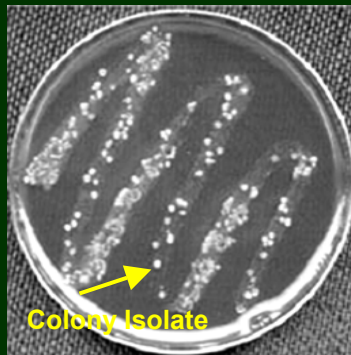


# BST? MST? CST?

## Do they help?



An example of the streak-plate method. A single bacterial colony isolate (single-species) is indicated.

# Watershed Assessment

Determine the watershed DA of the survey area.

Determine Land use.

Estimate population of the survey area.

Windshield survey possible pollution sources paying particular attention to the following:

- Percentage of impervious surface in the watershed

- Marinas and docking facilities.

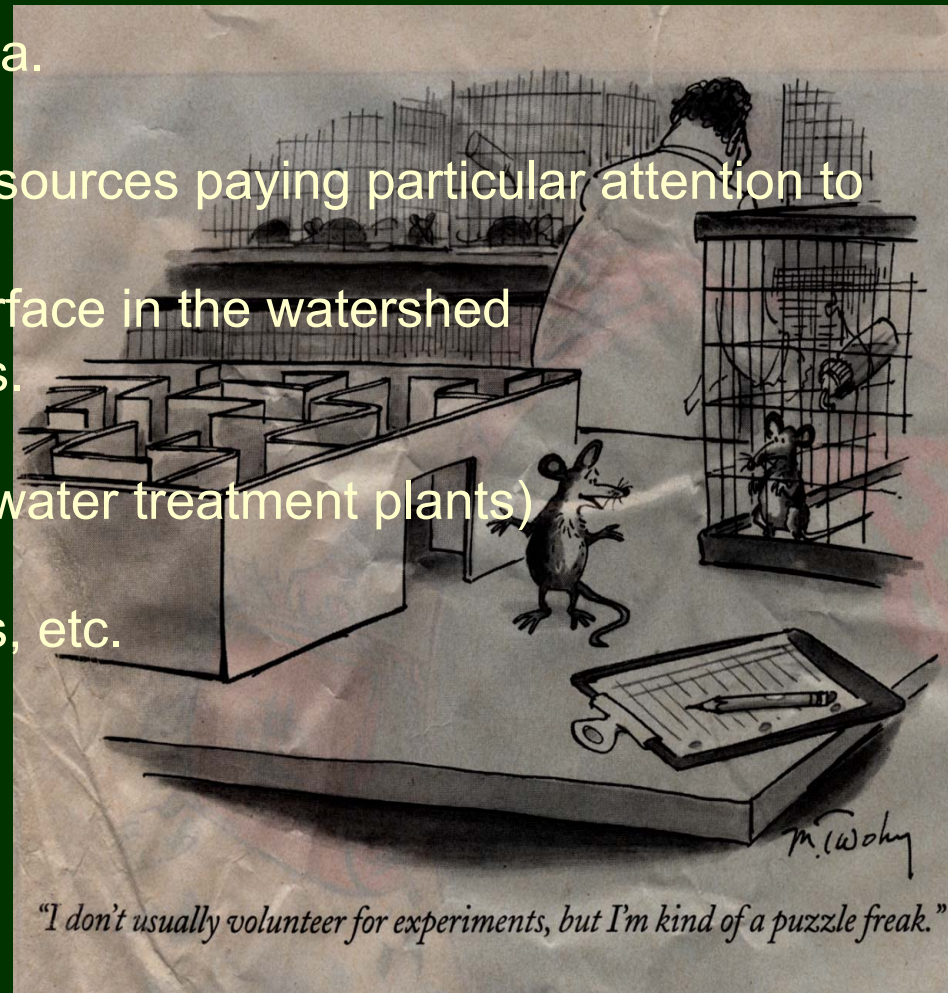
- Wastewater treatment plants

- Package plants (small wastewater treatment plants)

- Livestock Agriculture

- Domestic pets, i.e. cats, dogs, etc.

- Wildlife



# Watershed Assessment, cont'd. . . .

## Expand the inspection by going door-to-door.

Pay particular attention to the following:

- Septic systems (Look for Signs of Septic System Failure)
- Stormwater drains
- Ditches, canals

## Signs of Septic System Failure

\* **Wet spots or standing water** - Areas that remain wet after rain events or appear without precipitation should be investigated, since effluent can seep up from a failing system and puddle on the ground surface.

\* **Odd growth patterns** - Green lines of vegetative growth on the drainfield that follow the path of the pipes are normal; however, green patches not in line with pipes, random patches of lush growth, and bulls-eye patterns (dead areas surrounded by green growth) all indicate problems in the drainfield.

\* **Slow drains or backups** - If wastewater is slow to drain and the plumbing is working properly, there may be a problem with backup in the septic tank.

\* **Septic odor** - Easy to detect, odor is one of the best problem indicators

# Microbial Source Tracking Methods Classification

## Non-molecular Library-dependent Methods

Antibiotic Resistance Methods

Carbon Source Profiling (Nutritional Analysis)

FAME (Sterols/Fatty Acid Methyl Ester Analysis)

## Non-molecular Library-independent Methods

Fecal Bacteria Ratios

Streptococcal Population Profiles

Phenotypic Species-Specific (Host-Specific) Indicators

F+ coliphage serotyping

Enterotoxin Biomarkers

## Molecular Library-dependent Methods

Rep-PCR (Polymerase Chain Reaction)

PFGE (Pulsed-Field Gel Electrophoresis)

Ribotyping

RAPD (Randomly Amplified Polymorphic DNA)

tRFLP (Terminal Restriction Fragment Length Polymorphism)

## Molecular Library-independent Methods

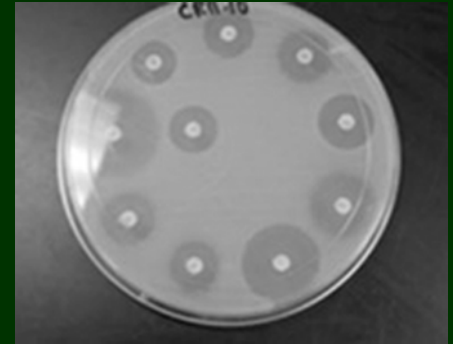
Bacteriophage Indicators

Virus (Human Pathogen) Indicators

Genotypic Species-Specific (Host-Specific)

Indicators

Bacterial Endemism and Co-Speciation



Antibiotic resistance profile analysis of an *E. coli* isolate.

The white disks on the plate are impregnated with different antibiotics and the clear zones of bacterial growth inhibition are measured to determine the resistance of each bacterial isolate.

# Chemical-Based Source Tracking Methods

## *Optical Brighteners/Detergents*

Whitening agents and optical brighteners are chemicals used in laundry detergents. They are often a large component of grey water discharge and have been used as indicators of sewage fallout with varying degrees of success

## *Caffeine*

Caffeine detection is a fairly new source tracking method. Caffeine is generally found in highest concentrations in highly urbanized areas; however, urbanization levels have not yet been matched to known caffeine concentrations in the environment.

## *Coprostanol*

Coprostanol is a byproduct of the breakdown of cholesterol and is present in human and some mammal species. Human secretion of coprostanol can be inconsistent but in general it is a good indicator of human fecal pollution.

## ***The Toolbox Approach or get a bigger hammer . . .***

Methods are complimentary methods rather than opposing.

No magic bullet. Better approach may be to combine several molecular and non-molecular methods. By using this “toolbox” approach, researchers can:

- Tailor source tracking methodologies to individual study needs and limitations;
- Increase statistical confidence by obtaining similar results from different methods;
- Expand research results by identifying sources from one method that the other was unable to identify; and supplement source tracking methods that give only qualitative or quantitative information.

# Monitoring and TMDL Modeling Techniques to Assess Bacterial Loading in Estuarine Environments and Improve Management Programs

## Cooperators and Participants

NCSU College of Design

NCSU CMAST

NCSU Water Quality Group

NC Division of Health, Shellfish Sanitation Program

Duke Marine Laboratory

NOAA/NOS Center for Coastal Environmental Health and Bimolecular Research

UNC-Institute of Marine Sciences

Carteret Craven Electric Cooperative

Jumping Run Creek Watershed Citizens

Croatan National Forest

Open Grounds Farm

USDA-CSREES

NC DENR Division of Water Quality, 319 Program

NC Clean Water Management Trust Fund

NC Wetland Restoration Program



# Project Objectives

1. Implement and <sup>call</sup> assess BST technology in <sup>104</sup> NC as part of watershed-based toolbox approach to reduce bacterial loading.
2. Review TMDL process using BST / Toolbox data.
3. Review / recommend management strategies.
4. Disseminate information to coastal local governments.
5. Incorporate information into coastal environmental management / policy curriculum.

← 2027  
← 2322  
← 1721  
← 2559  
← 946  
← 23130

5 4 3 2 1 2 6 \* 20 19 18 17 16 15 14 13 12 11 10 9 8 7 6 5 4 3 2 1



# Jumping Run

Total drainage area:

~800 acres

Mobile and RV

Housing

Low density single family residential

Industrial

No agriculture

Sources? **Dogs, cats, waterfowl, domestic wildlife, septic tanks.**

Transport vectors? **Ditches, surface runoff, surficial groundwater, airborne, direct deposit.**



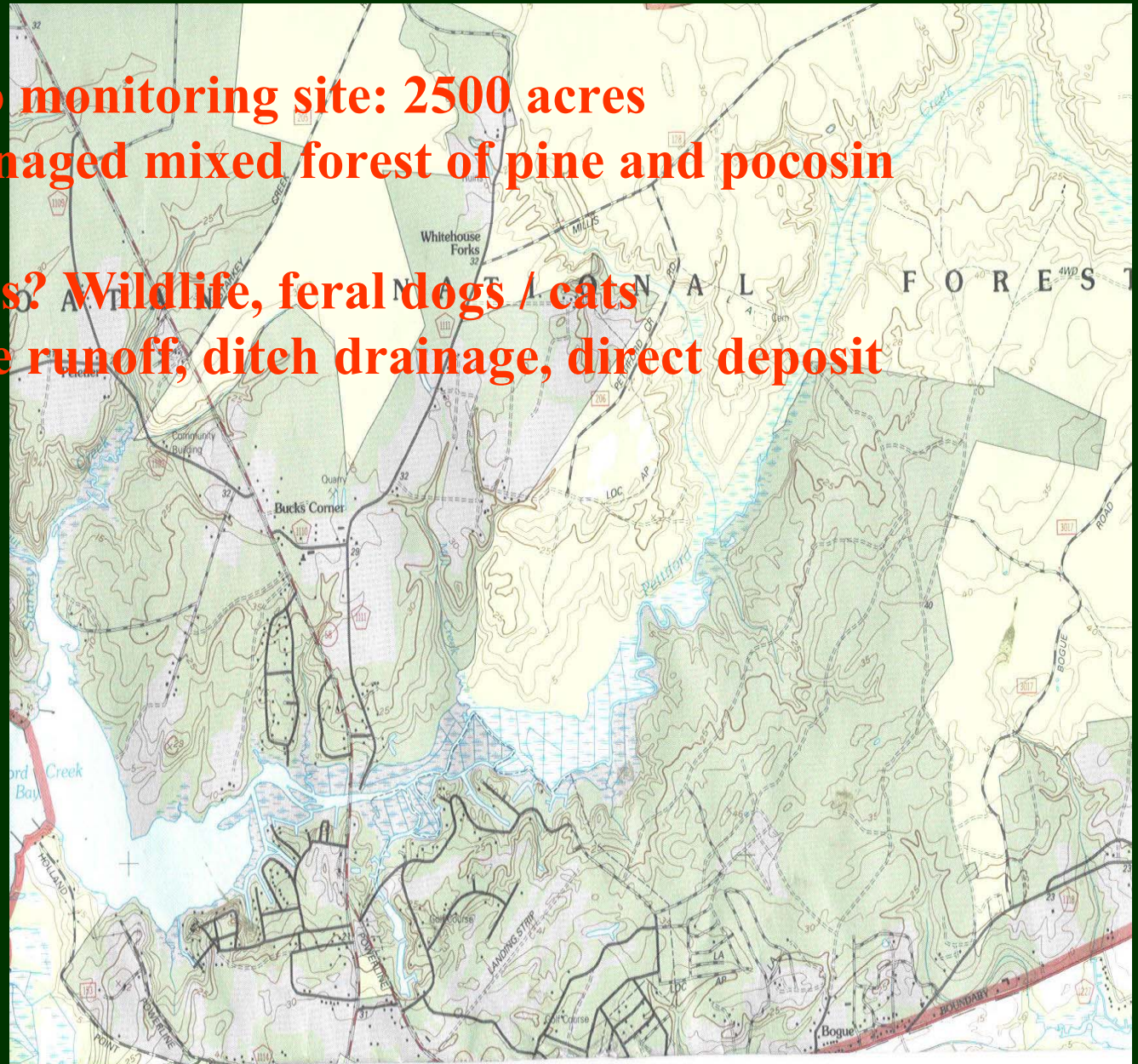
# Pettiford Creek-Croatan National Forest

**Drainage area to monitoring site: 2500 acres**

**Land cover: Managed mixed forest of pine and pocosin**

**Bacterial Sources? Wildlife, feral dogs / cats**

**Vectors? Surface runoff, ditch drainage, direct deposit**



# South River: Open Grounds Farm

Drainage Area to monitoring site: 3000 acres

Land cover: Cultivated row crop agriculture

Bacterial Sources? Wildlife, waterfowl and other birds, rodents, dogs  
Transport vectors? Surface runoff, ditch flow, direct deposit.



# Methods: Integration of watershed - based field assessment and laboratory techniques

## Watershed Assessment:

Land use / land cover surveys

Flow Monitoring

Time, travel, dilution studies

Rainfall / runoff measurements

Water sampling: Sound and base flow grabs,  
flow-weighted storm event for fecal and N, P, TSS

## Laboratory:

MPN quantification of fecal and water samples

e. coli speciation for fecal and water samples

Watershed-based fecal source library for MAR and DNA

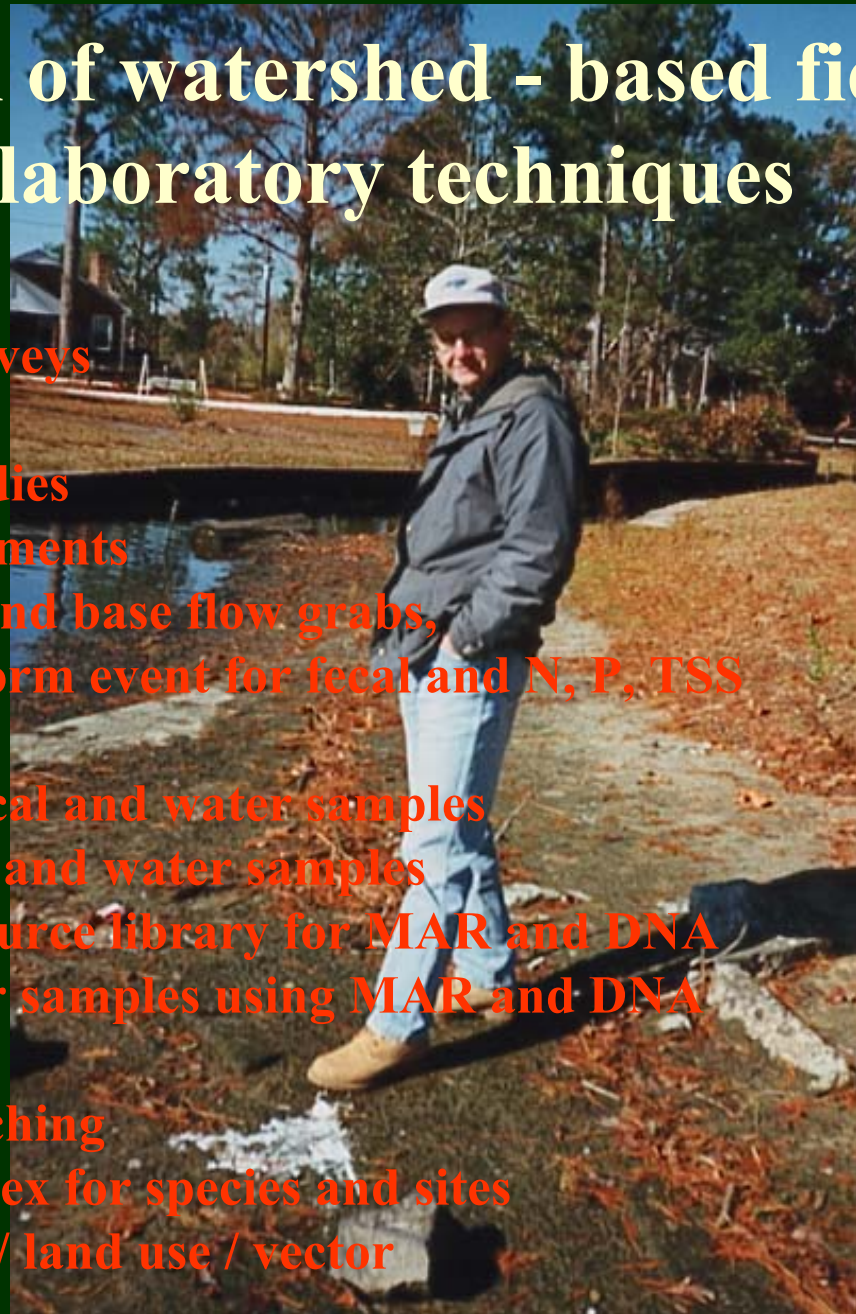
Characterization of water samples using MAR and DNA

## Analyses:

Spatial and seasonal matching

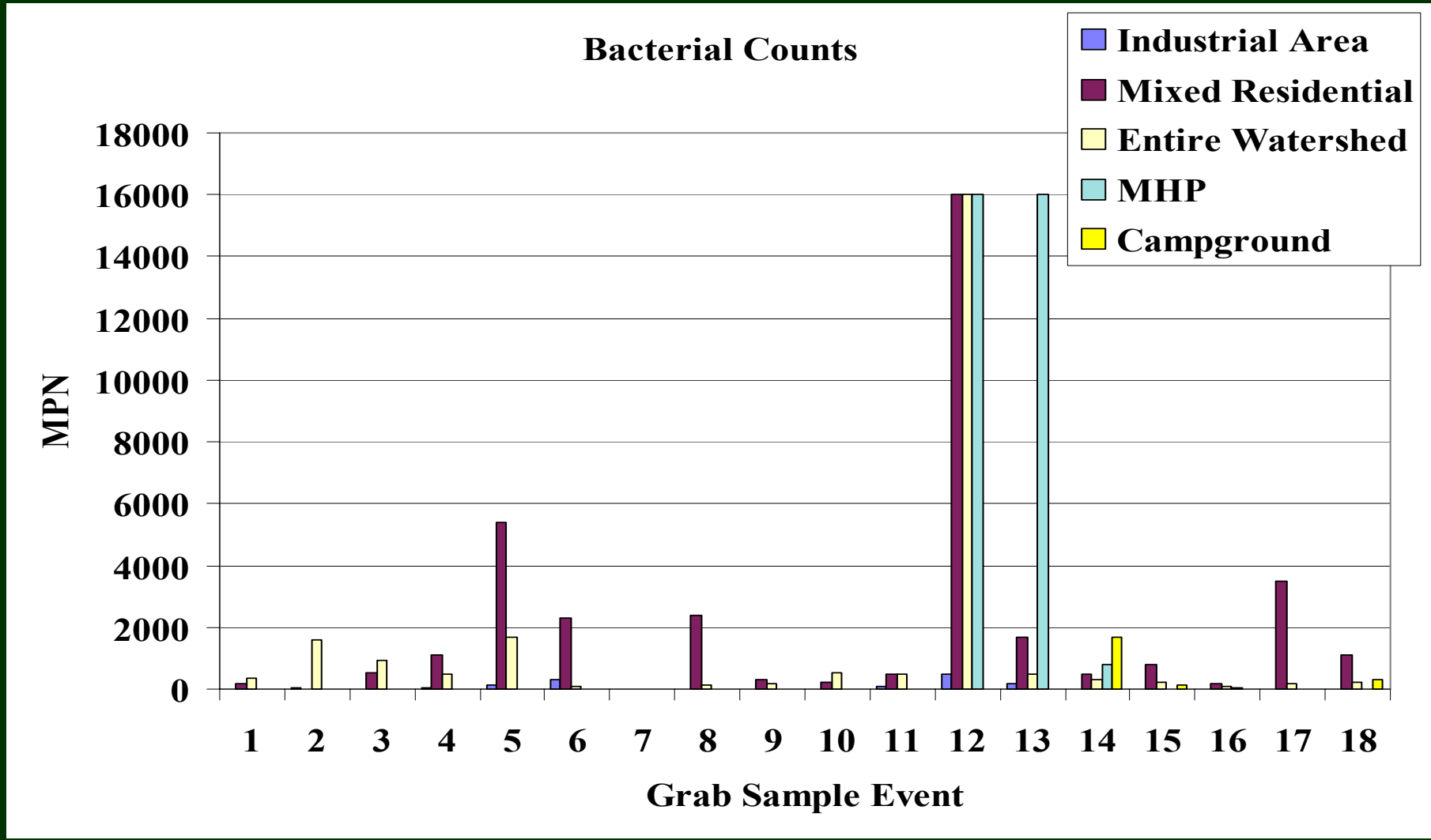
Resistance/ sensitivity index for species and sites

Loading analysis by type / land use / vector



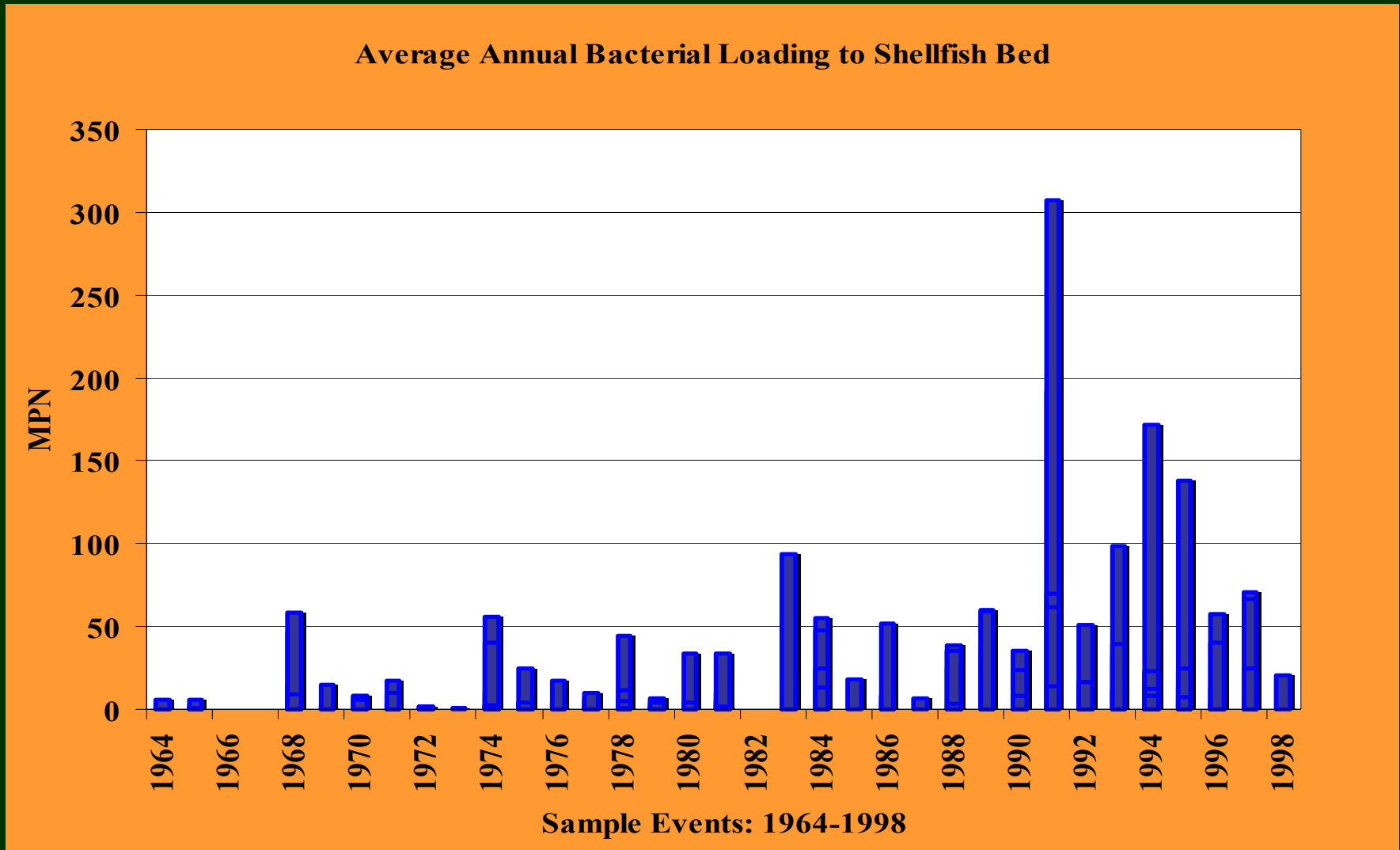
# Tailwater grab samples: Jumping Run

❖ Used to quantify bacterial densities and MAR/ DNA profile for the draw-down/ baseflow component of the hydrograph.



# Sound – based grab sample data: Jumping Run

❖ Will be used to quantify bacterial density and the MAR/ DNA profiles in the shellfish beds.



# Watershed Assessment: Jumping Run

❖ Used to spatially assess bacterial sources, transport vectors, and to ground truth land use / land cover

Location	Visits	SV	Dogs
MHP	47	0	23
Med. Density	66	2	60
Low Density	14	0	7
Campground	1	0	0
<b>Totals</b>	<b>128</b>	<b>2</b>	<b>90</b>

\*Does not include Roads

Location	Cats	Other	Imperviousness
MHP	6	3	40,000sft
Med. Density	47	25	185,000
Low Density	0	0	95,000sft
Campground	0	0	5,000sft
<b>Totals</b>	<b>53</b>	<b>28</b>	<b>325,000sft</b> or 7.5 acres





## Land Cover Change Assessment: Jumping Run

- ❖ Develop land use loading rates for TMDL assessment.
- ❖ Information is used to help direct and design restoration and management strategies for testing.





# Results: Dye/ Time and Travel Studies Jumping Run



❖ Information is using to determine ditch drainage patterns, water movement direction and timing, as well as dilution and dispersion

Date	Guage Ht	CFS	Time Travel
21-Jan-98	1.38'	8.4	n/a
22-Jan-98	1.18'	5.8	MHP-Outlet, 5hrs
04-Feb-98	3'	18.2	MHP-Gauge, 2.5hrs
23-Feb-98	1.7'	10.9	n/a
03-Mar-98	1.13'	3.21	Headwaters-Gauge, 3hrs.
04-Mar-98	1.10'	2.73	CmpGrnd to Outlet, 1.5hrs.
03-Apr	0.95	5.96	n/a
04/17/1998	0.9	10.4	n/a
04/30/1998	0.84	5.1	n/a
06/18/1998	0.76	4.6	n/a
07/14/1998	0.72	4.5	n/a



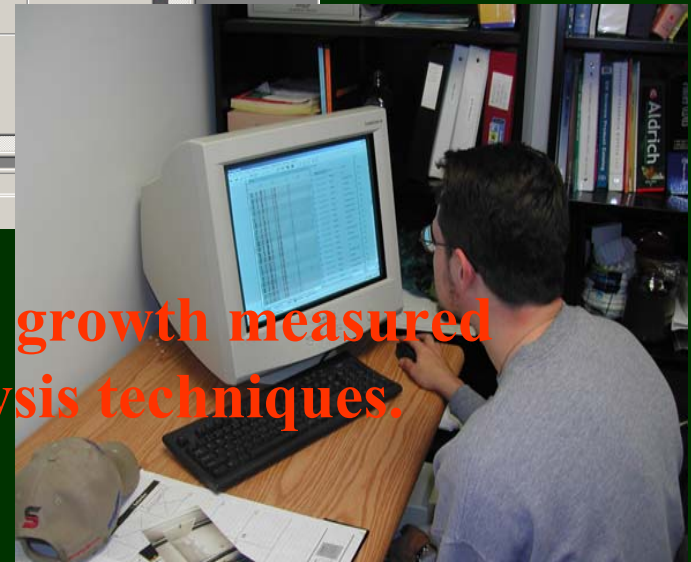
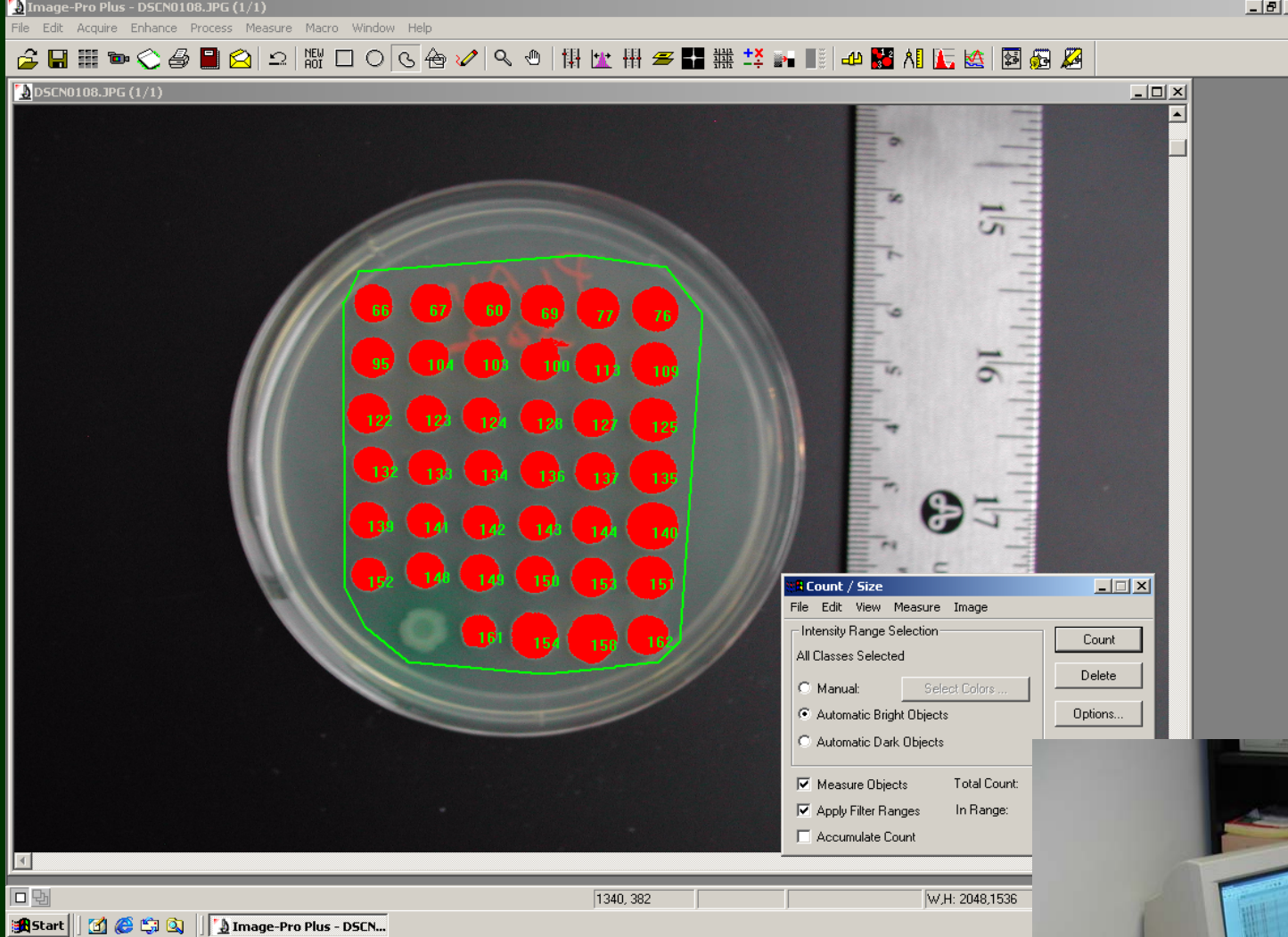
# Storm Event Monitoring



## MAR and DNA Data

MPN is calculated for both water / fecal library samples to link densities with water volume and flow.



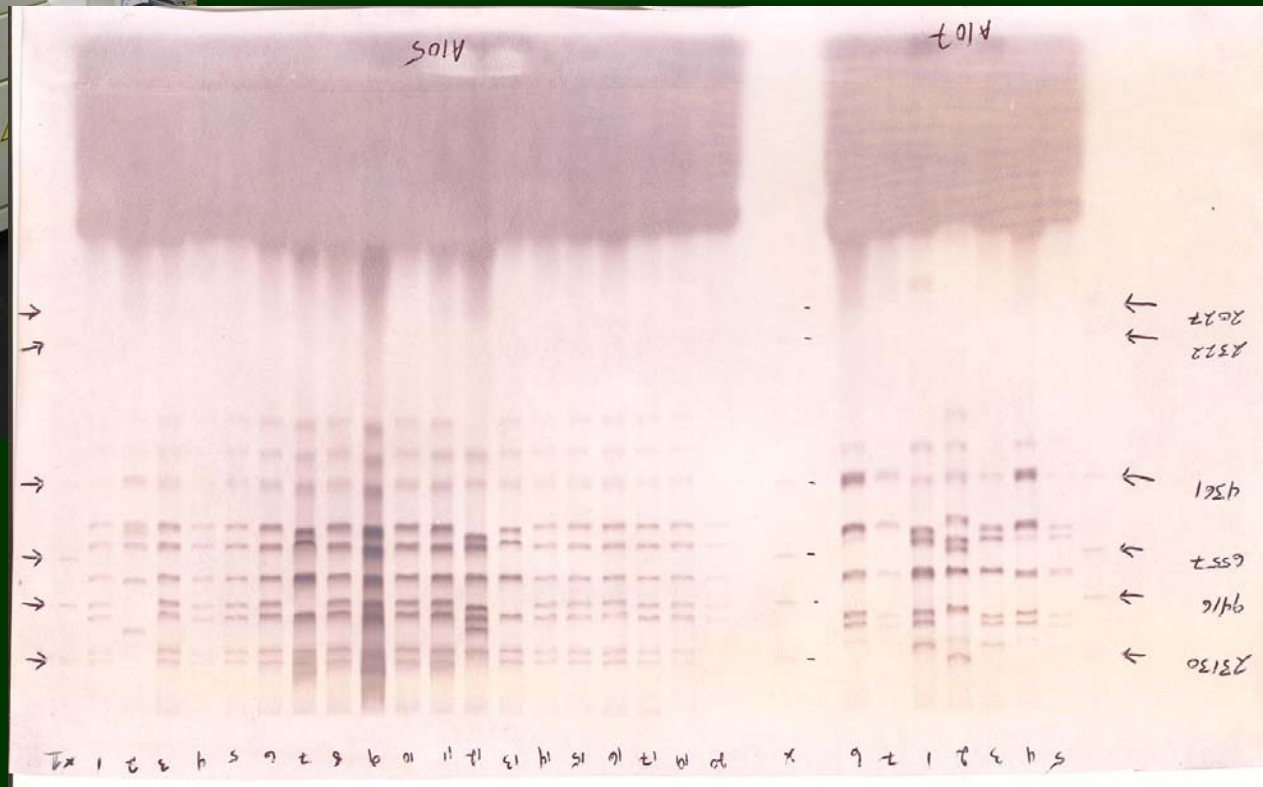


**Plates are digitally photographed and cell growth measured relative to two controls using image analysis techniques.**



**DNA analysis is conducted on all library and resistant water samples and 50% of the sensitive water samples.**

**Tracks of DNA material are delineated and banding patterns numbered to create a profile for each sample.**



## **Preliminary MAR Results**

**Jumping Run: 27 Water samples, Avg MPN: 3000 / 100ml, 178 isolates**

**Sensitivity index: 48 or 26 % were 10% resistant**

**4 or 2% were 20% resistant**

**7 or 0.7% were greater than 30% resistant**

**Croatan: 8 water samples, Avg MPN: 326 / 100ml, Number of isolates: 50**

**Sensitivity index: 8 or 16% were 10% resistant**

**1 or 2% were 20% resistant**

**None greater than 20%**

**Open Grounds: 6 water samples, Avg MPN 4300/ 100ml, 64 isolates**

**Sensitivity index: 17 or 26% were 10% resistant**

**1 or 1% were 20% resistant**

**1 or 1% were 40% resistant**

**In progress. . .**

**Conducting the matching and loading analyses.**

**Refining MAR profiling.**

**Adding new research components, of course!**



# Recommendations

The “toolbox” method appears to be the most effective use of these methodologies at present.  
***The best approach is to evaluate land uses and sources under investigation, and tailor the research methods to fit each individual situation.***

- Source types (human, non-human, livestock, domestic pets, or wildlife);
- Pollutant loading sources and delivery vehicles (point sources, nonpoint sources);
- Sample medium (marine, freshwater, groundwater, sediments, shellfish tissue);
- Level of source resolution needed (human vs. non-human or individual categories for all); and
- Cost for each method (usually estimated on a cost per isolate basis)
- Library building needs and capabilities.

**Best comprehensive source of information:**

**[Addressing microbial pollution: A reference for local governments](#)**

**Produced by NC NERR for NC DENR DWQ Coastal Nonpoint Source Program**

**<http://ncnerr.org/ccs/publications/index.html>**

**Contact Whitney Kurz at [whitney\\_kurz@ncnerr.org](mailto:whitney_kurz@ncnerr.org)  
North Carolina National Estuarine Research Reserve  
North Carolina Division of Coastal Management**

**phone 252.728.2170**

**fax 252.728.6273**

