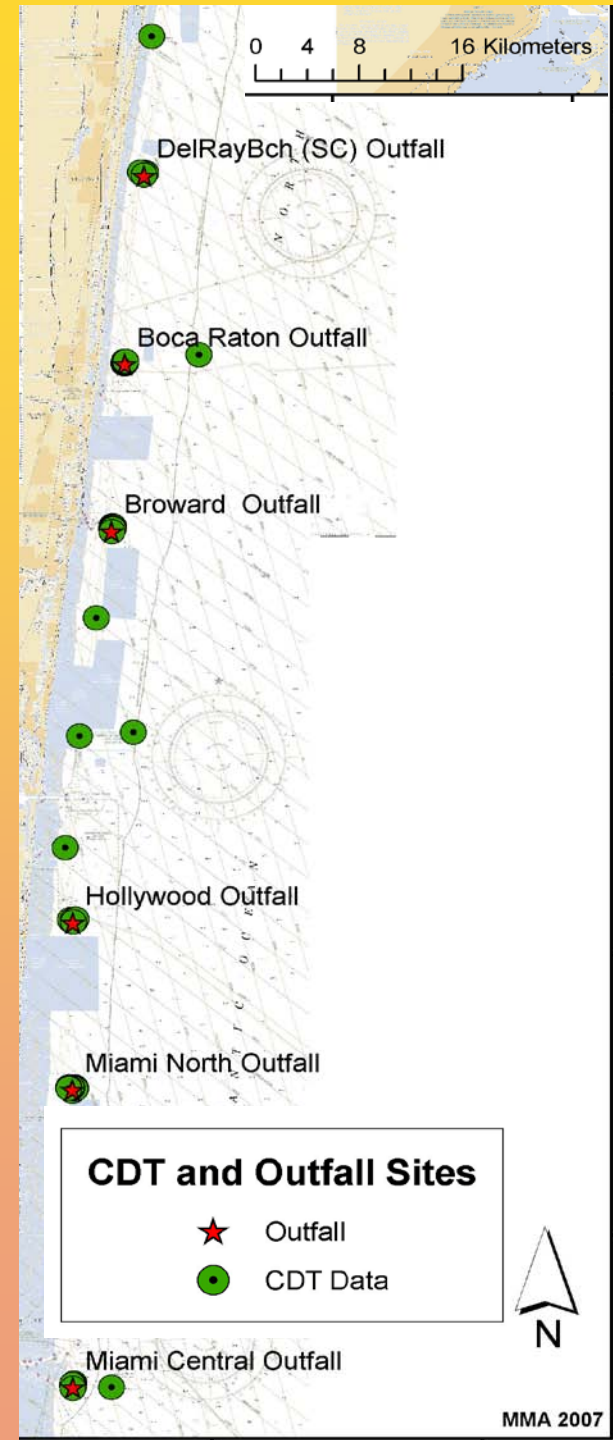


Florida Area Coastal Environment (FACE)

Water discharges to coastal waters including outfalls, inlets, & upwelling characterized for nutrient concentrations & microbiological water quality



Six treated wastewater outfalls off the Southeast Florida Coast



**Molecular Microbiology Sampling aboard
NOAA Ship Nancy Foster**

Multiple Assays Incorporated

enterococci

Escherichia coli / *Shigella* spp.

standard fecal
indicators

Bacteroides Fragilis Group

Bacteroides distasonis

alternative fecal
indicators

enterococci human marker

Bacteroides human markers

Bacteroides dog marker

source tracking
markers

E. coli O157:H7

Campylobacter jejuni

Salmonella spp.

Staphylococcus aureus

adenovirus

pathogens

intestinal & dermal
bacteria and virus

■ = Luminex

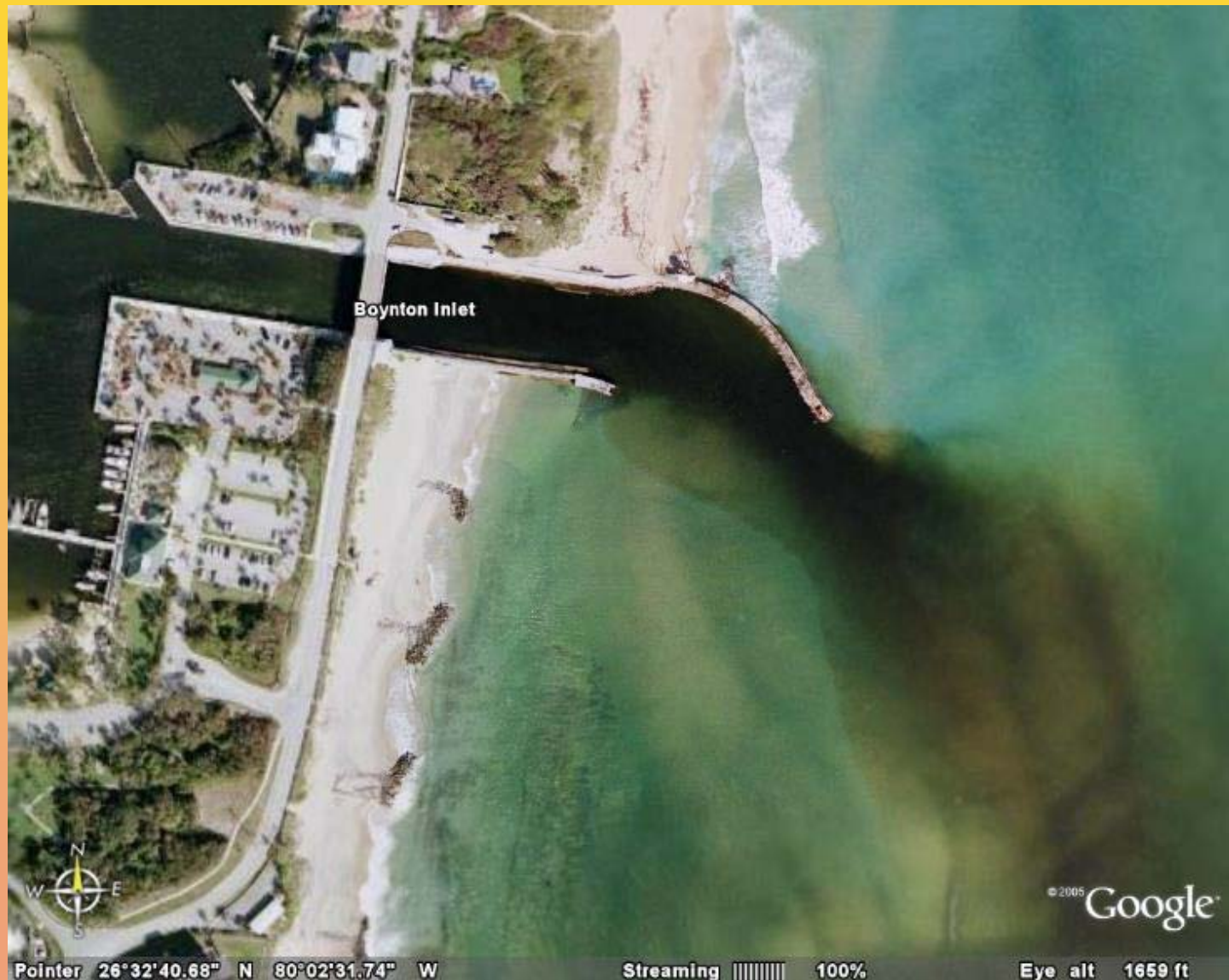
■+■ = qPCR & EC

Indicators and Pathogens in Wastewater Outfall Boils

assay	HWD	HWD	HWD btm	MC	MC btm	NMB	NMB btm
enterococci	+	+	+	-	+	-	+
human <i>Bacteroides</i> (HF8)	-	-	+	-	-	-	-
<i>S. aureus</i>	-	-	-	+	-	-	+
adenovirus	-	-	+	+	-	-	-
Norovirus	-	-	+	+	-	-	+
Phage MS2	+	-	-	+	-	+	+
Crypto/100L	55	17		236		8	
Giardia/100L	68	119		246		120	

Human enterococci, *Salmonella*, enterovirus: all negative

Tidal Discharges from Coastal Inlets



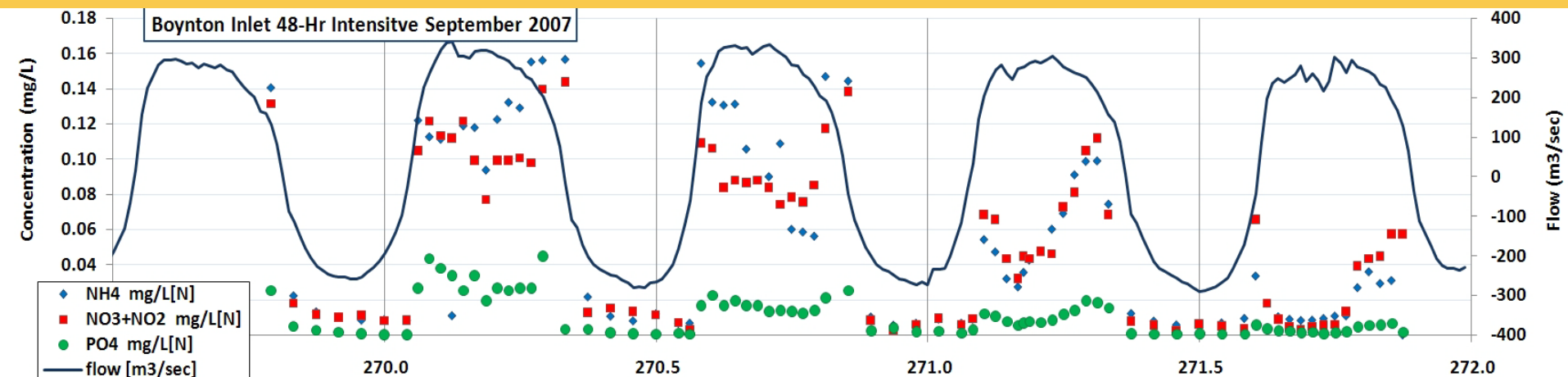
Satellite Photo Boynton Inlet

Export of microbial contamination with outgoing tide, Boynton Inlet 1st 48-hour intensive study, June 2007

assay	0400	1000	1600	2200	0400	1000	1600	2200	1400
human enterococci (esp)	+	-	-	-	+	-	-	-	+
human <i>Bacteroides</i> (HuBac)	+	-	-	-	+	-	+	+	+
<i>S. aureus</i>	+	-	-	-	+	-	+	+	+
adenovirus	+	-	-	+	-	-	+	-	+
norovirus	+	-	-	-	-				+
enterovirus	-	-	-	-	+				+
Cryptop/100L			6.3	<1	2.4				24.9
Giardia/100L			4.2	<1	1.2				19.4

inlet, outgoing tide; *Salmonella*, 0157:H7, *C. jejuni*: negative

Trends in elevated microbial contamination on outgoing tides correspond to trends in elevated nutrient levels



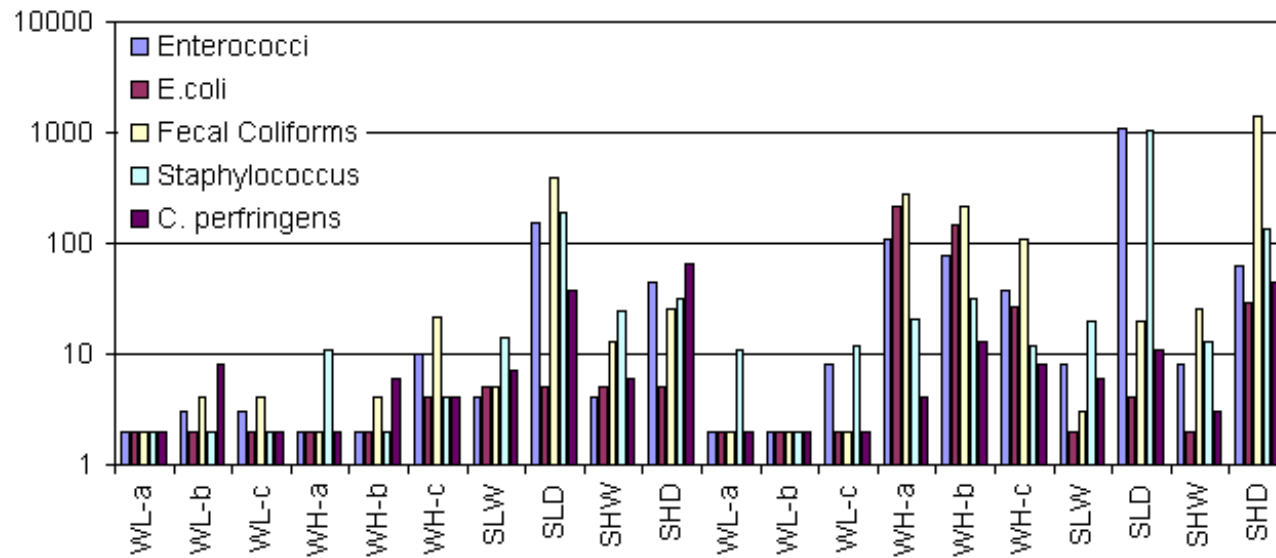
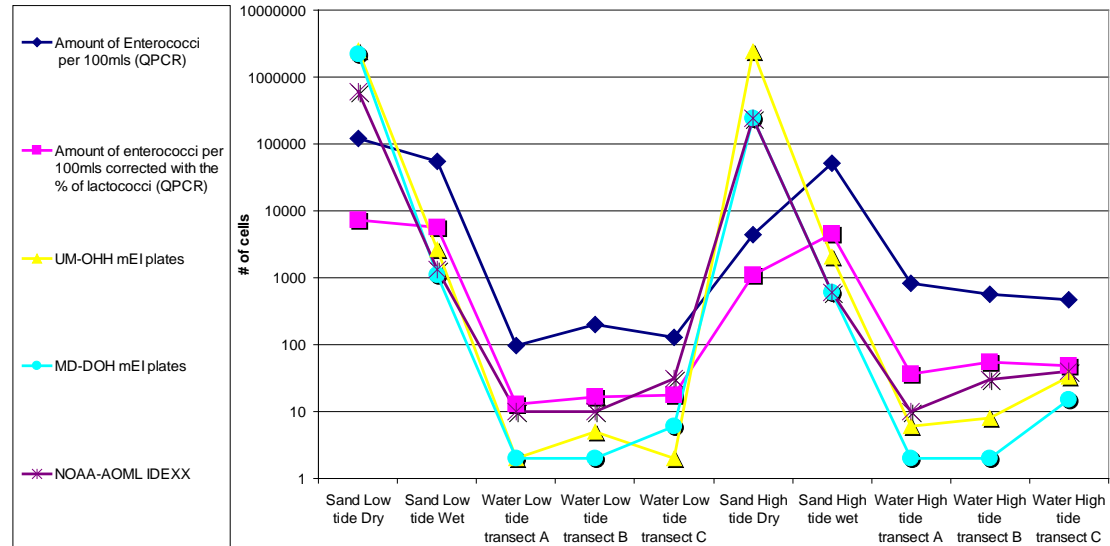
Nutrients and Flow, Boynton Inlet 2nd 48-hour Intensive study, September 2007

(Zheng et al.)

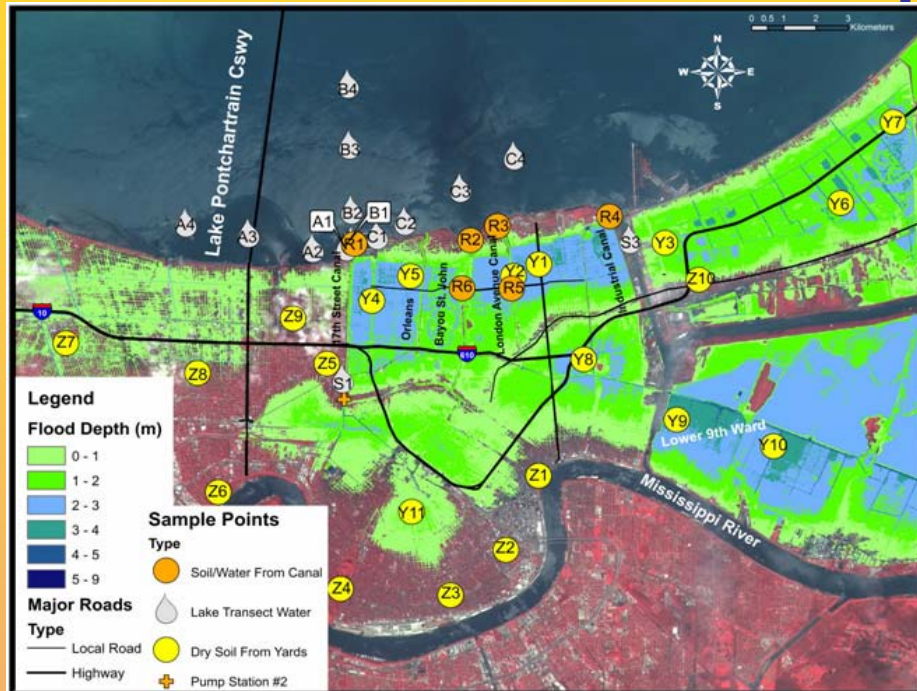
Beach Environmental Assessment and Characterization of Human Exposure Study (BEACHES)



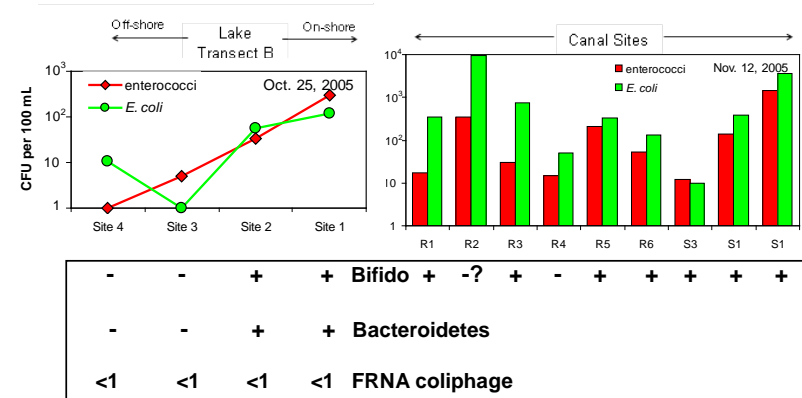
Enterococci results for Hobie beach 07/18/07



Impact of Hurricanes Katrina and Rita on the Microbial Landscape of New Orleans:

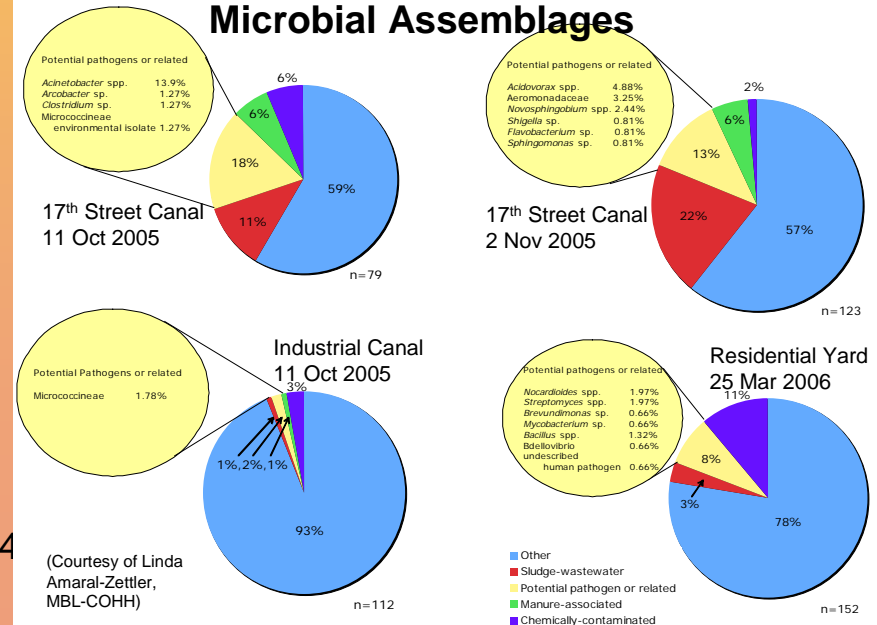


Traditional & Non-Traditional Indicators



Non-Traditional Indicators

Microbial Assemblages

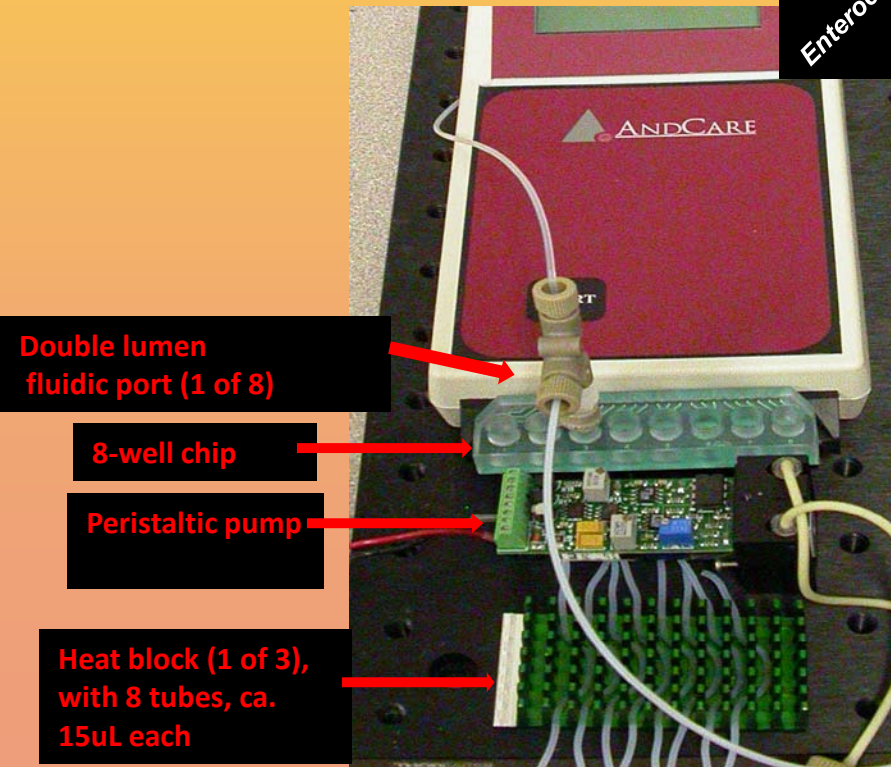
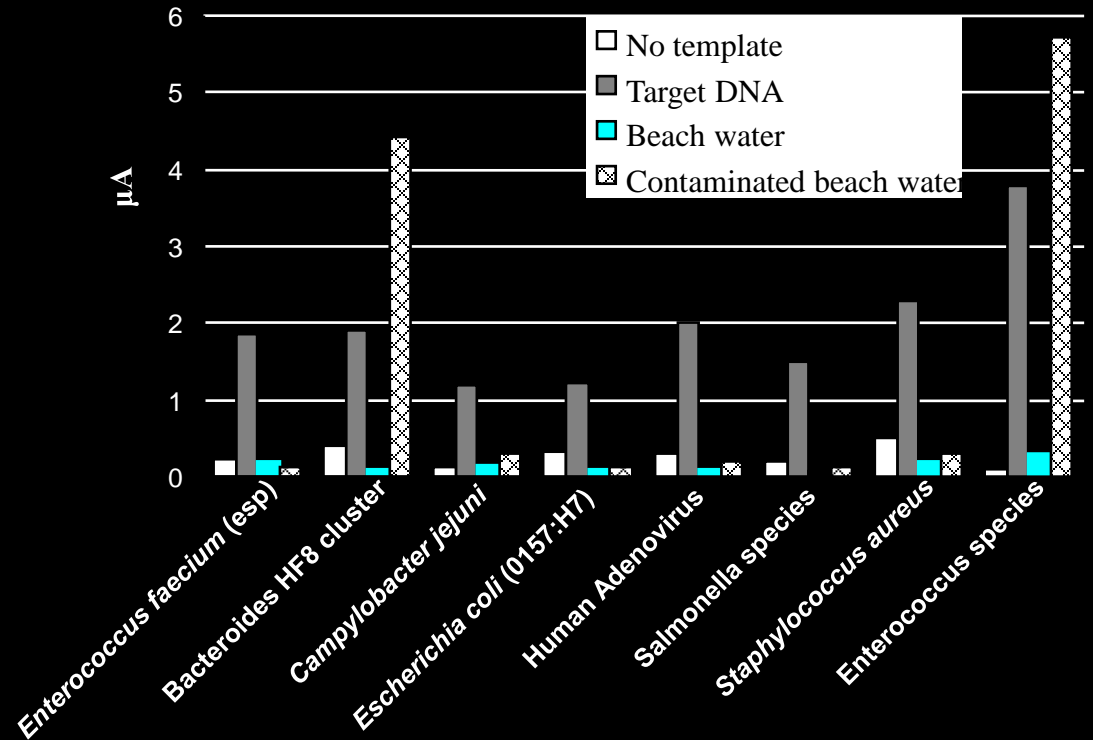


- Reported in: Sinigalliano et al., 2007, PNAS 104:9029-9034
- Reviewed in: Dobbs, 2007, PNAS 104:9103-9104

Portable & In-Situ Biosensors



Prototype handheld automated multiwell PCR analyzer and electrochemical detector, with PC interface



The Product:

biotechnological innovations to produce
timely & accurate reporting of ecosystem
health and environmental safety



apid, Reliable, Affordable
Biosensing

Health risks?

Questions?

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- NGI

