

# Dairy Manure-Borne Pathogens in the Environment and the Human Health Risk

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# Manure's Double-Edged Sword

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## Manure as Asset



Manure field-application is a cost-effective and sustainable approach for optimal soil tilth and fertility

## Manure as Liability



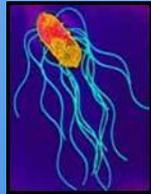
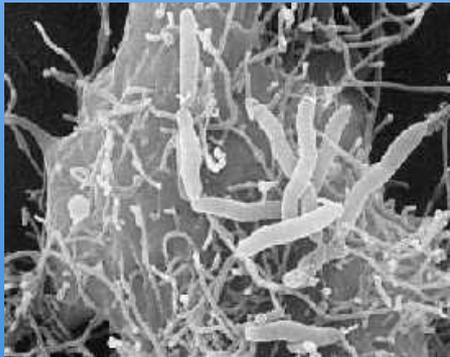
Manure may contain pathogens harmful to both humans and livestock

Societal goal: Maximize the beneficial uses of manure while minimizing environmental pathogen transmission

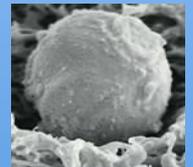
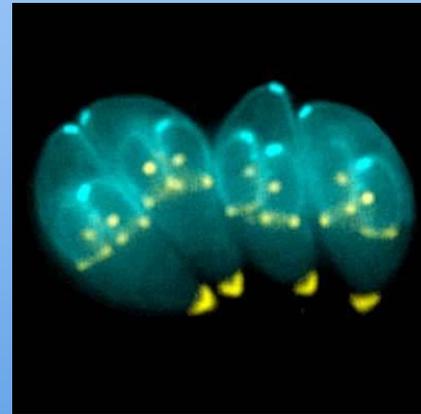
# Pathogens Present in Dairy Manure

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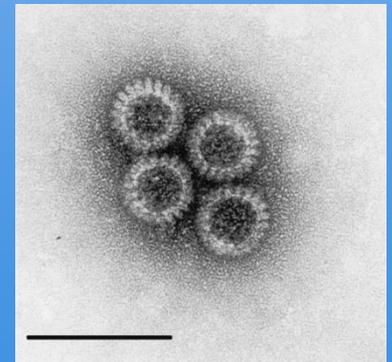
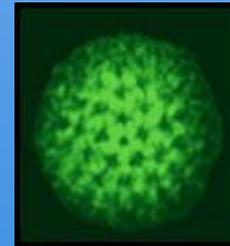
Bacteria (e.g., *Campylobacter*,  
*Salmonella*, *E. coli* O157:H7,  
*Listeria monocytogenes*)



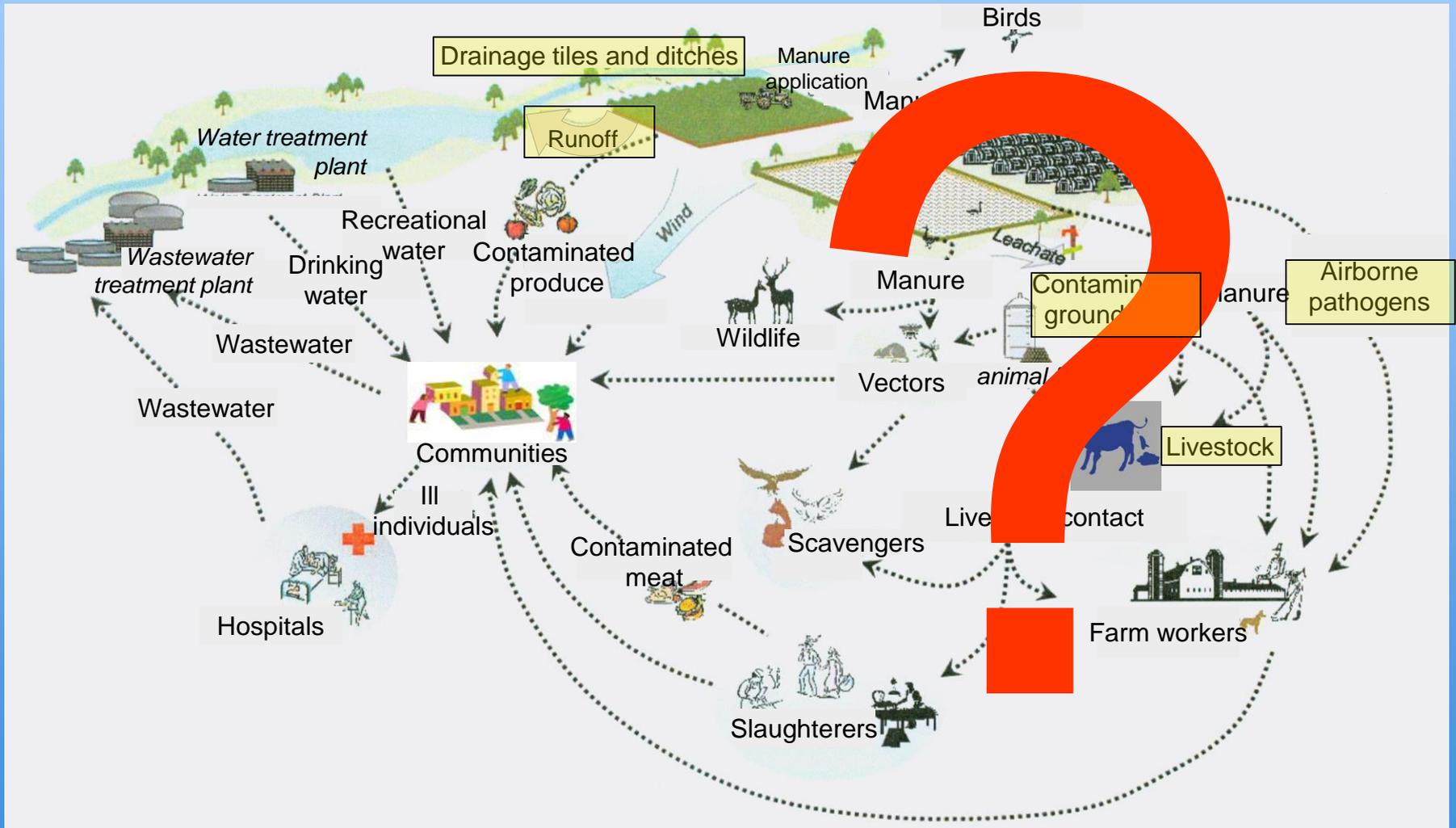
Protozoa (e.g.,  
*Cryptosporidium*,  
*Giardia*, *Toxoplasma*)



Viruses (e.g.,  
rotavirus)



# Human and Livestock Pathogen Movement in the Environment



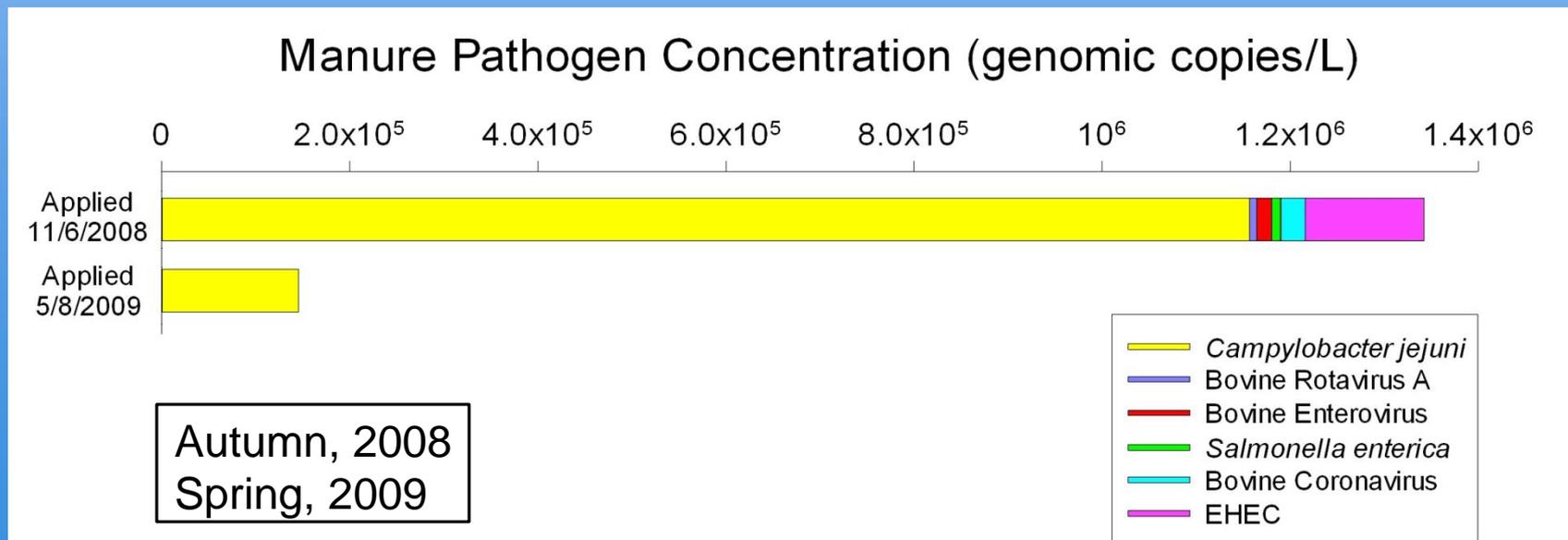
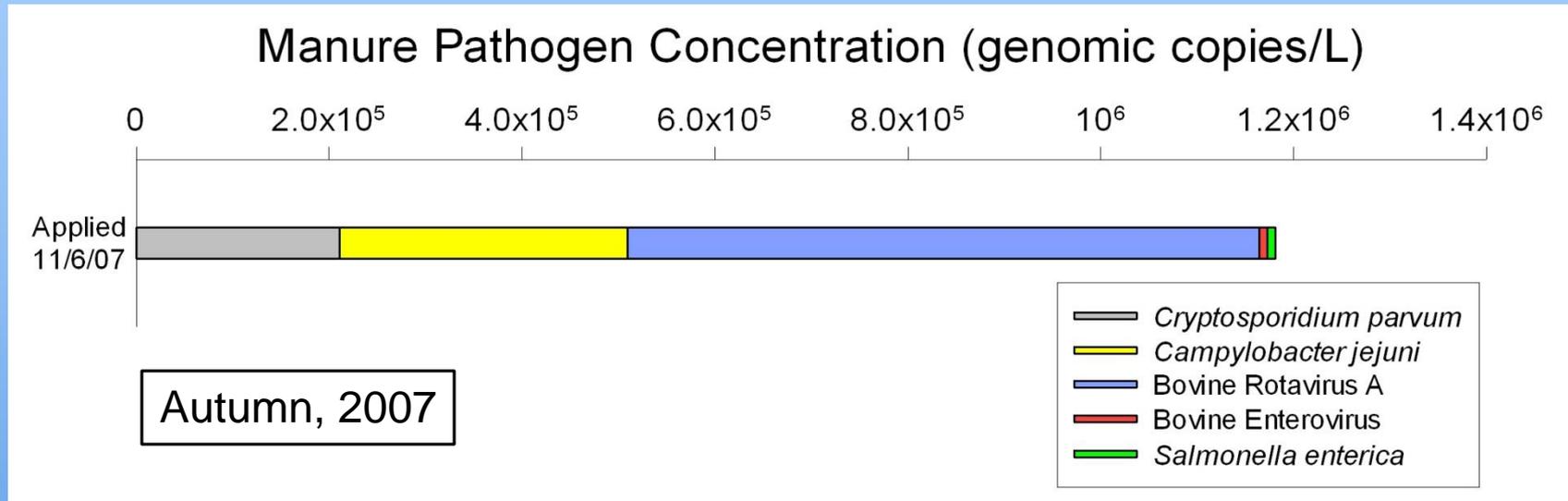
# Estimates of Enteric Illness Attributable to Contact with Animals and Their Environments in the United States

- 445,213 /3.2 million illnesses (14%) from animal contact

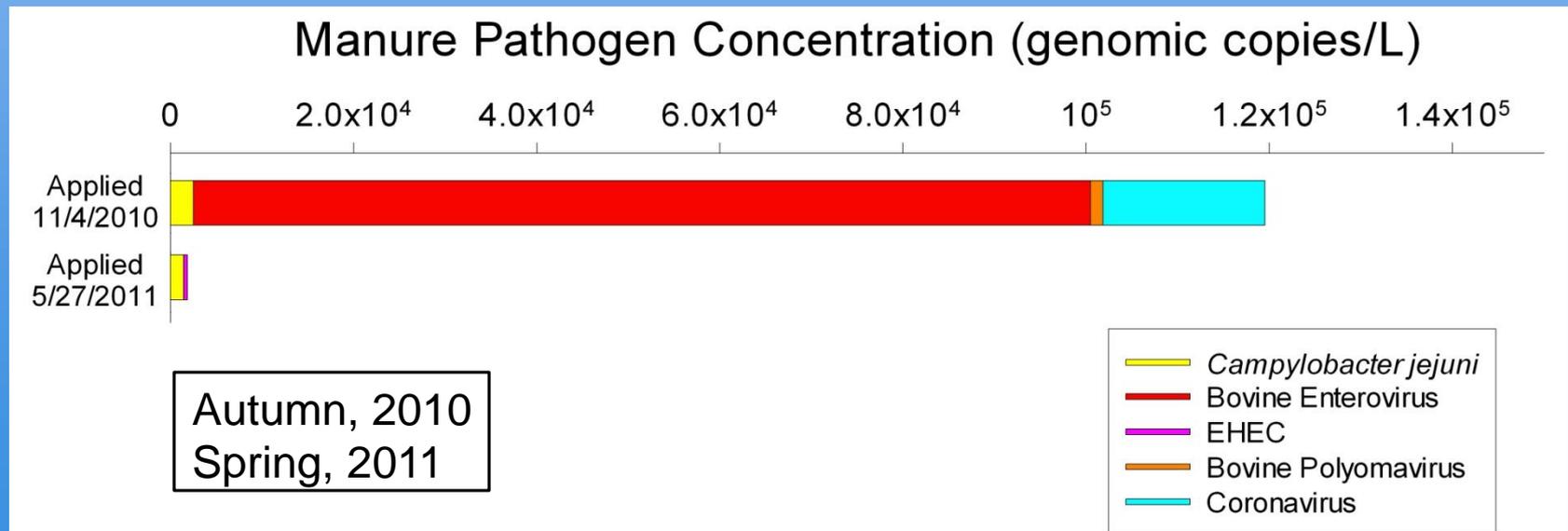
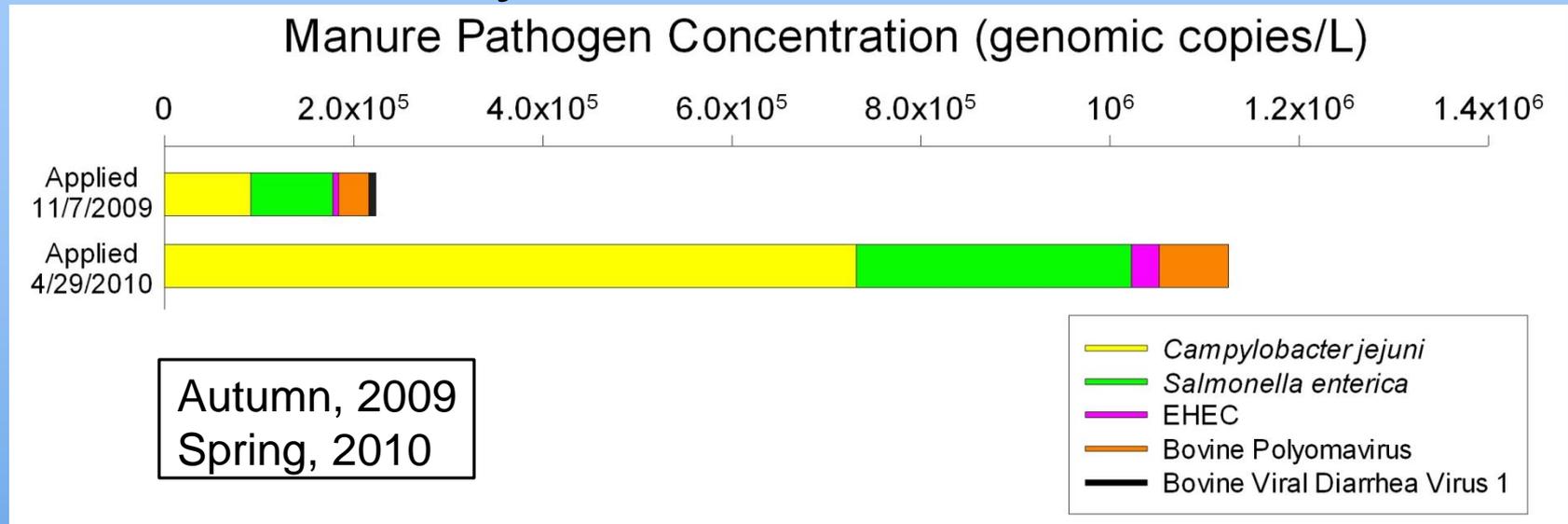
Organism	% from animal contact	Annual # illnesses	Annual # hospitalizations	Annual # deaths
Campylobacter species	17%	187,481	1,877	17
ST E. coli	14	16,057	230	2
Non-typhoid Salmonella	11	127,155	2,392	47
Cryptosporidium species	16	113,344	412	7

Centers for Disease Control and Prevention: Hale et al. 2012, CID, 54:S472-79.

# Pathogens in manure from a single farm by year and season



# Pathogens in manure from a single farm by year and season



# Brown Water Events in Northeast Wisconsin



- Groundwater recharge, especially spring snow melt, can generate brown water events
- Several outbreaks associated with these events e.g., EHEC, *Campylobacter jejuni*
- This well is code compliant, 123 ft deep, cased to 63 ft

*Photos courtesy of Chuck Wagner*

# Kewaunee County Cattle

- All cattle & calves in 2016 = 97,000
- Milk cows in 2013 = 45,500
- Milk cow herds in 2016 = 167
- Concentrated Animal Feeding Operations (CAFOs) 15 dairy, one beef
- Approximately 700 million gallons cattle manure per year



# Kewaunee County Septic Systems

- 4822 septic systems in the county
- 540 holding tanks, 155 abandoned

*Personal comm. Lee Luft, Kewaunee County Supervisor, March 7, 2017*

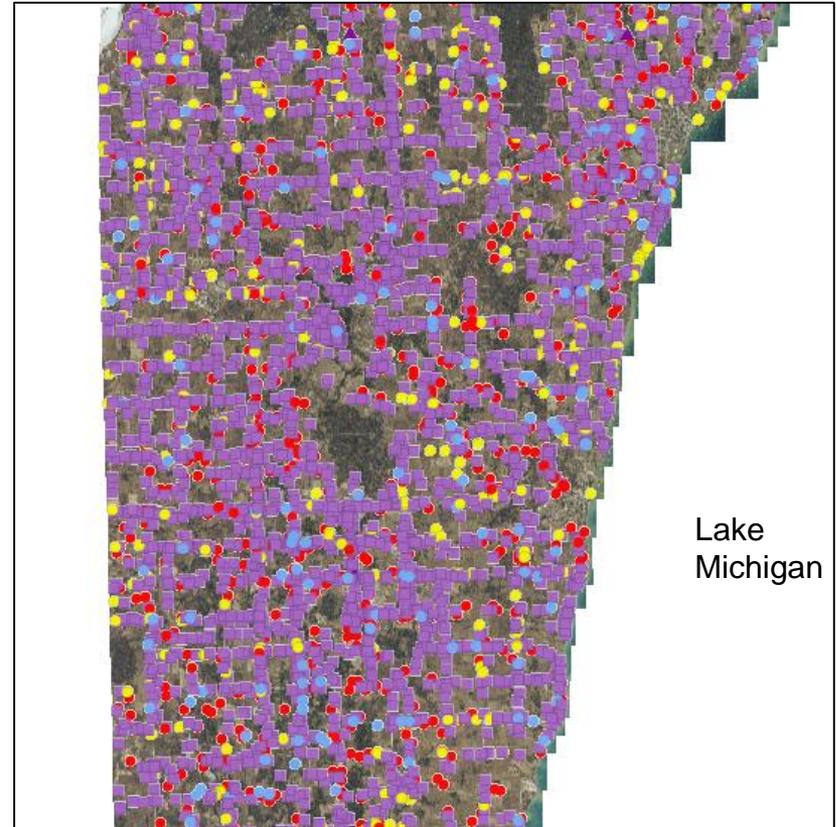
## Legend

Purple = replaced or inspected

Red = not inspected

Yellow = holding tank

Blue = abandoned system



Kewaunee County septic systems

Approximately 200 million gallons septic effluent per year released to the subsurface

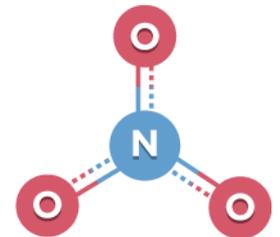
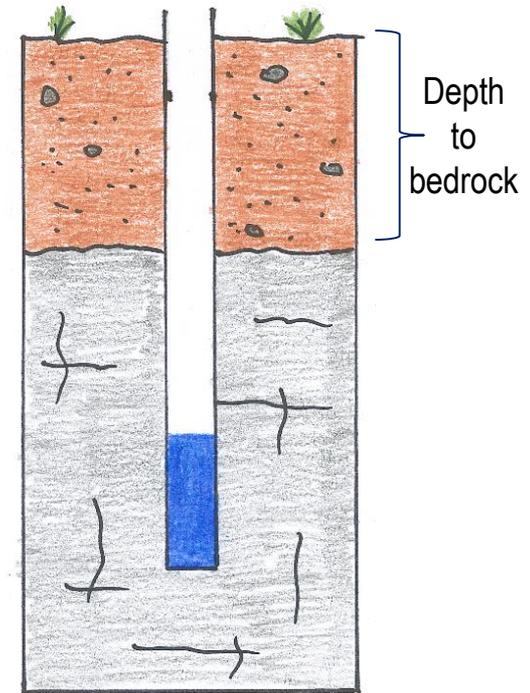
# Research Objectives

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1. Estimate county-wide contamination rate for nitrate and indicator bacteria as related to depth-to-bedrock
2. Determine source of fecal contamination using viruses and fecal markers
3. Identify risk factors for private well contamination using statistical models

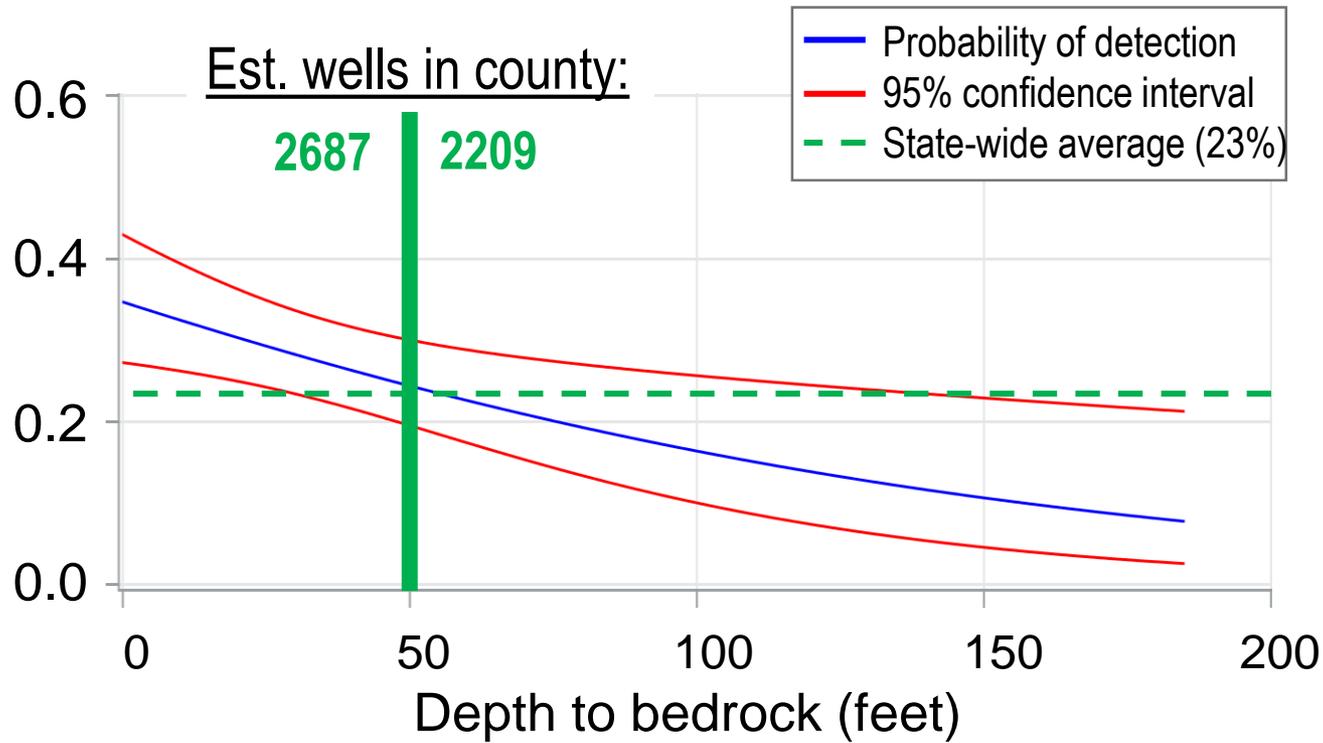
# Objective I: Total Coliform, *E. coli*, Nitrate

- County-wide randomized sampling of private wells – 4,896 in county
- Stratified by depth-to-bedrock: 0-5 ft, 5-20 ft, 20+ ft
- Participation rate ~ 50%
- Several day “Synoptic” sampling
- Recharge
  - November 2015
  - 317 wells in analysis
- No recharge
  - July 2016
  - 400 wells in analysis



# Effect of depth to bedrock on total coliform contamination

Probability of total coliform detection



**Fall, 2015**

n = 315

p = 0.009

# Objective 2: Determine fecal source

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- Randomized stratified sampling from 208 wells positive for total coliform, *E. coli*, or high nitrate ( $\text{N-NO}_3^- > 10 \text{ ppm}$ )
- Five sampling rounds:
  - April, August, November, 2016
  - January and March, 2017

# Study Sampling and Analyses

- Collected 138 samples from 131 household wells in Kewaunee County
- Pump ~800 L through hemodialysis filters
- qPCR for microbial genetic targets
  - Human-specific microbes
  - Bovine-specific microbes
  - Non-specific microbes (pathogens of both people and cattle)



# Microbes: Identifying the Fecal Source

(n = 138 samples from 131 wells) (red font indicates pathogenic)

Host	Microorganism	Wells	Concentration (gene copies/L)
	<b>Adenovirus A</b>	1	1
	<i>Bacteroidales</i> -like Hum M2	7	< 1 – 1050
Human- specific	Human <i>Bacteroides</i>	27	< 1 – 34
	<b><i>Cryptosporidium hominis</i></b>	1	qualitative
	<b>Rotavirus A</b> (G1 P[8])	7	qualitative
	<b>All</b>	<b>33</b>	

Not detected: [human-specific] adenovirus B & C, D, F, enterovirus, human polyomavirus, norovirus GI & GII  
[bovine-specific] coronavirus, bovine diarrheal virus 1 & 2

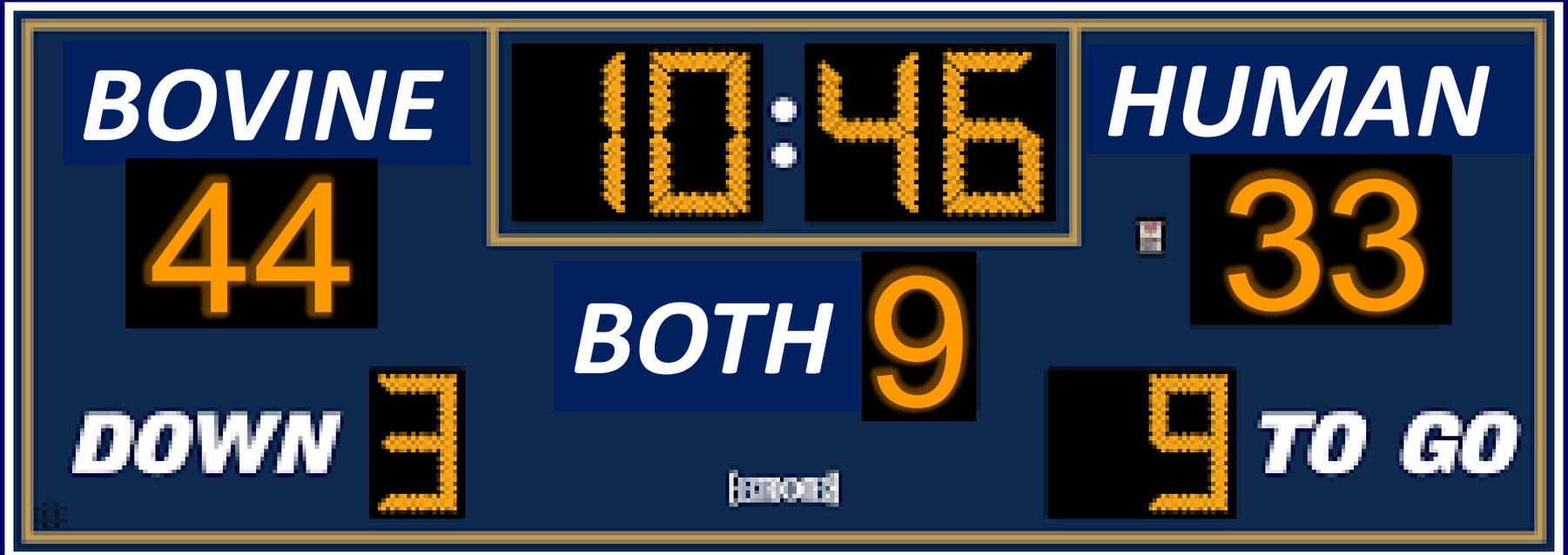
# Microbes: Identifying the Fecal Source

(n = 138 samples from 131 wells) (red font indicates pathogenic)

Host	Microorganism	Wells	Concentration (gene copies/L)
Bovine-specific	<i>Bacteroidales</i> -like Cow M2	2	29 - 915
	<i>Bacteroidales</i> -like Cow M3	4	3 - 49,818
	Bovine <i>Bacteroides</i>	36	< 1 - 42,398
	<b>Bovine polyomavirus</b>	8	< 1 - 451
	<b>Bovine enterovirus</b>	1	2
	<b>Rotavirus A</b> (G10 P[11])	12	qualitative
	<b>All</b>	<b>44</b>	

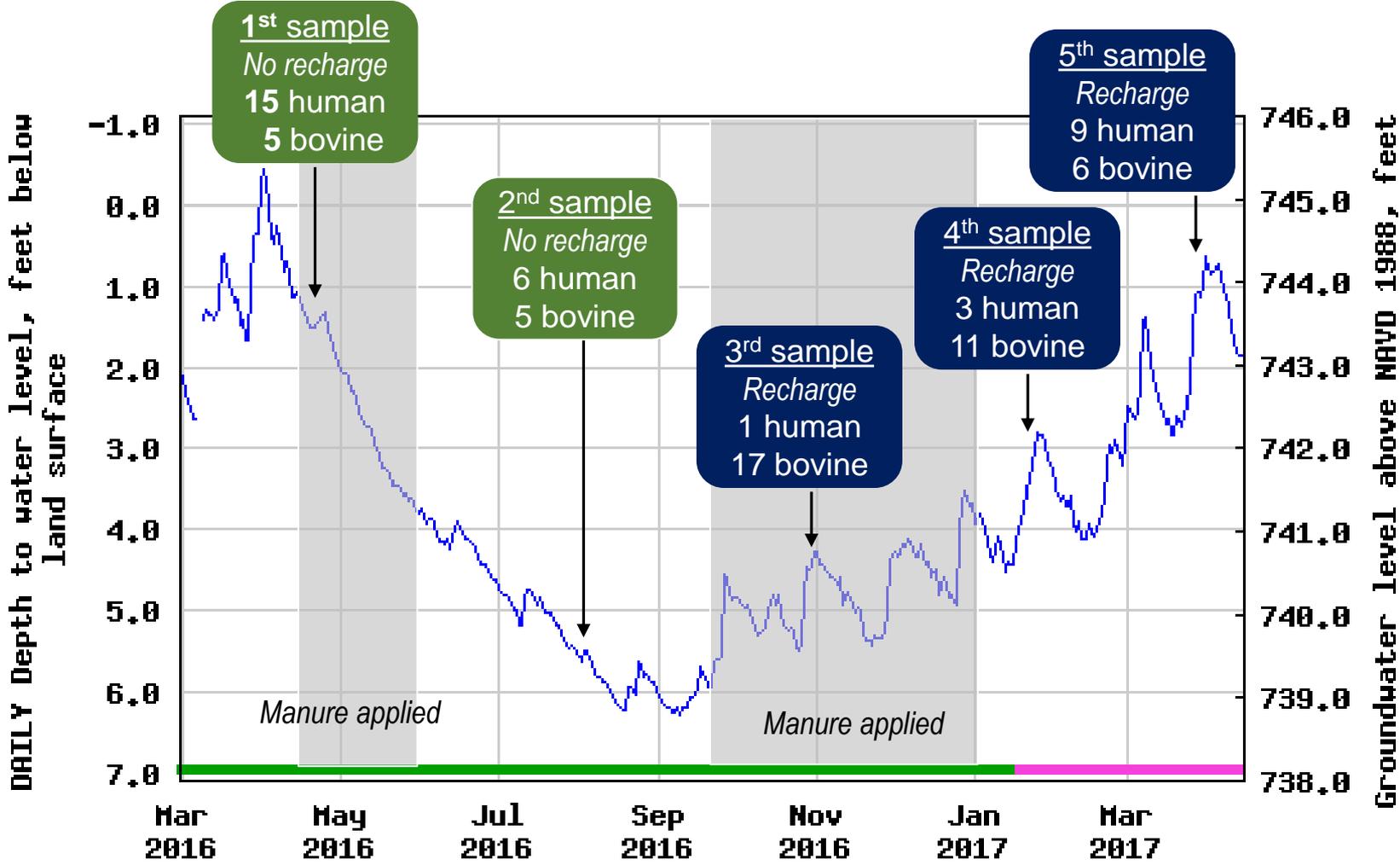
Not detected: [\[human-specific\]](#) adenovirus B & C, D, F, enterovirus, human polyomavirus, norovirus GI & GII  
[\[bovine-specific\]](#) coronavirus, bovine diarrheal virus 1 & 2

# Well Contamination Scoreboard



Host	Microorganism	Wells	Concentration (gene copies/L)
	<i>Campylobacter jejuni</i>	1	< 1
	<i>Cryptosporidium parvum</i>	13	qualitative
	<i>Cryptosporidium</i> spp.	16	< 1 – 3
	<i>Giardia lamblia</i>	2	< 1
	Pathogenic <i>E. coli</i> ( <i>eae</i> gene)	1	4
	Pathogenic <i>E. coli</i> ( <i>stx1</i> gene)	1	16
Non-specific	Pathogenic <i>E. coli</i> ( <i>stx2</i> gene)	1	1
	Pepper mild mottle virus	13	2 - 3811
	Rotavirus A ( <i>NSP3</i> gene)	17	< 1 – 4481
	Rotavirus A ( <i>VP7</i> gene)	7	< 1 – 732
	Rotavirus C	3	45 – 1301
	<i>Salmonella</i> ( <i>invA</i> gene)	3	< 1 – 13
	<i>Salmonella</i> ( <i>ttr</i> gene)	5	5 – 59
	<b>All</b>	<b>44</b>	
	<b>Total positive wells</b>	<b>79</b>	<b>&lt; 1 - 49818</b>

# Groundwater levels during sampling for pathogens & fecal indicators



— Daily median depth to water level    — Period of provisional data  
 — Period of approved data

No recharge    Recharge    Manure Applied

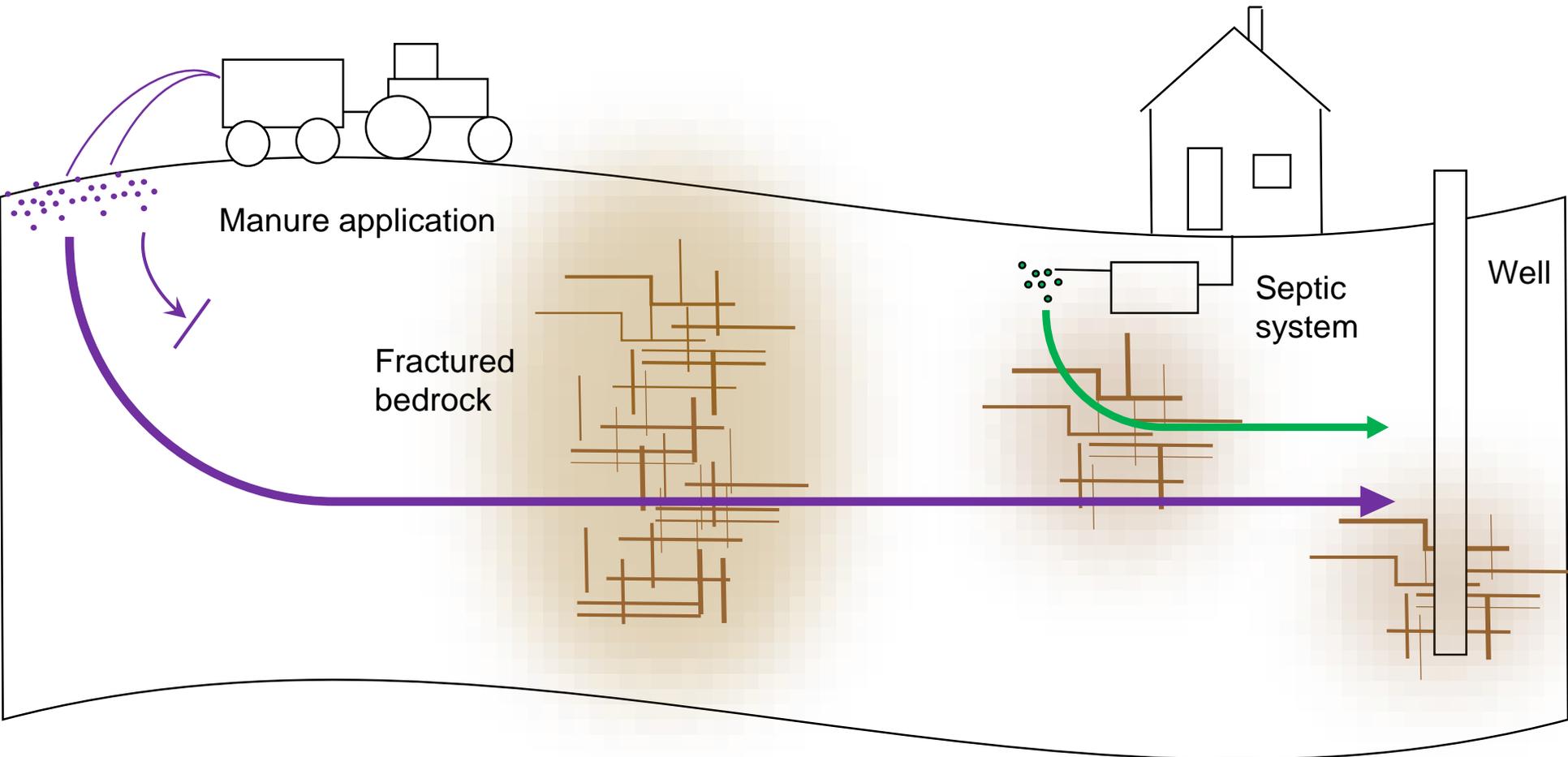
# Conceptual Model of Fecal Contamination in Kewaunee County - 1

## Bovine pathogen source

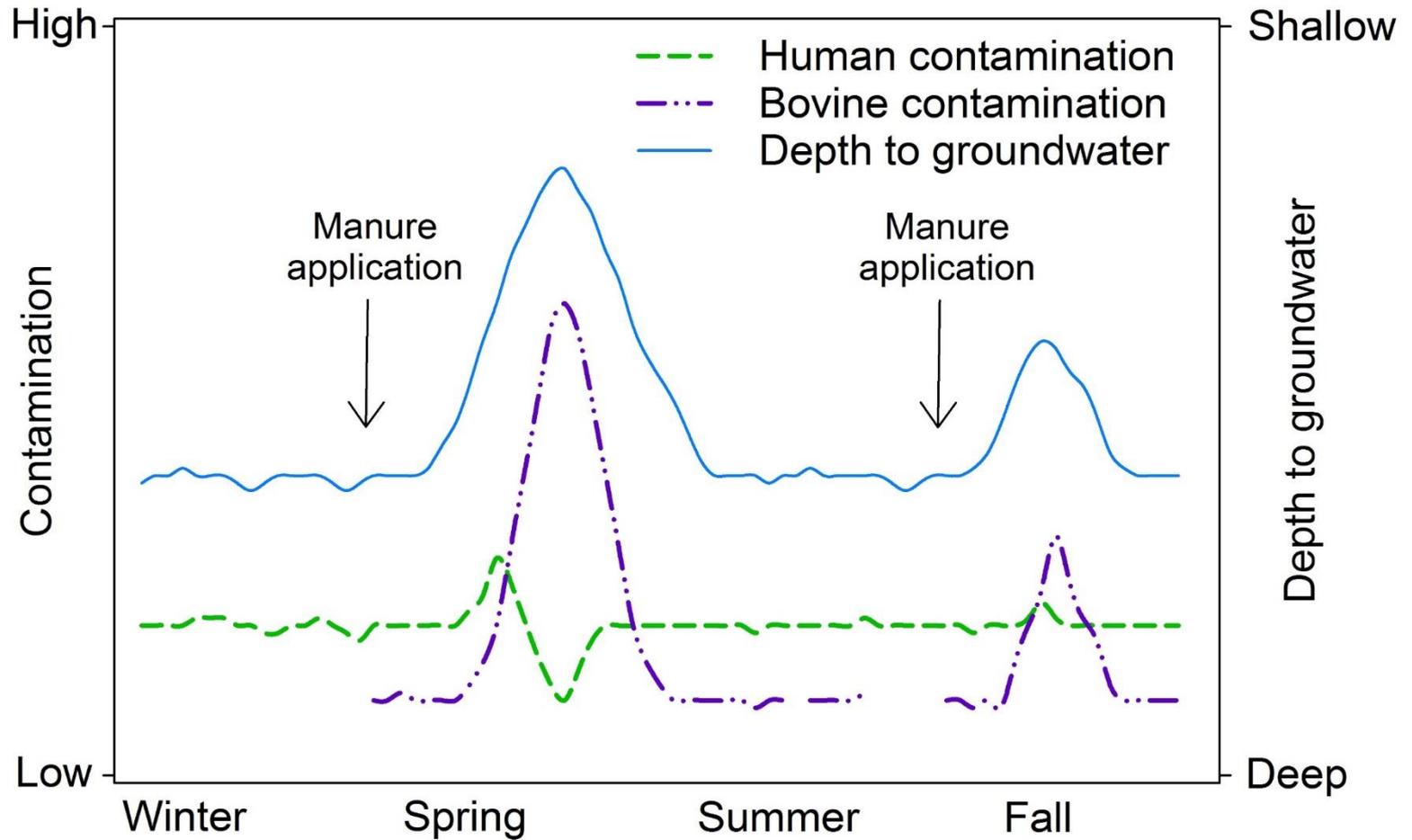
- Large fecal source
- Surface applied periodically
- Episodic infiltration

## Human pathogen source

- Small fecal source
- Sub-surface release continuously
- Continuous infiltration



# Conceptual Model of Fecal Contamination in Kewaunee County - 2



# Objective 3: Statistical Risk Factor Analysis

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## **Contamination Sources** – *Distance; count or acres within 750, 1500, 3000 ft of well*

### Bovine

Agricultural fields  
Manure storages

### Human

Septic systems (all, drain field, uninspected)  
Septage-applied fields

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## **Karst Features** – *Count within 750, 1500 and 3000 feet of well*

Sinkholes  
Bedrock features at the surface

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## **Precipitation & Groundwater Recharge** – *2, 7, 14, 21 days prior to sampling*

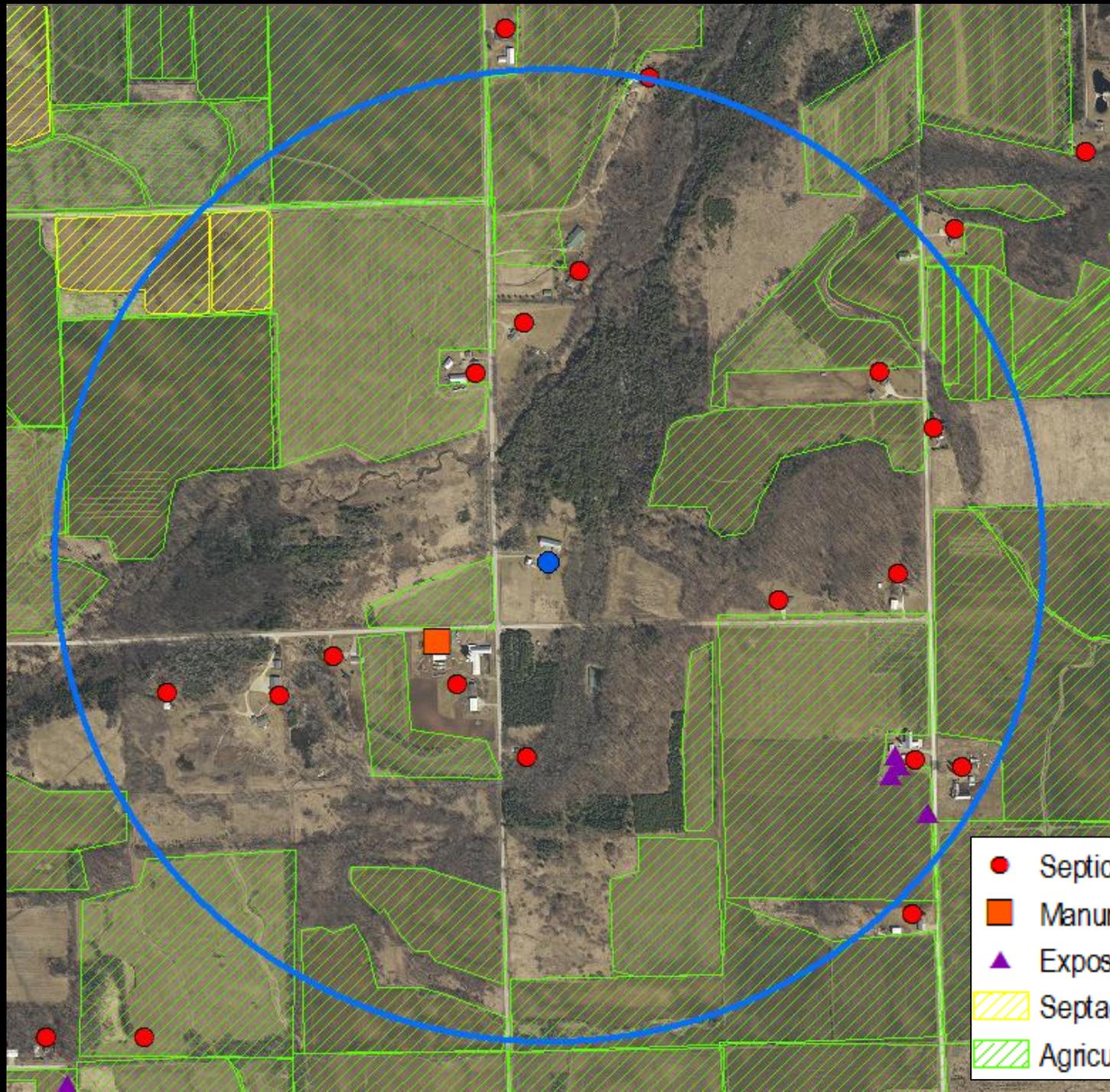
Precipitation (cumulative, no snowfall)  
Groundwater recharge (cumulative)  
Depth to groundwater (median & minimum)

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## **Well Construction & Siting**

Well depth	Well age
Casing depth	Length of casing into bedrock
Depth to groundwater	Length of casing below water table
Depth to bedrock	Elevation
Open interval length	Soil drainage

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- Septic system
- Manure storage
- ▲ Exposed bedrock & sinkholes
- ▨ Septage field
- ▨ Agricultural field

# Risk Factors for Human Fecal Contamination

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## Important factors

- Distance to nearest neighbor's septic system
- Number of neighbors' septic drain fields around well
- Precipitation
- Depth to groundwater
- Depth to bedrock

## Surprising unimportant factors

- Well construction variables (e.g., casing depth)

# Risk Factors for Manure Contamination

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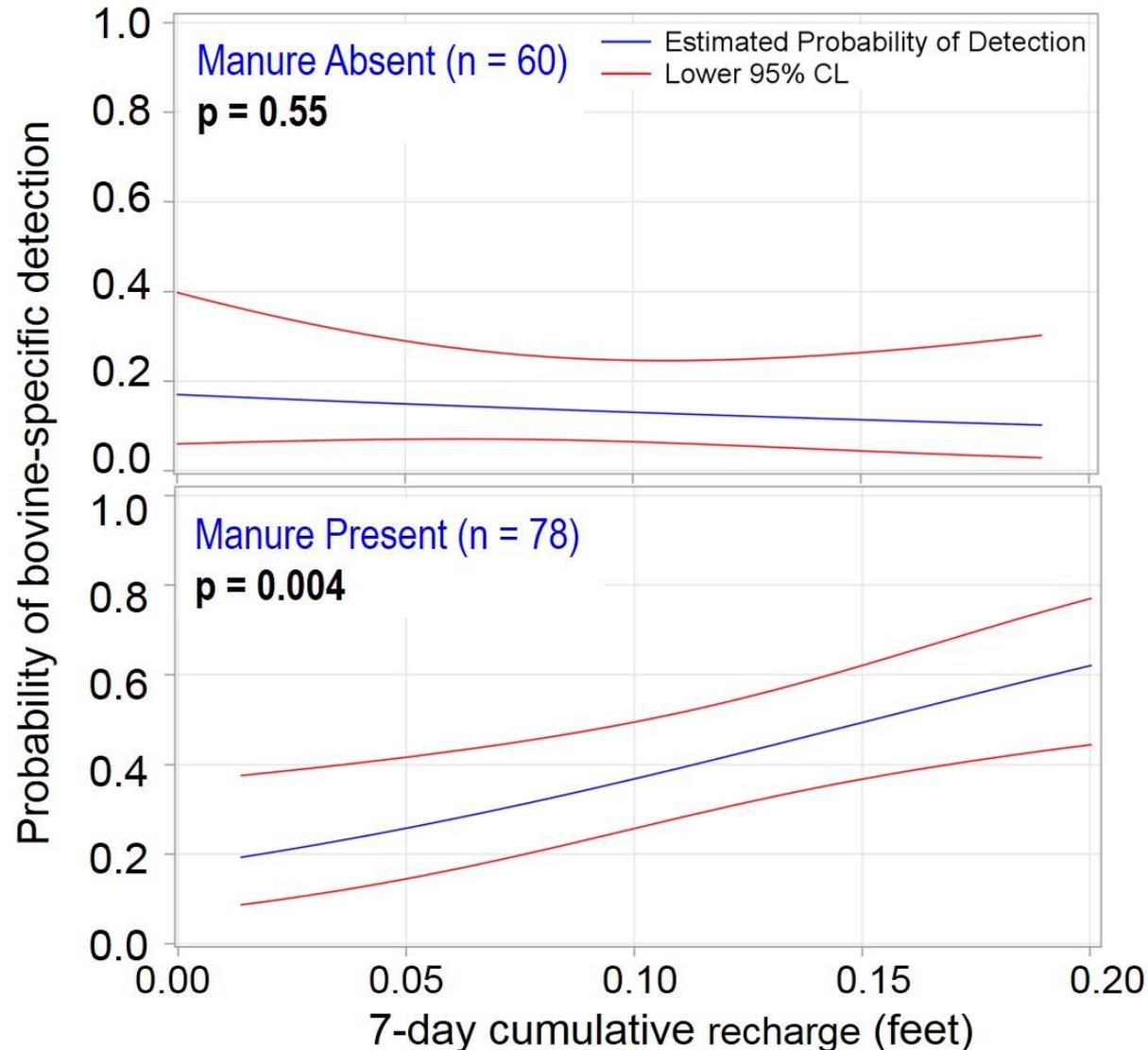
## Important factors

- Recharge
- Depth to groundwater
- Depth to bedrock
- Interactions between ag land use, manure application, and groundwater recharge

## Surprising unimportant factors

- Well construction variables (e.g., casing depth)

# Bovine-Specific Microbes in Private Wells by Previous 7 Days Cumulative Recharge



# Risk Factors for Total Coliform and High Nitrate Contamination

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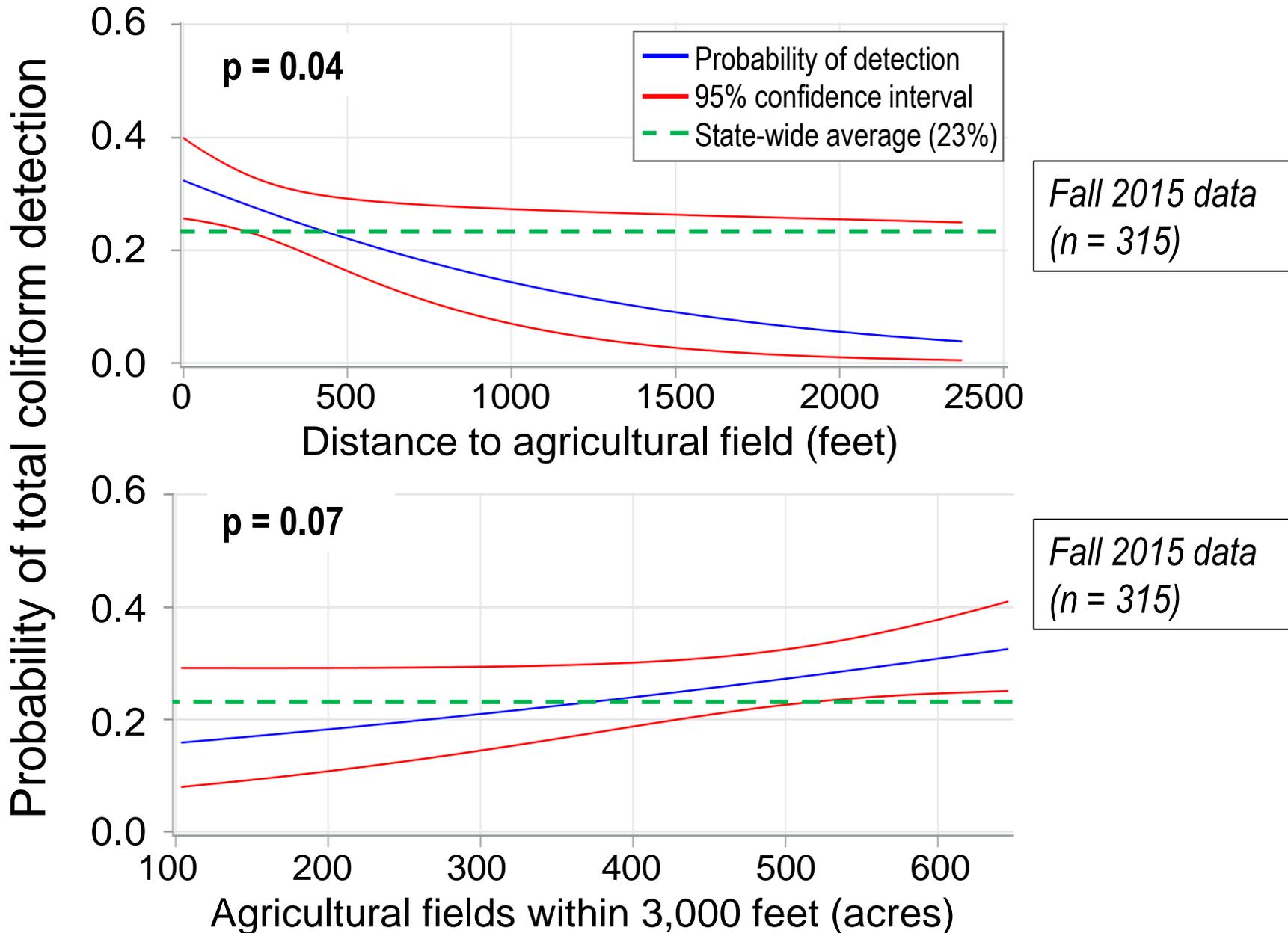
## Important factors

- Distance between well and agricultural field
- Number of acres of ag fields around well
- Number of sink holes around well
- Distance to nearest manure lagoon
- Recharge
- Depth to bedrock

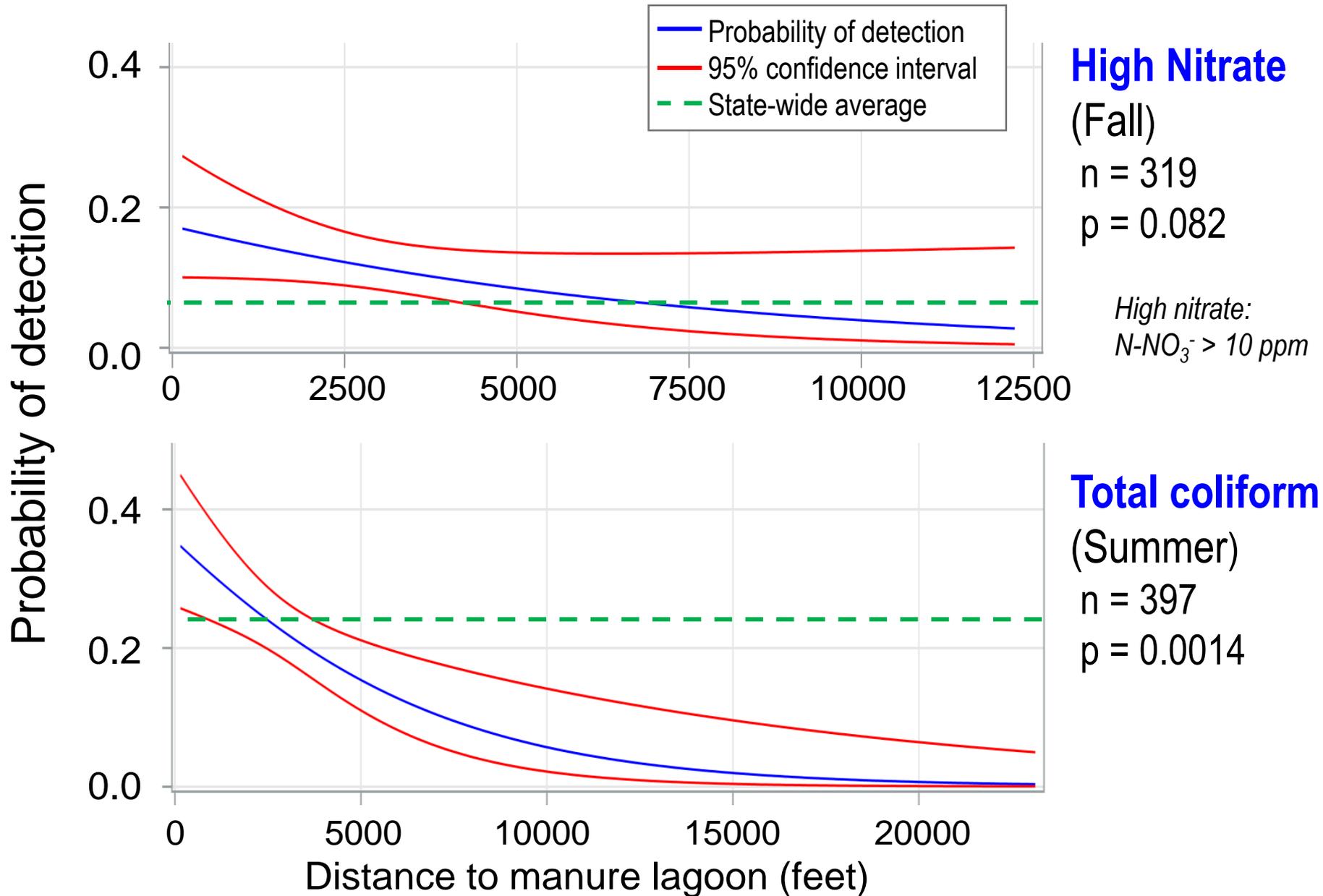
## Surprising unimportant factors

- Septic system variables
- Well construction variables (e.g., casing depth)

# Risk Factors for Total Coliform Contamination



# Manure Lagoon Association with Coliforms and Nitrate



# Do people get sick from drinking contaminated private well water?

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- Consider one pathogen: *Cryptosporidium parvum*
- Confirmed cryptosporidiosis cases in Kewaunee County reported to State:
  - 2 to 9 cases per year (2010 to 2016)
- Under-reporting of cryptosporidiosis cases in the USA is estimated to be 100-fold (Centers for Disease Control and Prevention, 2012)
- Therefore, in Kewaunee County there are likely 200 to 900 cryptosporidiosis cases per year

How many of these cases are from private wells?

# Estimate of Kewaunee County *Cryptosporidium parvum* infections from private wells

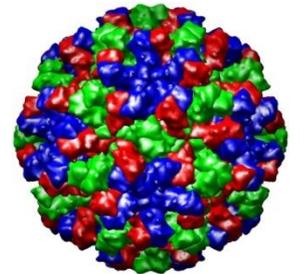
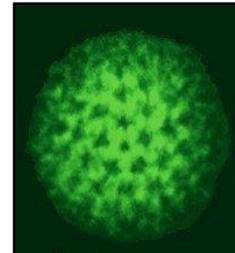
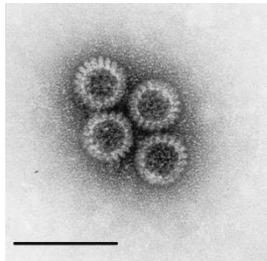
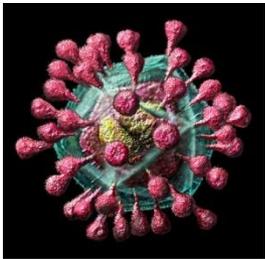
	People	Calves
Population using private wells	12,200	17,300
Wells contaminated by <i>C. parvum</i>	3.1%	3.1% (assumed)
Population exposed per day	380	540
Infections per exposure	10 infections per 10,000 people	85 infections per 10,000 calves
Total infections per year	140	1,700

# Solutions for Preventing Exposure to Manure-borne Pathogens in the Environment

Practices to Minimize Transport	Practices to Maximize Inactivation
Distance between livestock and waterways	Storage time
Vegetated treatment areas	Chemical treatment (e.g., lime)
Settling basins and wetlands	Thermophilic processes (e.g., aerated composting)
Manure storage and treatment lagoons	Anaerobic digestion

*From: Atwill et al. 2011 An Introduction to Waterborne Pathogens in Agricultural Watersheds, NRCS*

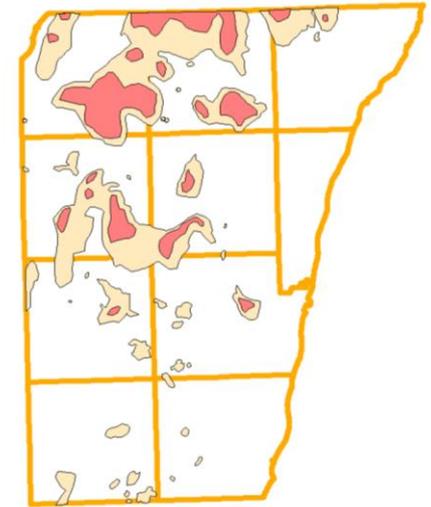
Questions?  
Comments?



# County-wide contamination rate; weighted by depth to bedrock

## Percent wells contaminated

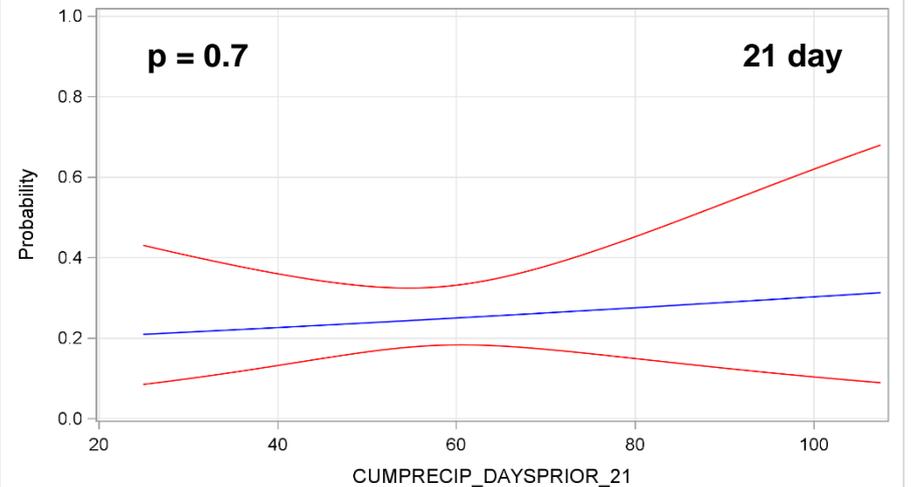
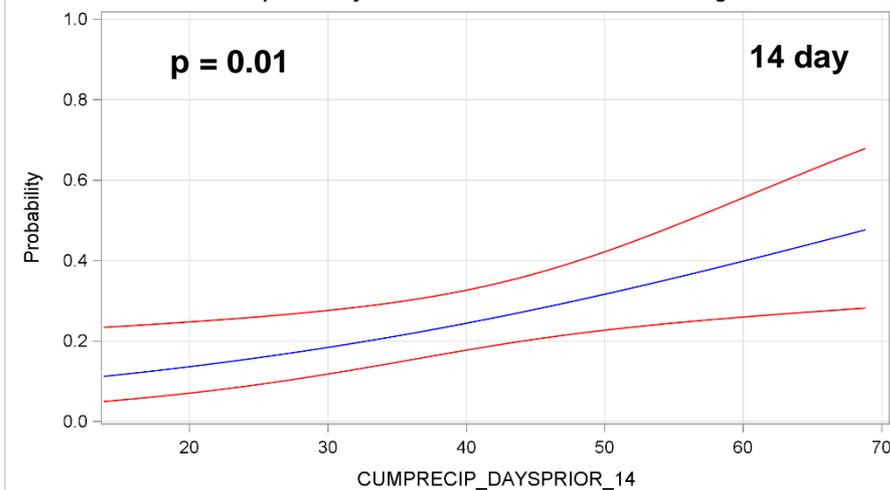
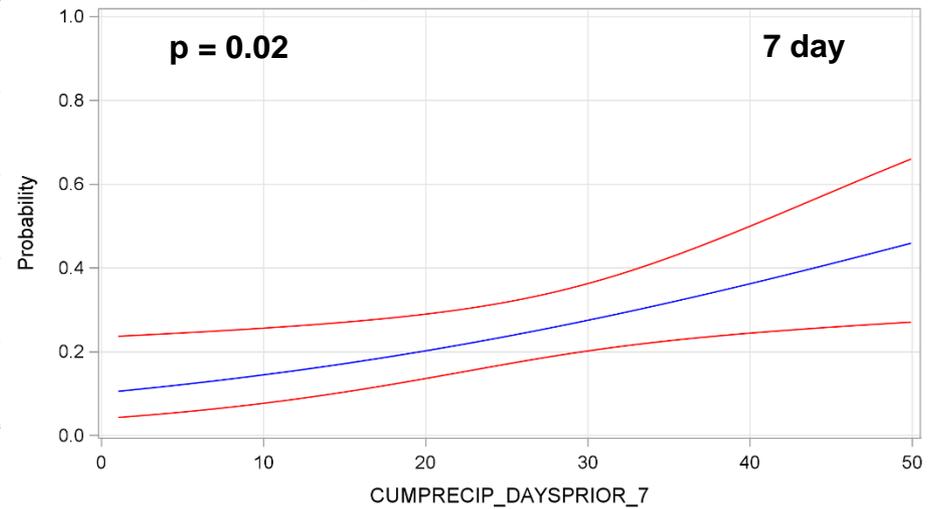
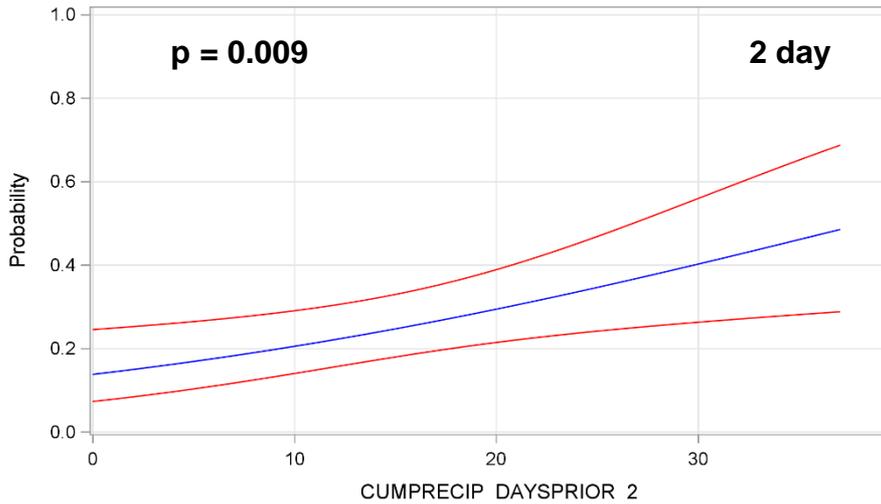
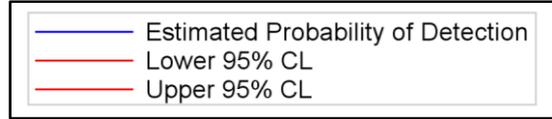
	Kewaunee County		Wisconsin*
	Recharge (n = 317)	No Recharge (n = 400)	(n = 534)
Total coliform	20.8	22.2	22.8
<i>E. coli</i>	0.4	1.2	2.6
High nitrate	7.4	6.8	6.6
Any of the 3 contaminants	26.4	27.6	NA



*High nitrate: exceeds health standard; N-NO<sub>3</sub><sup>-</sup> > 10 ppm*

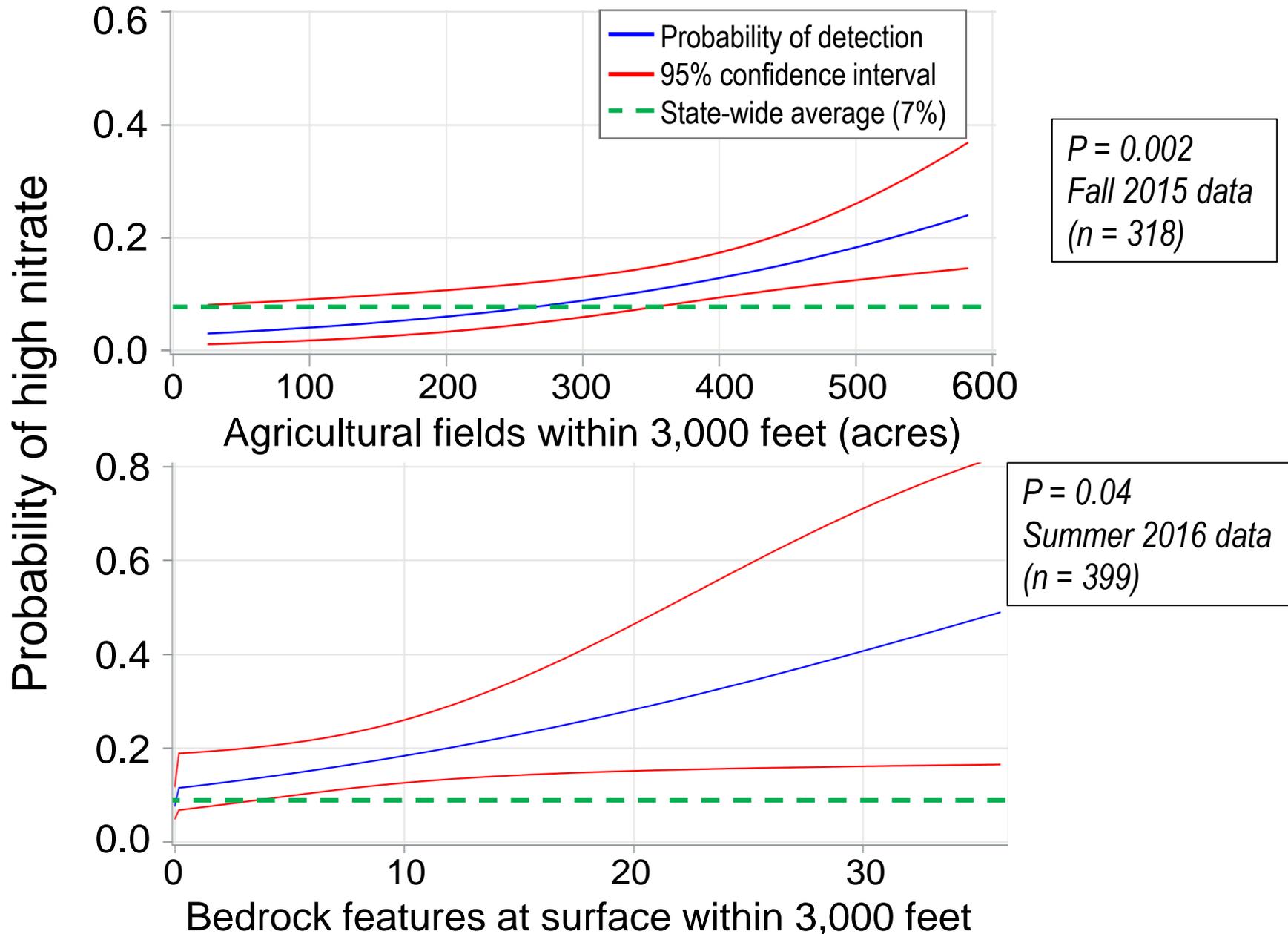
*\*private wells sampled; Information on the quality of water found at community water systems and private wells. United States GAO/RCED-97-123, June 1997*

# Probability of well contamination with human fecal markers as related to cumulative precipitation



Cumulative precipitation in millimeters

# Risk Factors for High Nitrate ( $N-NO_3^- > 10 \text{ ppm}$ ) Contamination



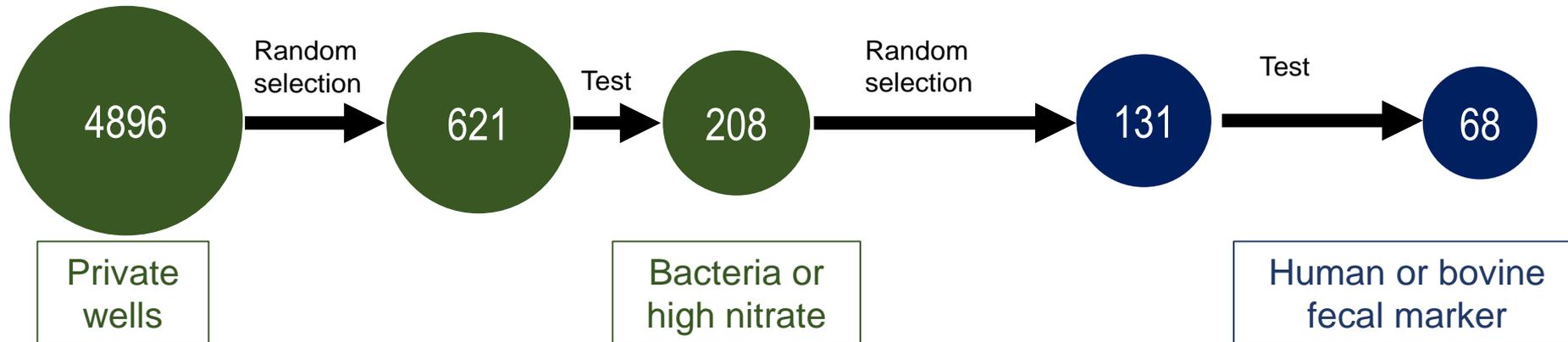
# Project objectives & study design

1. Measure total coliform, *E. coli*, nitrate



2. Determine fecal source

*Given  
contamination*



**Outcome:** County-wide occurrence as % wells contaminated

**Outcome:** Number of wells with human or bovine fecal markers

# Pathogens & fecal markers in Kewaunee County: Comparison to other studies



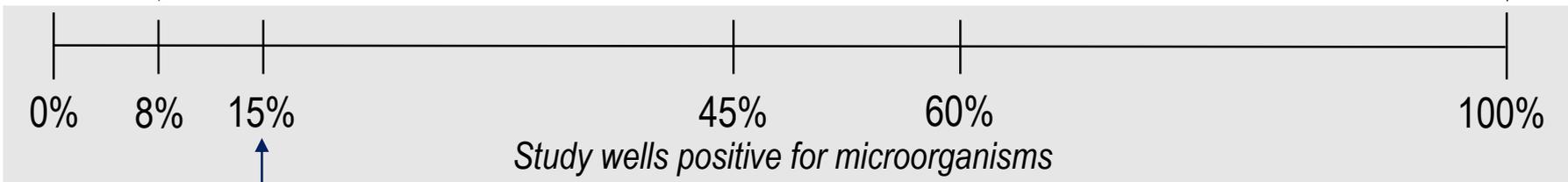
**Wisconsin 2003:**  
Private wells  
50 tested



**Ontario 2017:**  
Private wells  
11 tested



**Pennsylvania 2017:**  
Private wells  
5 tested



**Canada & USA 1990 – 2013**  
Public & private wells  
12,616 tested



**Kewaunee County**  
Private wells  
131 tested



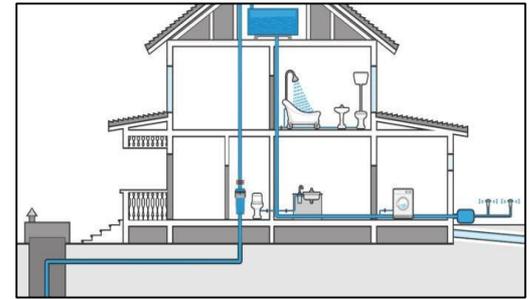
# From Farm Field to Household Well



Manure applied Oct 25, 2016



> 1 inch rain Oct 26, 2016



House near field

Farm field sampled Oct 27, 2016



Bovine Bacteroides  
Bovine enterovirus  
Bovine polyomavirus  
M2 Bacteroides-like  
M3 Bacteroides-like

Rotavirus A NSP3  
Rotavirus A VP7  
Rotavirus C

Tap water Oct 27, 2016

Bovine Bacteroides  
Bovine enterovirus  
Bovine polyomavirus  
M2 Bacteroides-like  
M3 Bacteroides-like  
Campylobacter jejuni  
Cryptosporidium  
Rotavirus A NSP3  
Rotavirus A VP7  
Rotavirus C



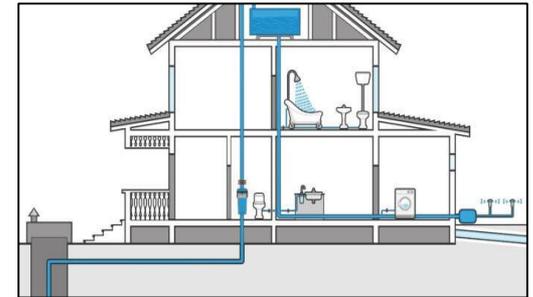
# From Farm Field to Household Well



Manure applied Oct 25, 2016



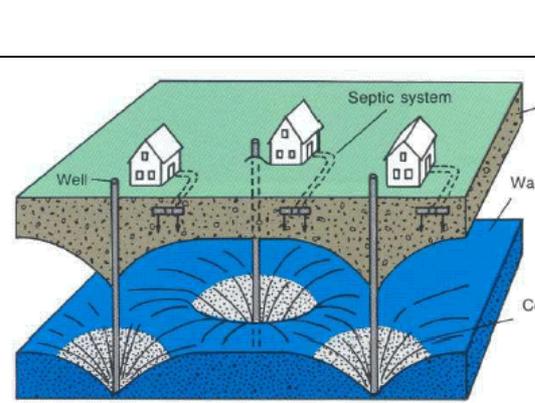
> 1 inch rain Oct 26, 2016



House near field

Neighbor's well sampled Oct 31, 2016

Tap water Oct 27, 2016



Bovine Bacteroides  
Bovine polyomavirus  
M2 Bacteroides-like  
M3 Bacteroides-like  
  
Rotavirus A NSP3  
Rotavirus A VP7  
Rotavirus C

Bovine Bacteroides  
Bovine enterovirus  
Bovine polyomavirus  
M2 Bacteroides-like  
M3 Bacteroides-like  
Campylobacter jejuni  
Cryptosporidium  
Rotavirus A NSP3  
Rotavirus A VP7  
Rotavirus C



# Summary

- Well contamination in the fractured dolomite aquifer in Kewaunee County results from both human and bovine fecal sources.
- Wells are contaminated with pathogens of significant concern: *Salmonella*, EHEC, *Cryptosporidium*, rotavirus.
- At depths to bedrock less than 50 feet total coliform and nitrate contamination rates generally exceed statewide averages.
- Risk factors for well contamination are: groundwater recharge, depth to groundwater, sink holes, precipitation, timing of manure application, agricultural land use, and the density of septic drain fields.

# We Thank...

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**Kevin Masarik**

UW-Stevens Point Watershed Science and Education

**Kevin Erb**

UW-Extension – Environmental Resources Center

**Leah Kammel & Laura Hubbard**

Upper Midwest Water Science Center

**Stephen Mael**

Wisconsin Geological & Natural History Survey

**Dustin Goering and Liz Houle**

NOAA National Weather Service

**Travis Engels**

Kewaunee County Land and Water Conservation

**Staff at the ERIC Lab, UW-Oshkosh**