

American Alligator Bacterial Source Tracking in Coastal South Carolina



Michelle A. Johnston, Ph.D.
John A. Knauss Marine Policy Fellow
November 10, 2010

Purpose

- To determine if American alligators (*Alligator mississippiensis*) are a source of fecal coliform and potential pathogens in coastal South Carolina surface waters



Outline

- Introduction
- Hypothesis
- Objectives
- Design and Methods
- Results
 1. Bacterial Identification
 2. Genotypic Fingerprinting
 3. Bacterial Source Tracking
- Conclusions
- Acknowledgements



Wilfredo Lee

Water Quality Degradation

- Half of population in coastal communities
- Bacterial pollution
- Various contributing sources
- Potential pathogen evaluation with fecal coliform indicators
- Helps detect unsafe water conditions



Indicator Limitations

- Regulatory microbiology
- Oversimplification
- EPA evaluation of methods
- No source identification
- Identification for management
 - Endotherms main source



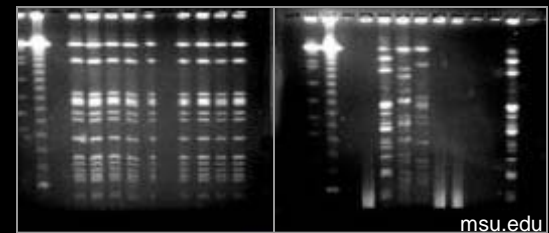
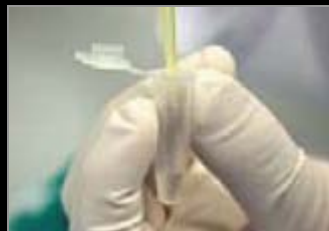
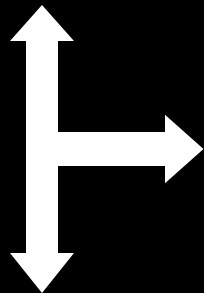
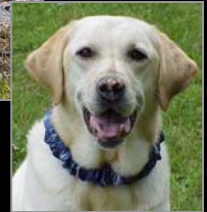
American Alligator



Alligator mississippiensis



Bacterial Source Tracking (BST)



Scat Fingerprints

Water

Hypothesis and Objectives

- H₁: The American alligator is a source of bacterial contamination in South Carolina surface waters
 - i. Determine if alligators contribute to fecal coliform bacterial water quality degradation
 - ii. Determine density of alligator fecal coliforms
 - iii. Evaluate bacterial identification tools and BST fingerprinting tools
 - iv. Determine fingerprint reproducibility and stability
 - v. Evaluate BST database stability

Design and Methods: Study Area



Alligator Sampling Techniques

Distress Call



Alligator Sampling Techniques

Snare and Roll



Alligator Sampling Techniques

Cloacal Swab



Alligator Sampling Techniques

Water Sample, Tag, and Release



- Palmetto Bluff
– 2006 (n=31)



Research Component 1

Bacterial Identification

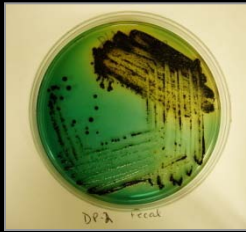


=



Sample Processing

Culture



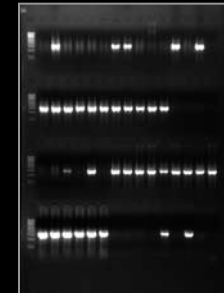
DNA Extraction



PCR



PCR Products

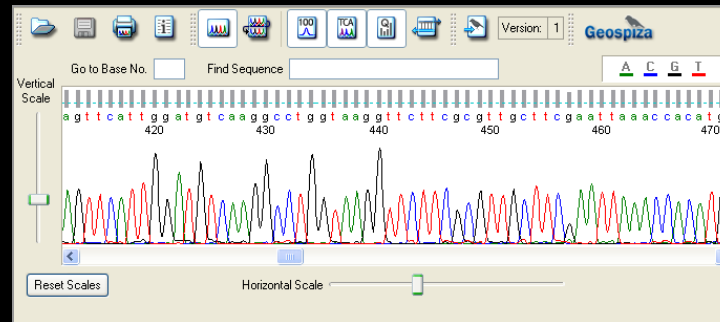


**Sequencing
Identification**



Nat. Center for Biotech. Info.,
Basic Local Alignment Search Tool

16S rRNA gene Sequencing

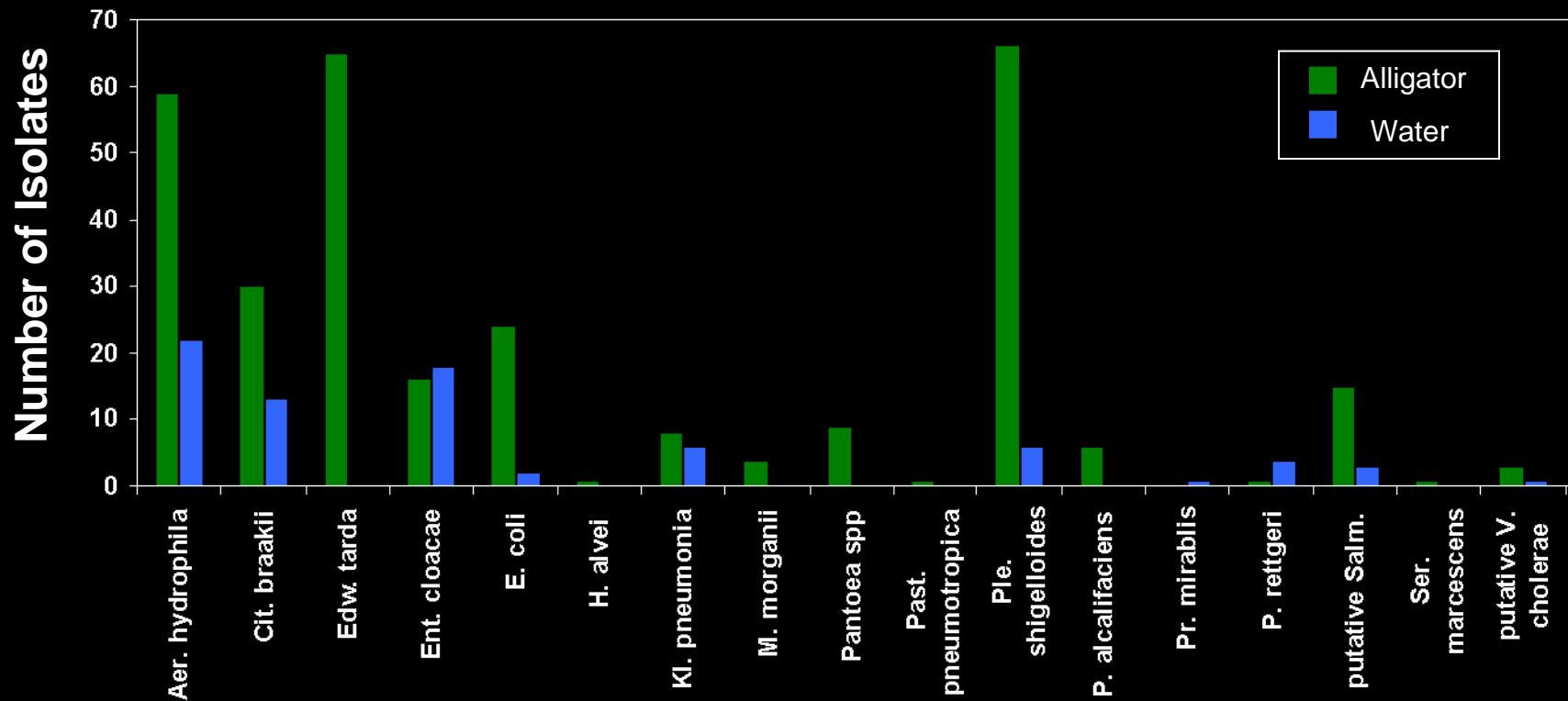


Purification



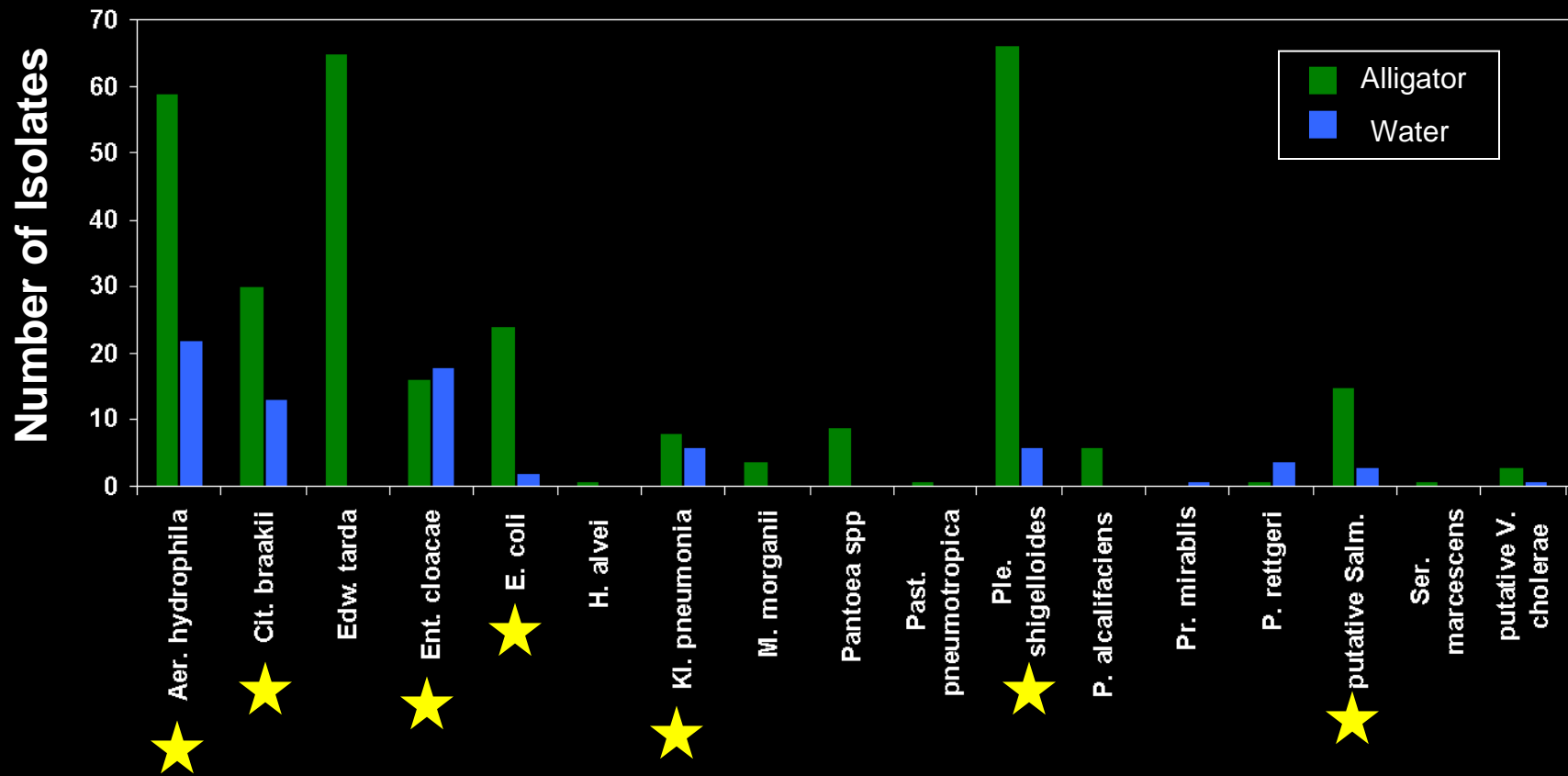
Bacteria Identification Results

2006 Identified Bacteria from Alligator and Water Isolates



Bacteria Identification Results

2006 Identified Bacteria from Alligator and Water Isolates



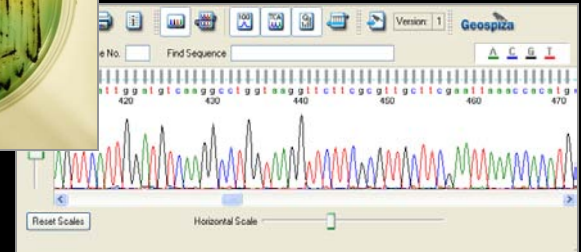
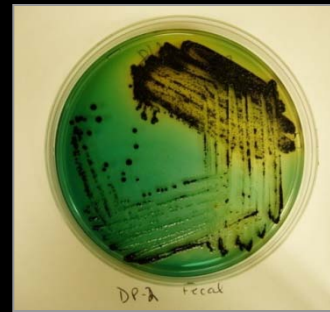
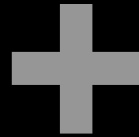
Bacterial Density Results

- Fecal Coliform Most Probable Number (MPN):

Species	Fecal Coliform MPN (Density/g feces)	Source
Alligator	8.0×10^9	Johnston et al., 2010
Duck	3.3×10^7	Schueler and Holland, 2000
Human	1.3×10^7	Schueler and Holland, 2000
Dog	2.3×10^7	Schueler and Holland, 2000
Turtle	1.6×10^6	Harwood et al., 1999
Cow	2.3×10^5	Schueler and Holland, 2000

Research Component 2

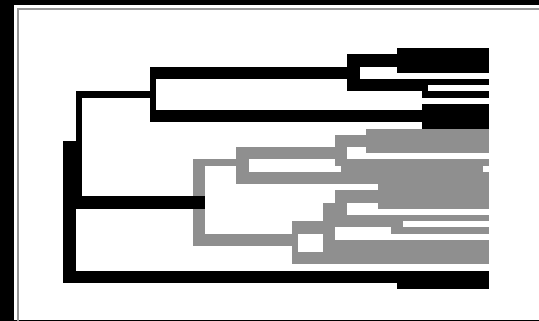
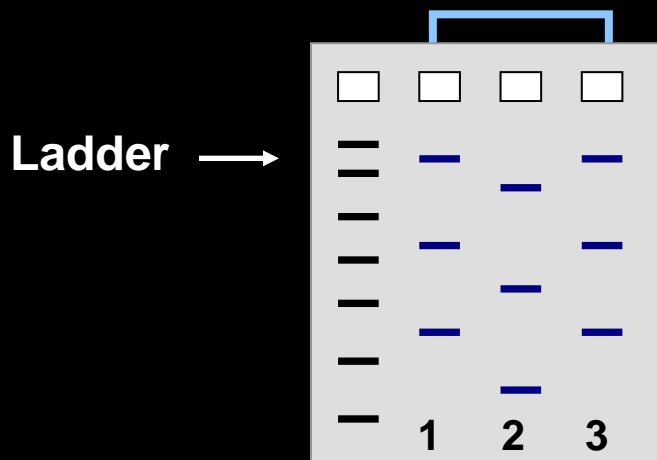
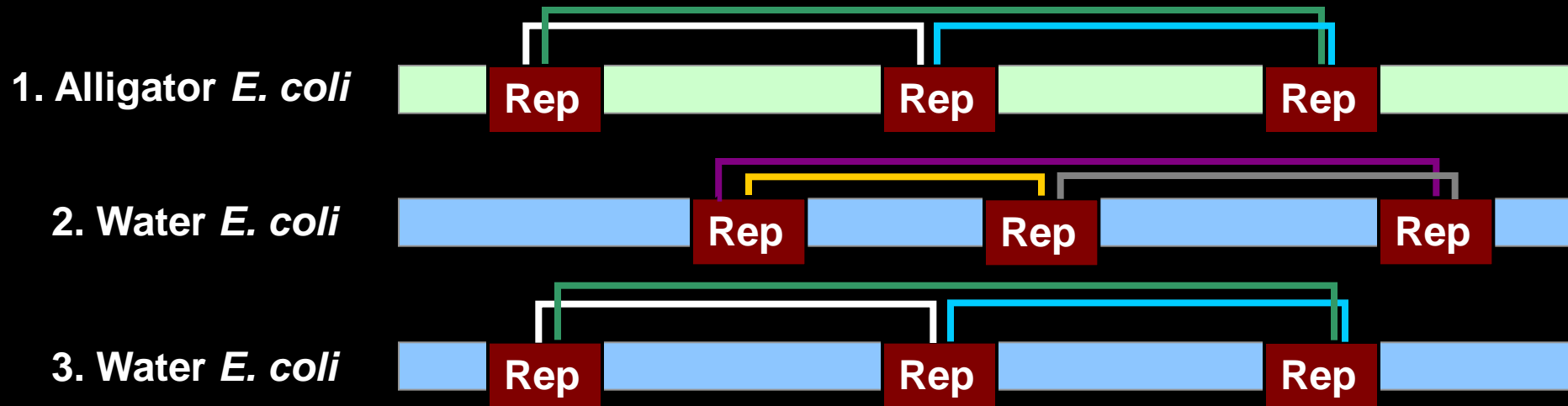
Genotypic Fingerprinting



Genotypic Fingerprinting

Source	Isolate ID	No.	Source	Isolate ID	No.
Alligator	<i>Aeromonas hydrophila</i>	31	Chicken	<i>Escherichia coli</i>	6
	<i>Aeromonas punctata</i>	1	Cow	<i>Escherichia coli</i>	5
	<i>Aeromonas veronii</i>	24	Dog	<i>Escherichia coli</i>	5
	<i>Citrobacter freundii</i>	92	Horse	<i>Escherichia coli</i>	10
	<i>Enterobacter aerogenes</i>	3	Pig	<i>Escherichia coli</i>	5
	<i>Enterobacter cloacae</i>	21	Raccoon	<i>Escherichia coli</i>	1
	<i>Escherichia coli</i>	41	Sea Turtle	<i>Citrobacter freundii</i>	3
	<i>Klebsiella planticola</i>	5		<i>Enterobacter aerogenes</i>	1
	<i>Klebsiella pneumoniae</i>	10		<i>Escherichia coli</i>	1
	<i>Plesiomonas shigelloides</i>	78		<i>Salmonella enterica</i>	1
	<i>Salmonella enterica</i>	7			
	Total	313		Total	38

Genotypic Fingerprinting



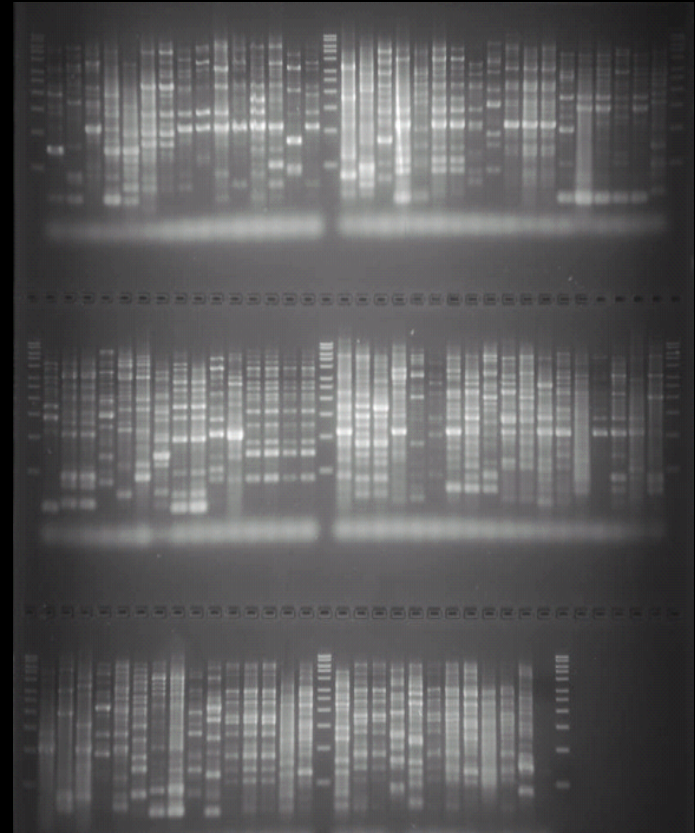
Repetitive Extragenic Palindromic-PCR

- REP-PCR Primers for entire genome:

Primer	Sequence
REP1R-I	5'- III ICG ICG ICA TCI GGC -3'
REP2-I	5'- ICG ICT TAT CIG GCC TAC -3'
ERIC1R	5'- ATG TAA GCT CCT GGG GAT TCA C -3'
ERIC2	5'- AAG TAA GTG ACT GGG GTG AGC G -3'
BOX A1R	5'- CTA CGG CAA GGC GAC GCT GAC G -3'
(GTG) ₅	5'- GTG GTG GTG GTG GTG -3'
Source:	Versalovic et al., 1994

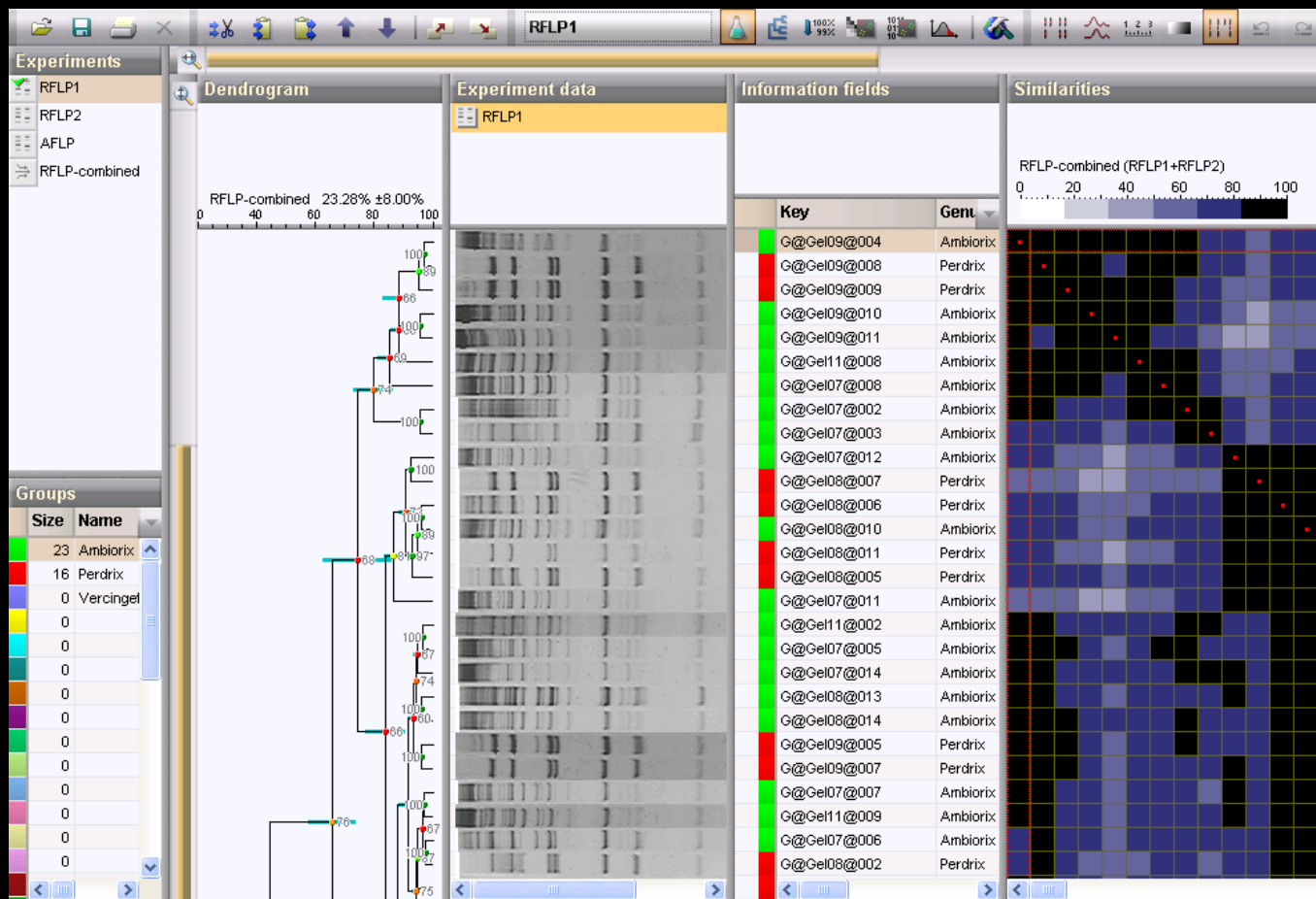
Research Component 2: Methods

- REP-PCR
- Gel Electrophoresis
- EtBr stain and image



Research Component 2: Results

- Gel Compar II Similarity matrix (Pearson Correlation)



Research Component 2: Results

- Percentage of alligator isolates assigned to each group using Jackknife analysis

Percentage (%) of Alligator Isolates Correctly Assigned to Each Source Group

Isolate Group	<i>Aer. spp.</i>	<i>Cit. freundii</i>	<i>Ent. spp.</i>	<i>E. coli</i>	<i>Kl. spp.</i>	<i>Ple. shigelloides</i>	<i>Salm. enterica</i>
<i>Aer. spp.</i>	80.4	3.3	4.6	0.0	0.0	1.3	0.0
<i>Cit. freundii</i>	7.1	90.2	4.6	2.4	13.3	1.3	0.0
<i>Ent. spp.</i>	7.1	2.2	63.6	2.4	0.0	2.6	0.0
<i>E. coli</i>	3.6	2.2	4.6	90.2	20.0	1.3	0.0
<i>Kl. spp.</i>	0.0	0.0	4.6	2.4	66.7	0.0	0.0
<i>Ple. shigelloides</i>	1.8	2.2	18.2	2.4	0.0	93.6	0.0
<i>Salm. enterica</i>	0.0	0.0	0.0	0.0	0.0	0.0	100.0

Values in boldface represent the rate of correct classification (RCC). Average RCC was 84%.

Research Component 2: Results

- Percentage of *E. coli* isolates assigned to each group using Jackknife analysis

Percentage (%) of *E. coli* Isolates Correctly Assigned to Each Source Group

Source	Alligator	Chicken	Cow	Dog	Horse	Pig	Raccoon	Sea Turtle
Alligator	95.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Chicken	0.0	100.0	0.0	0.0	0.0	0.0	0.0	0.0
Cow	2.4	0.0	100.0	0.0	0.0	0.0	0.0	0.0
Dog	0.0	0.0	0.0	100.0	0.0	0.0	0.0	0.0
Horse	0.0	0.0	0.0	0.0	100.0	0.0	0.0	0.0
Pig	0.0	0.0	0.0	0.0	0.0	100.0	0.0	0.0
Raccoon	0.0	0.0	0.0	0.0	0.0	0.0	100.0	0.0
Sea Turtle	2.4	0.0	0.0	0.0	0.0	0.0	0.0	100.0

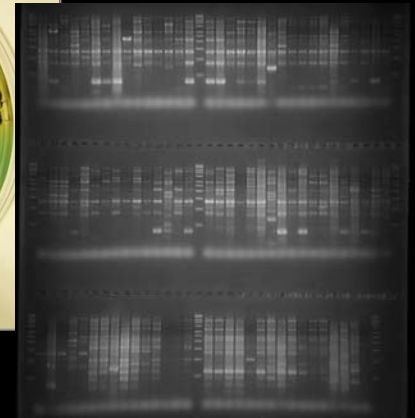
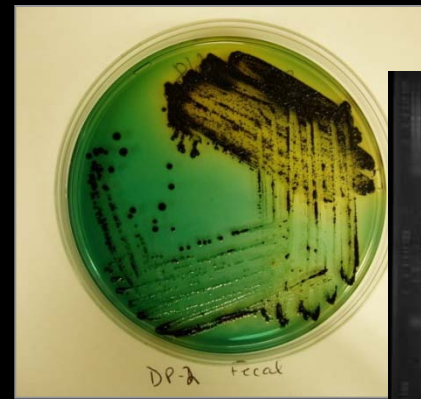
Values in boldface represent the rate of correct classification (RCC). Average RCC was 99%.

Research Component 3

Bacterial Source Tracking



+



=



Alligator Sampling

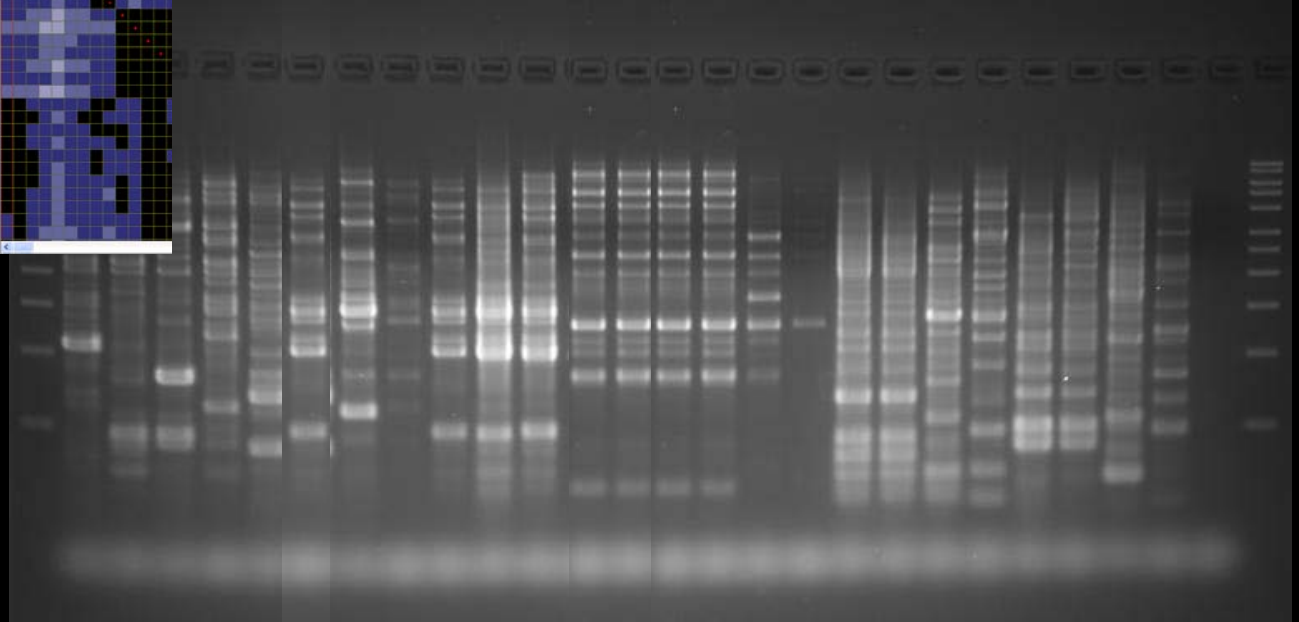
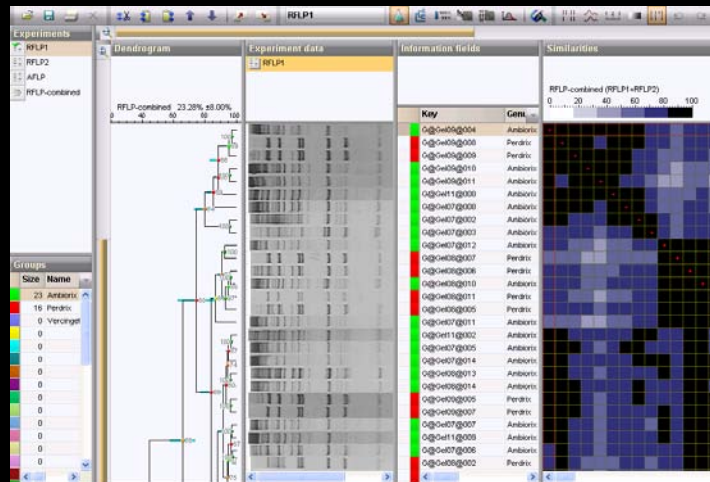


- Palmetto Bluff
 - 2006 (n=31)
 - 2009 (n=4)
- Spring Island
 - 2009 (n=4)



Research Component 3: Results

- Generated 705 alligator and water fingerprints



Research Component 3: Results

- 2006 matches

2006 Bacteria	Water to Alligator Matches	Total Water Isolates	Total Percent (%) Matched
<i>Aer. spp.</i>	2	23	9
<i>Cit. freundii</i>	7	19	37
<i>E. coli</i>	9	24	38
<i>Ent. spp.</i>	3	28	11
<i>Kl. spp.</i>	2	9	25
<i>Ple. shigelloides</i>	3	7	43
<i>Salm. enterica</i>	0	0	0
Pooled	26	110	24

Research Component 3: Results

- 2009 matches

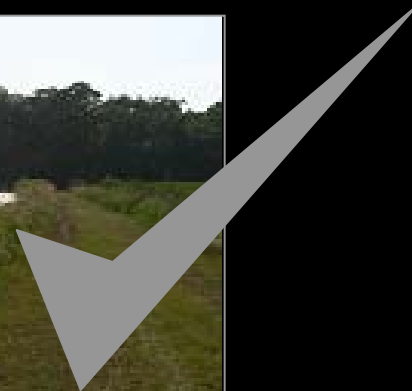
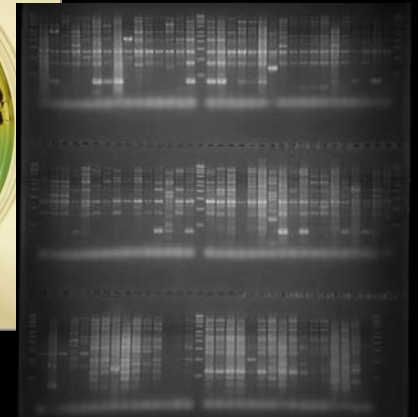
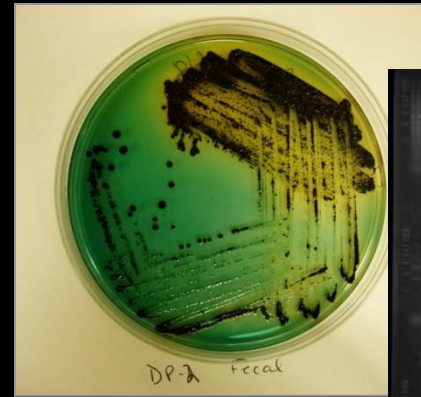
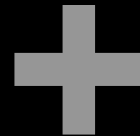
2009 Bacteria	Water to Alligator Matches	Total Water Isolates	Total Percent (%) Matched
<i>Aer. spp.</i>	0	Unknown	0
<i>Cit. freundii</i>	5	Unknown	2
<i>E. coli</i>	4	Unknown	1.5
<i>Edw. tarda</i>	2	Unknown	1
<i>Ent. spp.</i>	0	Unknown	0
<i>Kl. spp.</i>	0	Unknown	0
<i>Ple. shigelloides</i>	6	Unknown	2
<i>Salm. enterica</i>	0	Unknown	0
Pooled	17	272	6

Research Component 3: Results

- MPN from 2009:

Species	Fecal Coliform MPN (Density/g feces)	Source
Alligator	9.0 x 10⁹	Current study
Duck	3.3 x 10 ⁷	Schueler and Holland, 2000
Human	1.3 x 10 ⁷	Schueler and Holland, 2000
Dog	2.3 x 10 ⁷	Schueler and Holland, 2000
Turtle	1.6 x 10 ⁶	Harwood <i>et al.</i> , 1999
Cow	2.3 x 10 ⁵	Schueler and Holland, 2000

Conclusions



Conclusions

- First study to identify poikilotherms as a source of bacterial pollution in coastal waters
- Alligator fecal coliforms denser than endotherms
- REP-PCR valid BST technique for indicator bacteria and potential pathogens



Conclusions

- Management practices:
 - Hunting
 - Tagging
 - Vegetated buffers



Alligators



Alligators live in many of South Carolina's coastal waters. They eat fish, turtles, and other small animals. Large alligators, however, occasionally catch larger animals such as raccoons, deer, dogs, etc.

- ▶ Alligators are protected under S.C. Regulation 123-151.
- ▶ It is unlawful to feed alligators; this includes throwing fish or fish parts into the water.
- ▶ Do not approach alligators no matter how big or small. "Gators" can move fast!
- ▶ Keep pets on a leash and away from the water.
- ▶ Do not allow children to throw toys or rocks into the water. To an alligator, a splash means food.



DNR

Conclusions

- Potential Collaborations:
 - Caimans: Universidade Federal de Mato Grosso do Sul, Campo Grande, Mato Grosso do Sul, Brazil



- GEL Engineering, Charleston, SC

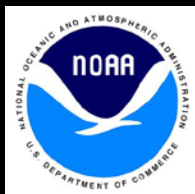
Conclusions

- Cold-blooded sources:
 - Fecal coliforms
 - Potential pathogens
- Concentration of sources
 - Loss of habitat
- Management practices to strive for human and animal balance



Acknowledgements

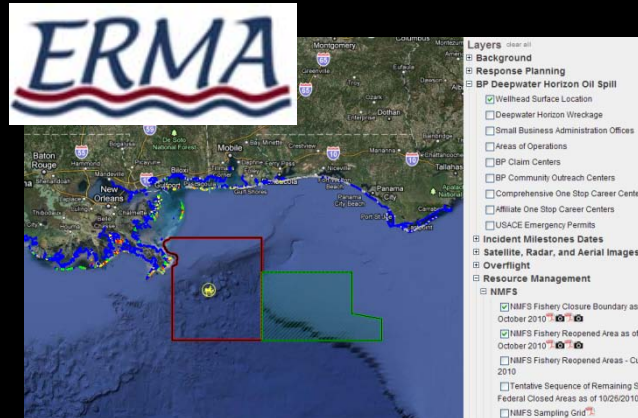
- This project is supported by NOAA Center for Coastal Environmental Health and Biomolecular Research (CCEHBR), USC, the NOAA/Sea Grant Research Fellowship, USC Arnold Fellowship, Slocum-Lunz Foundation Grant, and a South Carolina Wildlife Federation Scholarship
- **Committee:**
 - Dr. Geoff Scott, NOAA CCHBR-USC
 - Dr. Sean Norman, USC
 - Dr. Dwayne Porter, USC
 - Dr. Heath Kelsey, NOAA-UMCES
- Molecular Microbial Ecology Lab, GIP Lab, NOAA/CCEHBR, Palmetto Bluff Conservancy, SCDNR, Cypress Gardens, and Spring Island - Low Country Institute



Office of National Marine Sanctuaries



Blue Ocean Film Festival, MBNMS



Deepwater Horizon Oil Spill Response



IEEE Conference, Buenos Aires, Argentina



Montana LIVE Dive, TBNMS



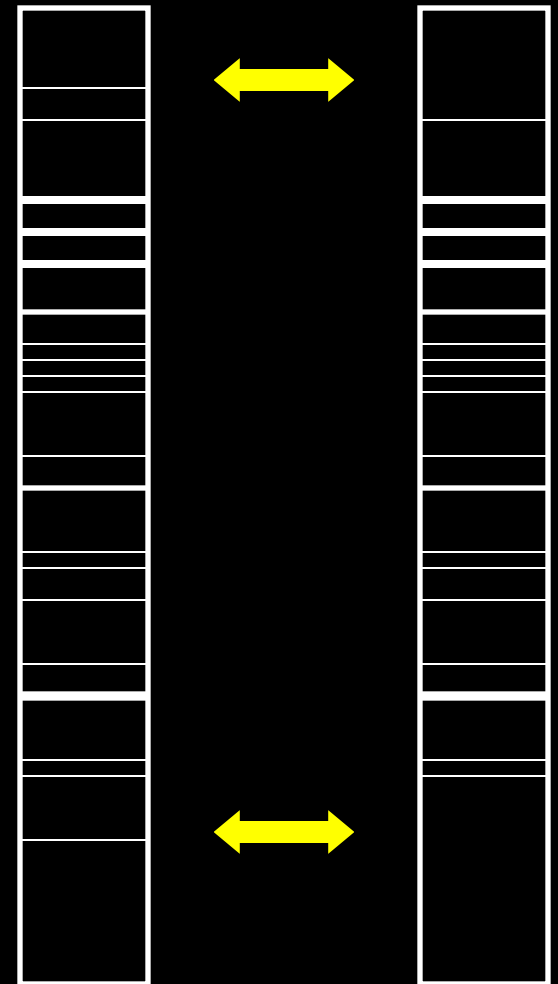
Aquarius 2010 Mission, FKNMS

Later Alligator



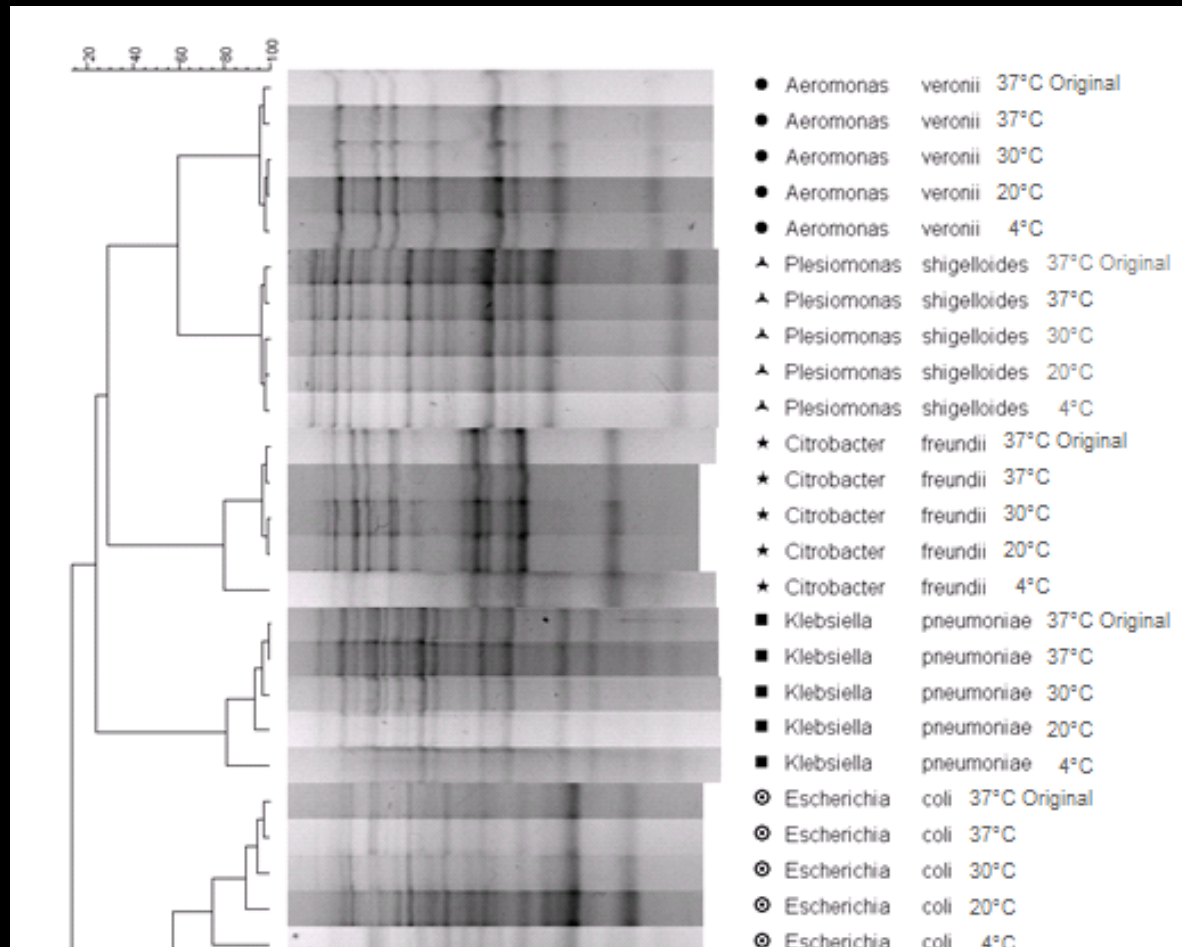
Research Component 2: Results

- 90% similarity
 - Repeated control runs
 - 2 band difference
- $\geq 90\%$ similarity cutoff value
 - Matching bands from same source

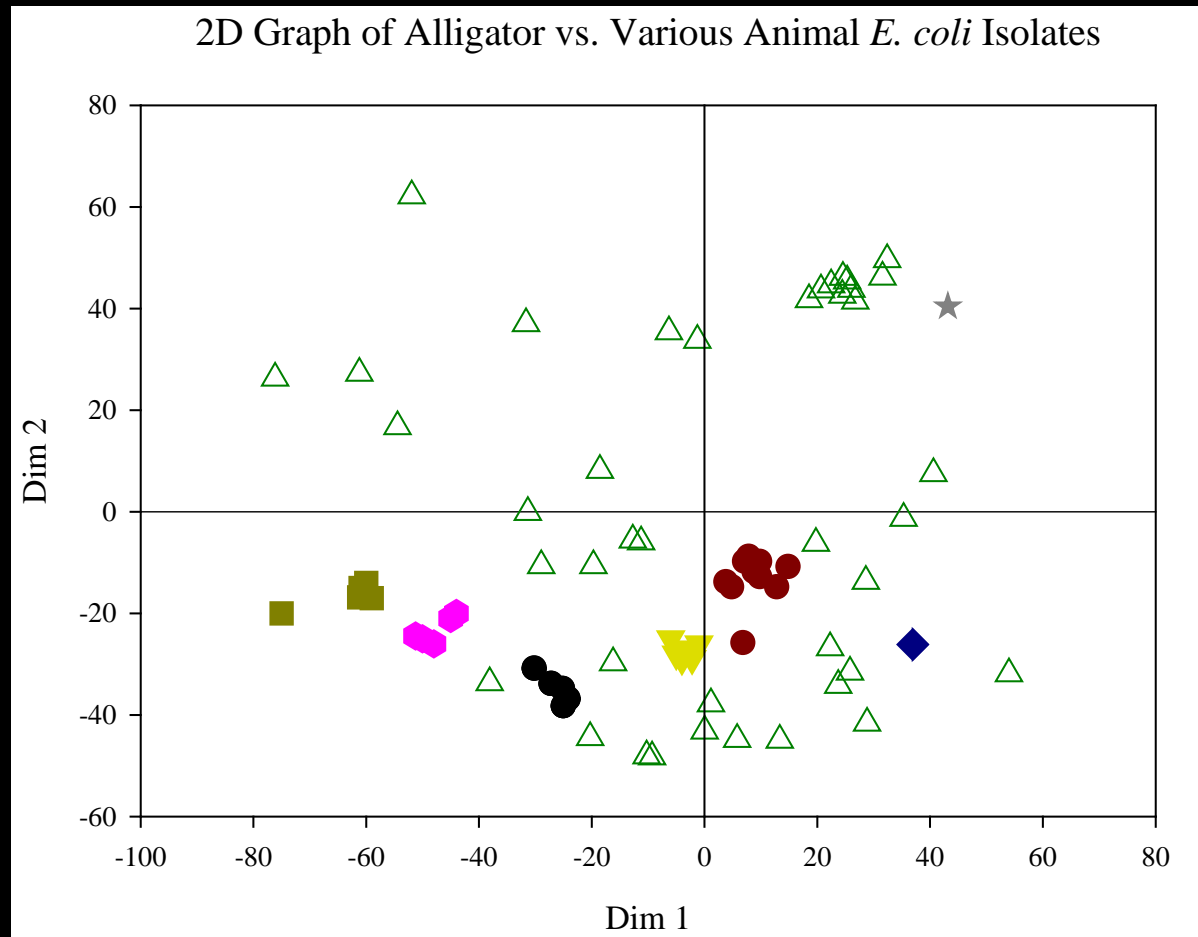


Research Component 2: Results

- Fingerprint stability



Research Component 2: Results



E. coli isolates represent alligator ▲ , chicken ▼ , cow ○ , dog ■ , horse ● , pig ◆ , raccoon ★ , and sea turtle ◆ .

Results

