

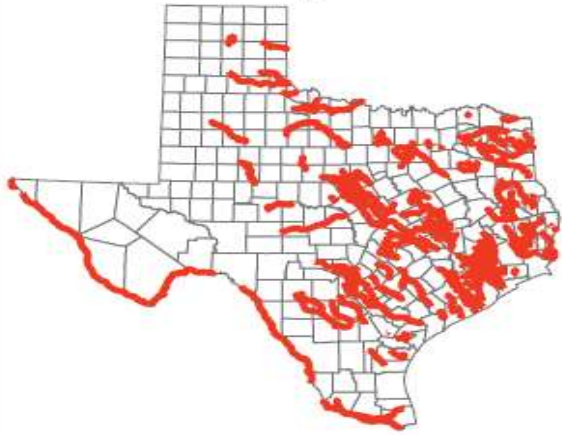
Using Bacterial Source Tracking to Develop Watershed Restoration Plans

**Kevin Wagner, George DiGiovanni,
Terry Gentry, Elizabeth Casarez**

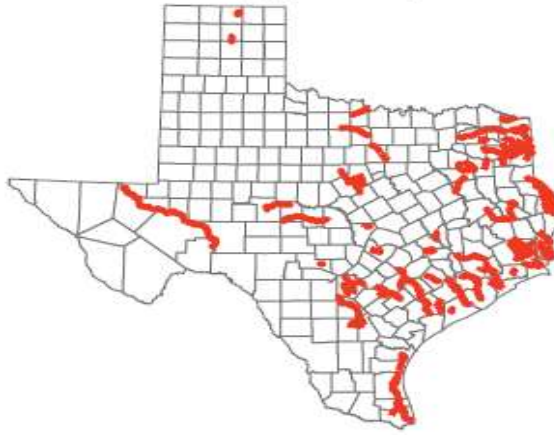
Bacteria/Pathogens

The #1 Cause of Water Quality Impairment in Texas

Bacteria Impairment



Dissolved Oxygen Impairment

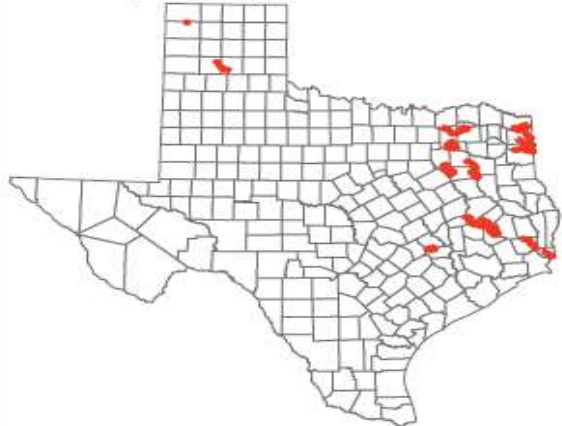


Toxicity Impairment

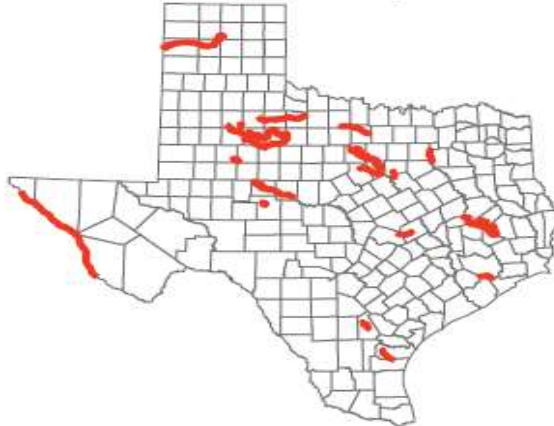


WATER QUALITY IMPAIRMENTS IN TEXAS

pH Impairment



Dissolved Solids Impairment Nitrate and Nitrite Impairment

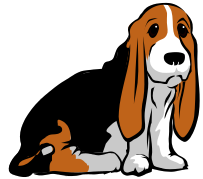
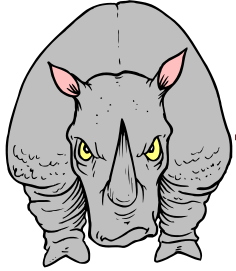


Where did the Bacteria (*E. coli*) Come From?

- **Potential sources**
 - Humans
 - Domesticated animals
 - Wildlife
 - ~140 mammals
 - ~650 birds
- **Methods for determining sources**
 - Source survey
 - Modeling
 - Bacterial source tracking (BST)



PREMISE BEHIND BST

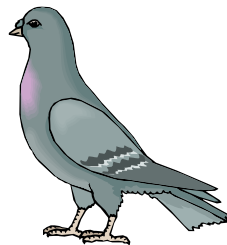
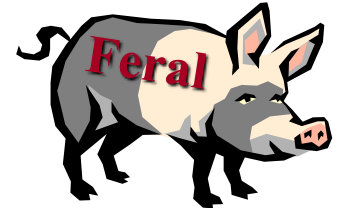


Different guts → Different adaptations

→ Different *E. coli* strains →

Genetic Differences

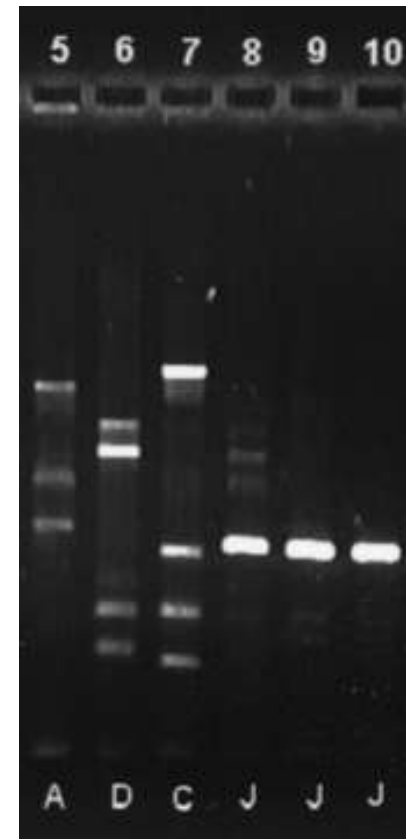
Phenotypic Differences



History of BST in Texas

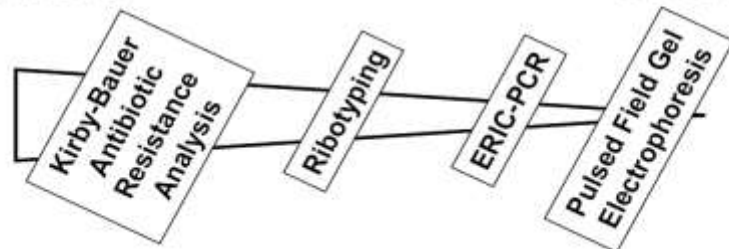


- Lake Waco/Belton Project (2002-2006)
 - Evaluated utility & methods
 - Recommended 2-method composite
 - Enterobacterial repetitive intergenic consensus sequence-polymerase chain reaction (ERIC-PCR)
 - RiboPrinting® (RP)
- TMDL Task Force Report – 2007
 - Confirmed ERIC-RP as recommended method
- Required BST Library Development



Lowest
Resolution

Highest
Resolution



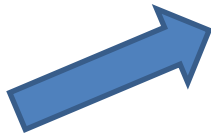
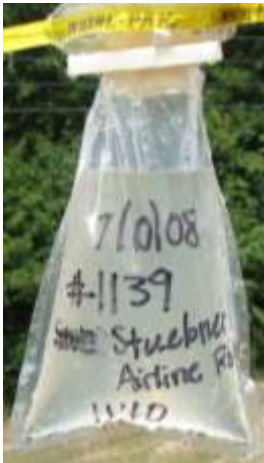
Methods

Known Sources

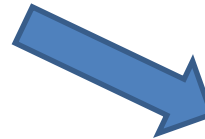
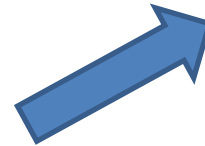


Isolate
E. coli

Unknown Source

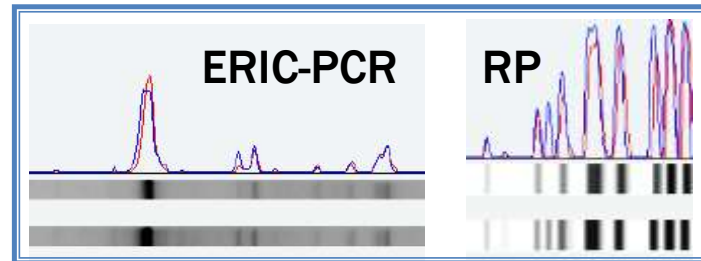


DNA
Fingerprint



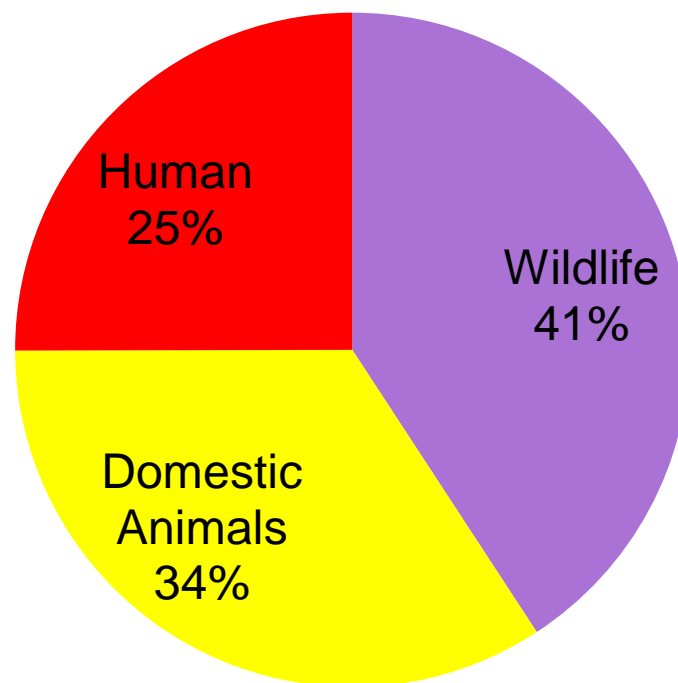
Add to
Library

Compare
to Library
for Source
ID

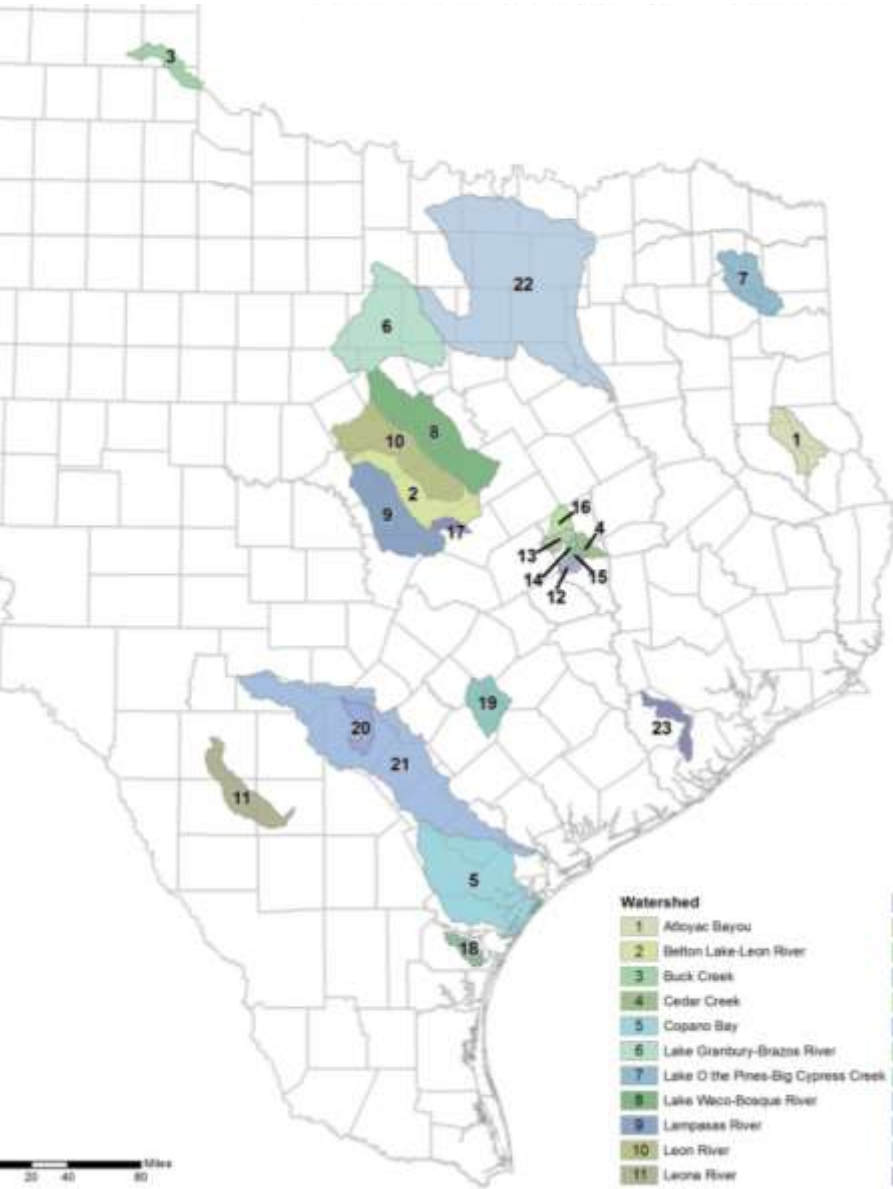


Texas *E. coli* BST Library

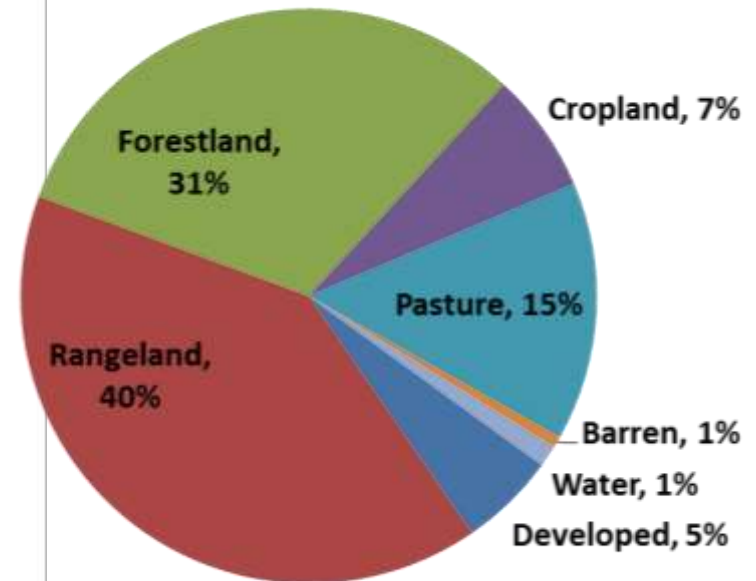
- **Contains**
 - **1,669 *E. coli* isolates**
 - **From 1,455 different fecal samples**
 - **Representing >50 animal subclasses**
 - **Collected from 13 watersheds (& growing) across Texas**



Texas BST Studies To Date

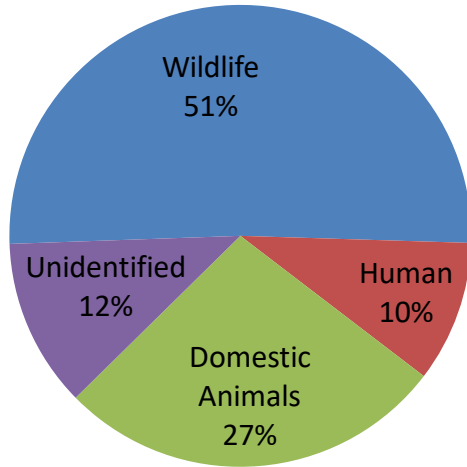


Typical Landuse in 11 BST Watersheds



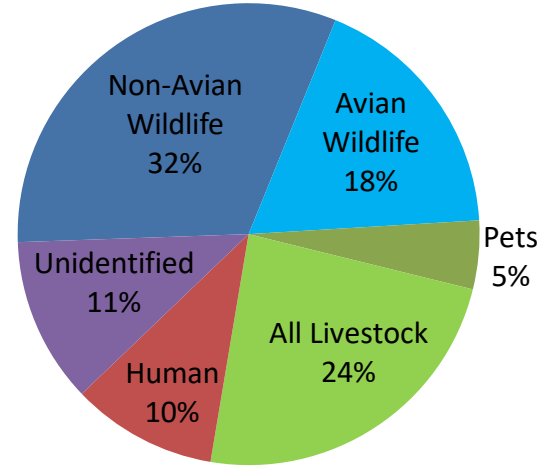
3-Way Split

(averages based on findings in 11 watersheds)



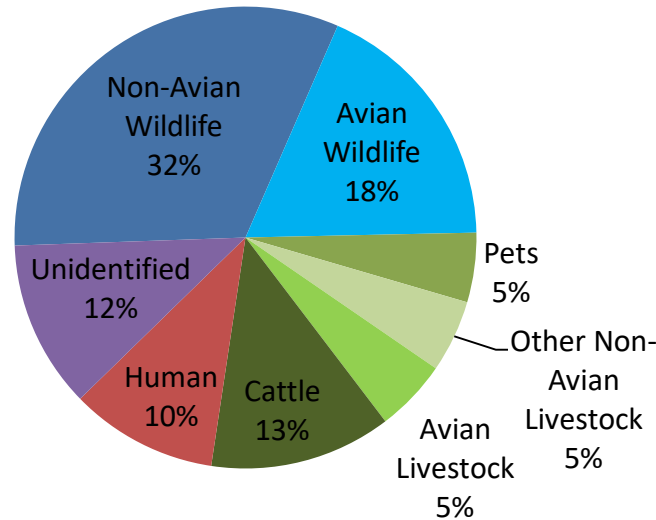
5-Way Split

(averages based on findings in 10 watersheds)



7-Way Split

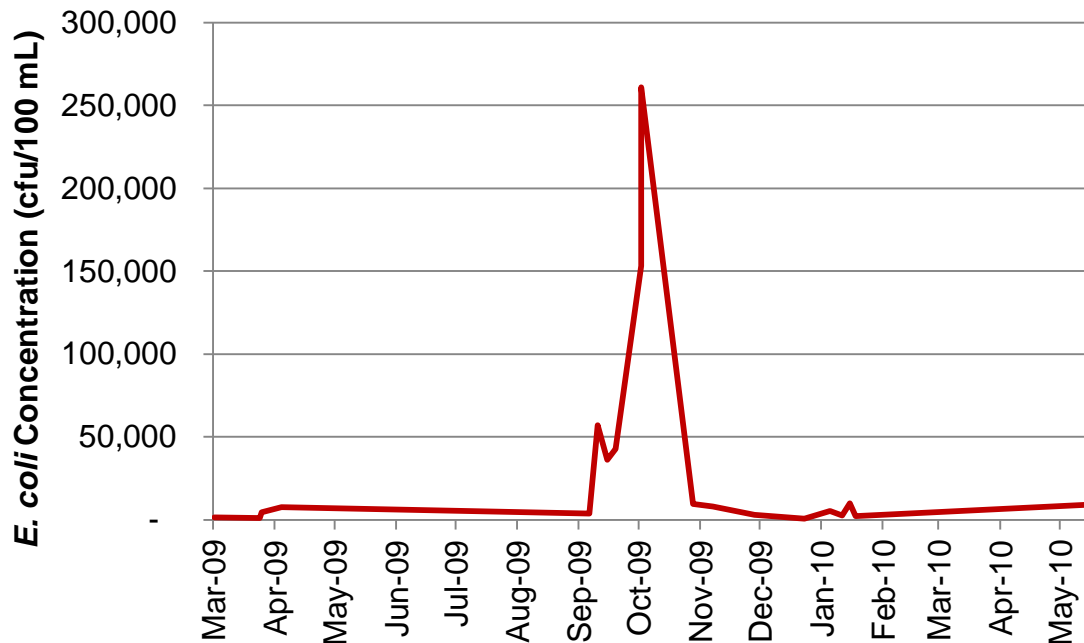
(averages based on findings in 7 watersheds)



Impacts of wildlife on *E. coli* runoff

Site	Fecal Coliform (#/100 mL)	<i>E. coli</i> (cfu/100 mL)	Reference
Ungrazed pasture	10,000		Robbins et al. 1972
Ungrazed pasture	6,600		Doran et al. 1981
Control plots		6,800	Guzman et al. 2010
Pasture destocked >2 mos.		1,000-10,000	Collins et al. 2005
Ungrazed pasture		6,200-11,000	Wagner et al. 2012
Pasture destocked >2 wks.		2,200-6,000	Wagner et al. 2012

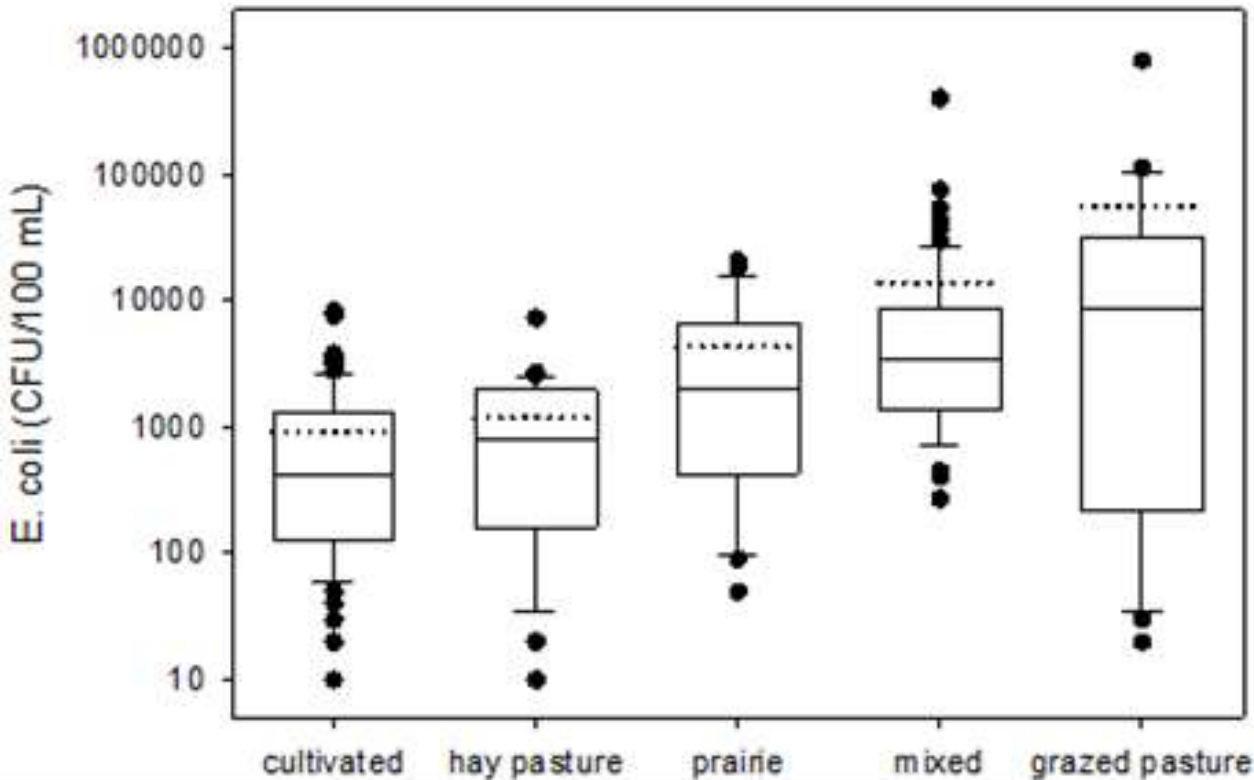
***E. coli* concentrations at ungrazed site BB1 (2009-2010)**



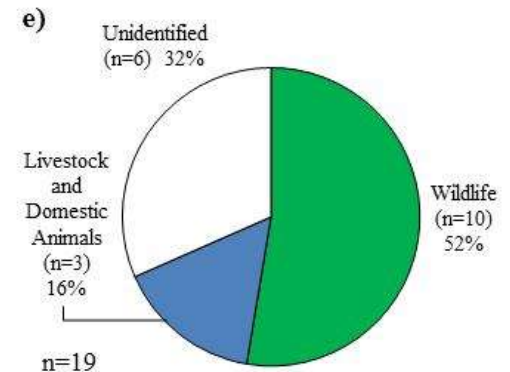
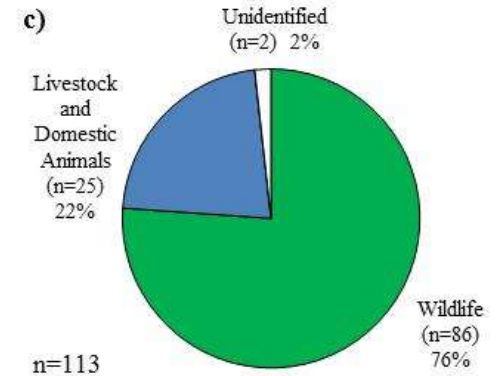
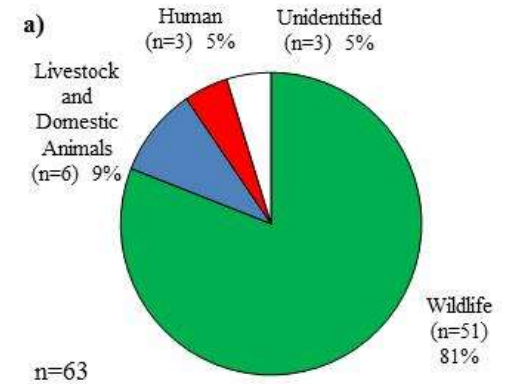
**Wildlife contributed
>80% of *E. coli*
loading at grazed
sites in 2009**

Increasing *E. coli* with increasing wildlife habitat

Edge-of-field runoff *E. coli* concentrations (Harmel)



Soil *E. coli* sources (Gregory)



Summary & Implications of BST Findings

Summary:

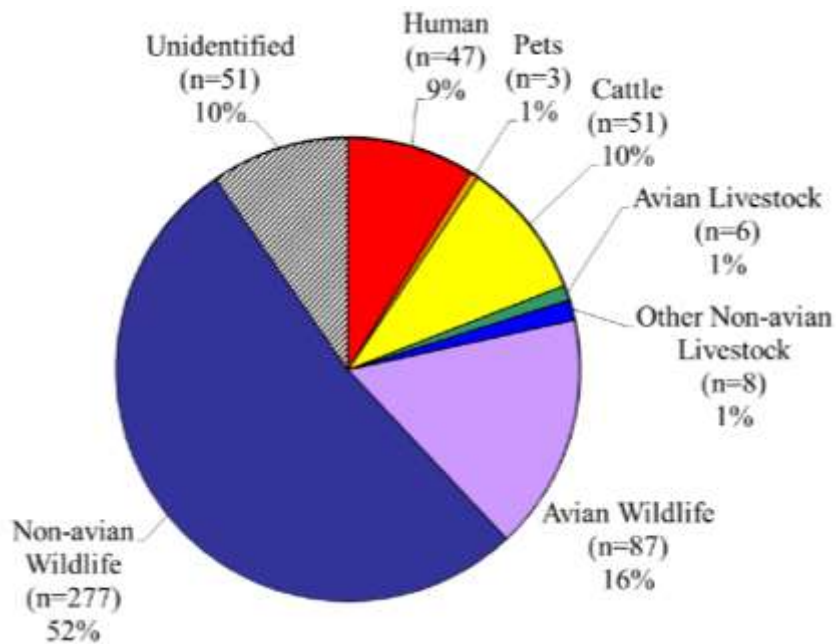
- BST performing well (100% 3-way RCC; 92% 7-way RCC)
 - Proving to be useful tool for identifying significant bacteria sources
- Wildlife = source of 50% of isolates in predominately rural watersheds
 - Edge of field monitoring confirms significance of background sources

Implications:

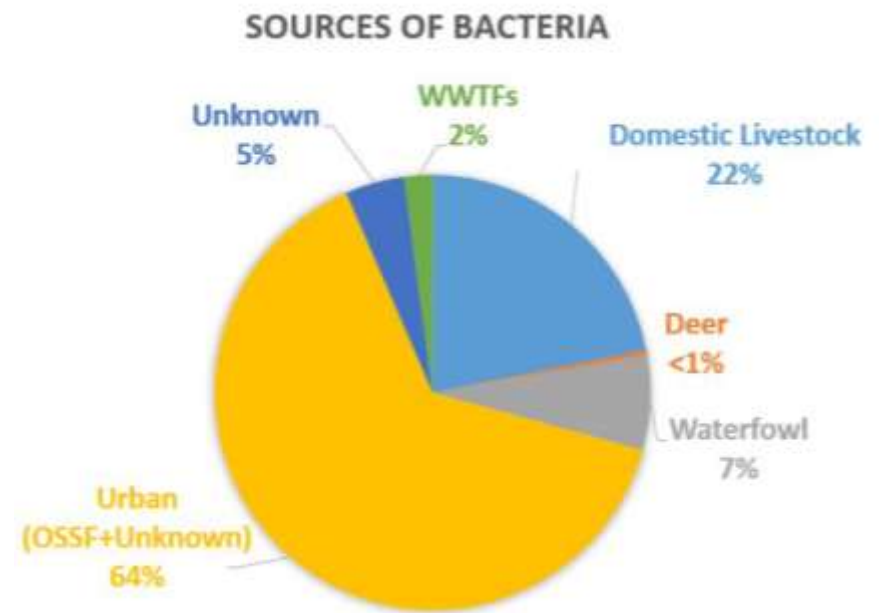
- Background/wildlife loadings need to be considered when:
 - Applying water quality standards
 - Developing tmdls and watershed based plans
- Ignoring background concentrations may lead to:
 - Nonattainment of water quality standards
 - Inaccurate load allocations and reductions

Integrating Modeling & BST: Arroyo Colorado Case Study

BST Results

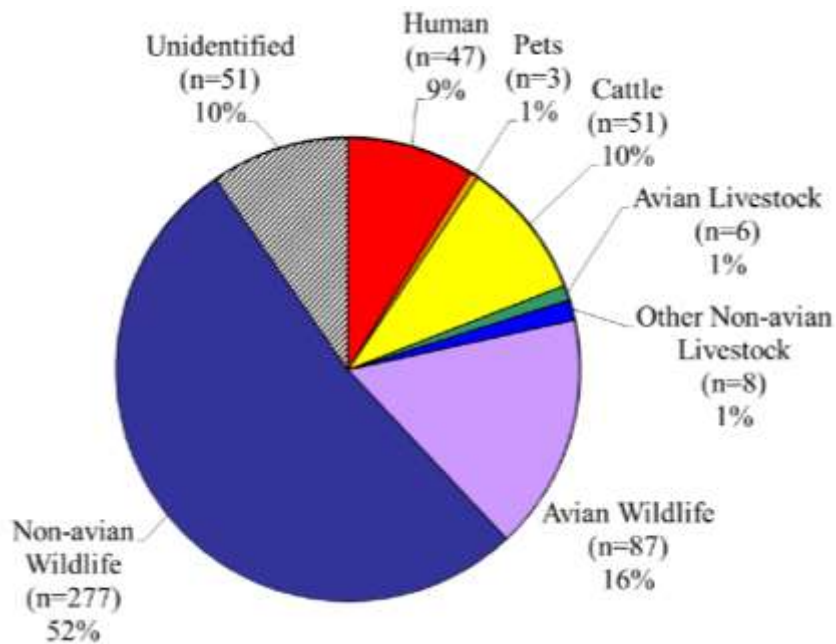


Initial SWAT Model Results

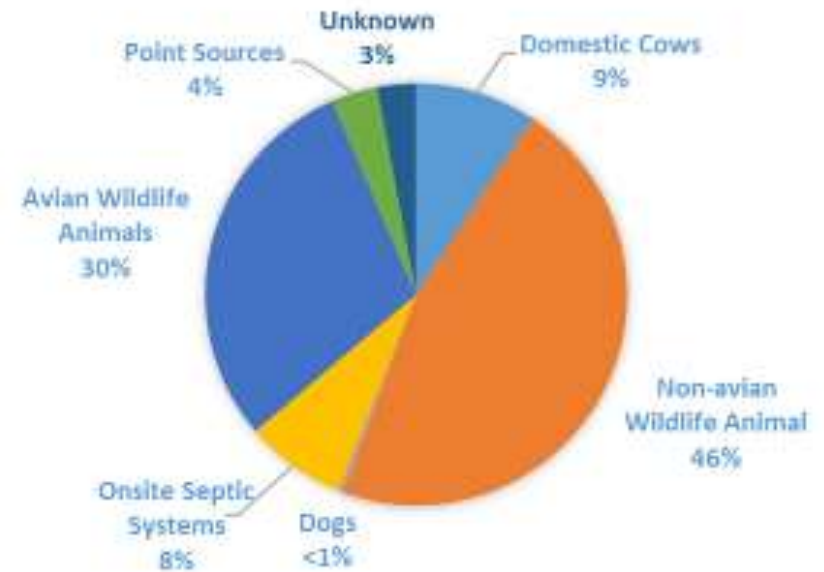


Calibrated/validated SWAT with BST

BST Results



Final SWAT Model Results

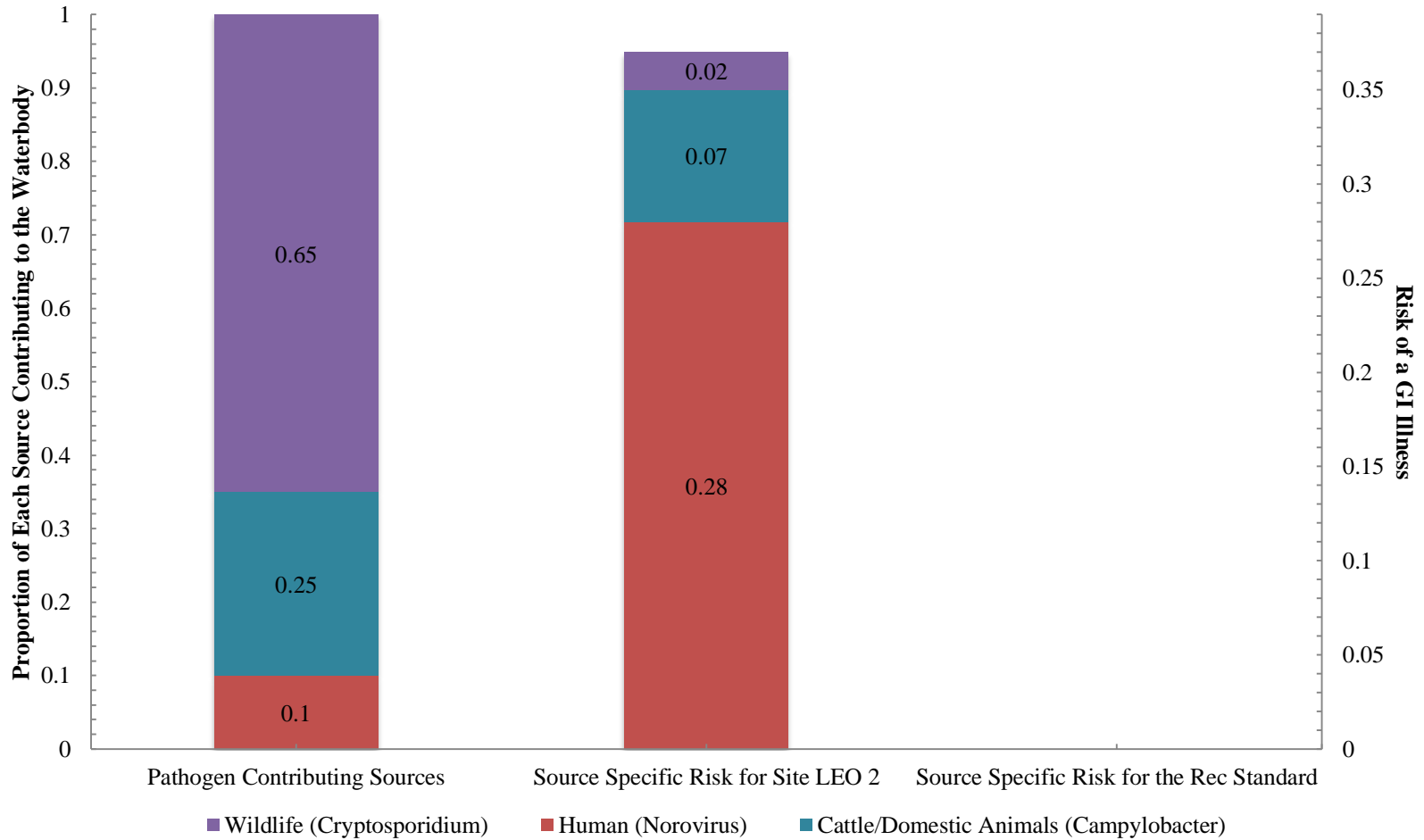


Future uses of BST:

Quantitative Microbial Risk Assessment

- EPA 2012 recreational water quality criteria provided tools for developing site-specific criteria:
 - epidemiological studies
 - quantitative microbial risk assessment
 - use of alternative indicators or methods

Walnut Creek QMRA Case Study: Risk of GI Illness ≠ BST Percentages



QMRA Findings & Implications

- Human and non-human fecal sources have different potential risks for a GI illness
 - Proportion of a single source contributing to the overall *E.coli* concentration not an indicator of overall human health risk
- Risk driven by human source
- Management toward reducing human sources
 - Compliance & maintenance of WWTPs, sanitary sewer systems, wastewater collection systems & infrastructure

Questions?

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