

## Objectives

- Discuss the advantages and disadvantages between culture based methods and molecular panels for gastrointestinal pathogen detection
- Compare results of traditional laboratory procedures to results from multiplex testing of gastrointestinal pathogens
- Explore issues with culture independent testing.

## Disclosure and Acknowledgements

- In 2013 UNMC received money from BioFire to perform testing for FDA approval for the GI panels. I personally was not paid nor my salary was paid but money was paid to our Pathology department.
- My salary is paid by the ELC supplemental Ebola grant from the CDC.

## Issues Associated with Culture Independent Testing with Gastrointestinal Panels

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## Gastrointestinal Disease

- Major cause of Morbidity and Mortality World-wide
- Diarrhea disease global concern with 2 billion cases a year
- 1.5 million deaths of children less than 5 of age.



Do we need a multiplex panel for gastrointestinal pathogens?

- Overlapping symptomatology
- **Norovirus:** diarrhea, nausea, vomiting, stomach pain, fever
- **Salmonella:** diarrhea, stomach cramps, fever
- **Yersinia enterocolitica:** diarrhea, fever, abdominal pain, vomiting
- **Cryptosporidium:** diarrhea, stomach cramps, vomiting, fever
- Constant exposure to potential agents of gastroenteritis
- Each agent treated different—Some we can treat with antibiotics, relieve patient anxiety, rapid public health response



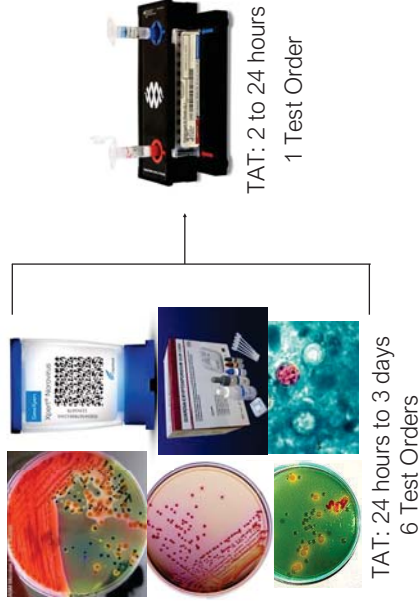
## Treatment for GI infections

- Oral/intravenous rehydration
- Antimicrobials maybe used in severe cases of gastroenteritis caused by certain pathogens.
- Timely and accurate diagnosis of stool pathogens is essential.

## Gold Standard Method for Identification

- Stool culture which require a 48 to 72 hours to complete.
- Enzyme-Linked Immunoabsorbant assay (EIA)based antigen detection.
- Microscopic detection
- Individual real-time PCR assay

Do we need a multiplex panel for gastrointestinal pathogens?



## Syndromic Testing

Allows the Health Care Professional to cast a wide net.



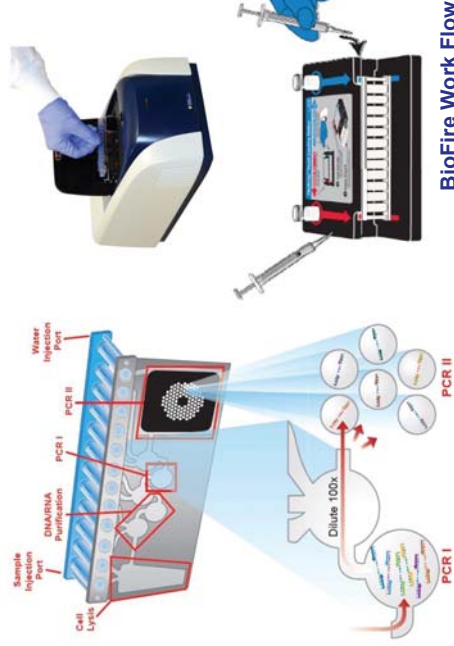
## Multiplex (Syndromic) Panels

- Reduced Sample Size
- Broad coverage without ordering specific tests.
- Enhanced ability to pick up co-infections
- Higher Through Put
- And in some cases higher sensitivity

## BioFire FilmArray™ Gastrointestinal Panel (GIP)

Bacteria	Diarrheagenic <i>E. coli</i> /Shigella	Parasites	Virus
<i>Campylobacter</i>	Enterogastric <i>E. coli</i> (EAGC)	<i>Cryptosporidium</i>	Adenovirus F 40/41
<i>Campylobacter</i>	Enteropathogenic <i>E. coli</i> (EPEC)	<i>Cyclospora cayentensis</i>	Astrovirus
<i>Campylobacter</i>	Enterotoxigenic <i>E. coli</i> (ETEC)	<i>Entamoeba histolytica</i>	Norovirus GI/GII
<i>Salmonella</i>	Shiga-like toxin-producing <i>E. coli</i> (STEC)	<i>Giardia lamblia</i>	Rotavirus A
<i>Vibrio</i>	<i>E. coli</i> O157		Sapovirus
<i>Vibrio cholera</i>	<i>Shigella</i> /Enterohemorrhagic <i>E. coli</i> (EHEC)		
<i>Yersinia enterocolitica</i>			

## The FilmArray Pouch



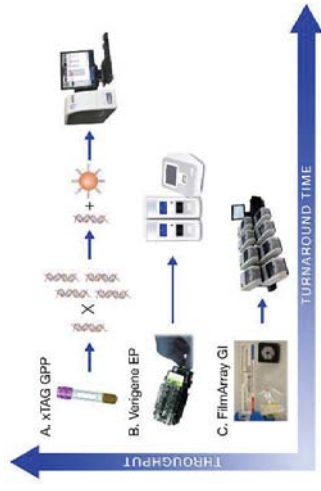
## BioFire FilmArray™ GIP

- The GIP had a sensitivity of 100% for 12/22 targets
- All other targets had a sensitivity of >94.5%
  - Figures for *Vibrio*, *V. cholera*, and *E. histolytica* were not calculated due to low incidence rates
- Specificity for all analytes was >97%

Analyte	No. of detections <sup>a</sup>	
	C	FA
<i>Campylobacter</i> spp.	35	58
<i>C. difficile</i>	165	204
<i>C. jejuni</i>	16	16
<i>S. enteritidis</i> spp.	31	37
<i>Vibrio</i> spp.	0	2
<i>V. cholera</i>	0	1
<i>Y. enterocolitica</i>	1	1
EAE	83	109
EPEC	317	348
ETEC	22	31
STEC	33	38
<i>E. coli</i> O157	3	4
<i>Shigella</i> spp. (EHEC) (culture) <sup>b</sup>	49	51
<i>C. parvum</i> spp.	18	24
<i>C. cryptosporidis</i>	19	19
<i>E. histolytica</i>	0	0
<i>G. lamblia</i>	20	27
Adenovirus F 40/41	44	55
Astrovirus	7	8
Norovirus GI/GII	55	70
Rotavirus A	6	18
Sapovirus	6	59

## Multiplex Panels for Gastrointestinal Pathogens

- FIVE FDA-cleared GI Panels
- **Hologic** - *Salmonella*, *Shigella*, *Campy*, Shiga toxin positive *E. coli* (STEC)
- **BD Max** - *Salmonella*, *Shigella*, *Campy*, STEC, *Vibrio*, *Y. enterocolitica*, Norovirus, Rotavirus
- **Nanosphere** - *Salmonella*, *Shigella*, *Campy*, STEC, *Vibrio*, *Y. enterocolitica*, Norovirus, Rotavirus
- **Luminex** - *Salmonella*, *Shigella*, *Campy*, STEC, O157, ETEC, *V. cholera*, *Y. enterocolitica*, *C. difficile*, Adenovirus, Norovirus, Rotavirus, *Cryptosporidium*, *E. histolytica*, *Giardia*
- **BioFire** - Most comprehensive to date, 14 bacterial targets, 5 viral targets, 4 parasitic target



FilmArray™ GI Panel		BIO FIRE	
Run Summary	Sample ID: ██████████	Run Date: 01/22/2015	Controls: Passed
Detected:	<i>Yersinia enterocolitica</i>		
Result Summary			
Not Detected	<i>Campylobacter</i>		Bacteria
Not Detected	<i>Clostridium difficile</i> toxin A/B		
Not Detected	<i>Plesiomonas shigelloides</i>		
Not Detected	<i>Shigella</i> spp.		
Not Detected	<i>Vibrio</i>		
Not Detected	<i>Yersinia enterocolitica</i>		
Not Detected	Diarrheagenic <i>E. coli</i> /Shigella		
Not Detected	Enterogastric <i>E. coli</i> (EAGC)		
Not Detected	Enteropathogenic <i>E. coli</i> (EPEC)		
Not Detected	Enterotoxigenic <i>E. coli</i> (ETEC)		
Not Detected	Shiga-like toxin-producing <i>E. coli</i> (STEC) var f/af2		
Not Detected	Shiga-like toxin-producing <i>E. coli</i> (STEC) var f/af2		
Not Detected	Shiga-like toxin-producing <i>E. coli</i> (EHEC)		
Not Detected	Parasites		
Not Detected	<i>Cryptosporidium</i>		
Not Detected	<i>Cyclospora</i>		
Not Detected	<i>Entamoeba histolytica</i>		
Not Detected	<i>Giardia lamblia</i>		
Not Detected	Virus		
Not Detected	Adenovirus F 40/41		
Not Detected	Astrovirus		
Not Detected	Rotavirus A		
Not Detected	Sapovirus		
Run Details	Run Name: C:\Users\JL11	Run Date: 01/22/2015	Run Time: 11:33:33
	Serial No.: 02919255	Lot No.: 248175	Operator: Amanda Starny (panata)
			Instrument: ZFA00353

## Use of BioFire FilmArray™

- **Advantages**
  - Increased sensitivity
  - Rapid turnaround time
  - Streamlines ordering procedures
- **Disadvantages**
  - Increased cost
  - Public health reporting necessitates organisms in culture

## BioFire FilmArray™ GIP

TABLE 6 Results of discrepant analysis

Analyte by FilmArray GI Panel result	Total no.	Inconclusive	FA correct	FA incorrect
<i>Campylobacter</i> spp.	24	5	19 <sup>d</sup>	
<i>C. difficile</i> toxin A/B	41		41	
<i>P. shigelloides</i>	15		15	
<i>Salmonella</i> spp.	6		6	

- In total, 199 of 237 GIP false positives were confirmed by an alternative method, confirming the positive result from the GIP
- Occasional cross reactivity between both ETEC and *Giardia* with commensal organisms
- Conventional culture routinely misses *P. shigelloides*, *Salmonella*, *Vibrio*, and *C. upsaliensis*
- Effective for the diagnosis of pathogens that were previously not routinely tested for such as diarrheagenic pathotypes of *E. coli*, Sapovirus, and Astrovirus

## Implementation

- Stool culture and *Cryptosporidium*/*Giardia* EIA were discontinued and all other traditional testing was maintained
- Testing maintained in-house
  - Full ova and parasite (including modified acid fast)
  - Rotavirus
  - Norovirus
  - Adenovirus
- Also implemented an *Aeromonas* only culture
- Guidance document was written and circulated with results of all GIP testing

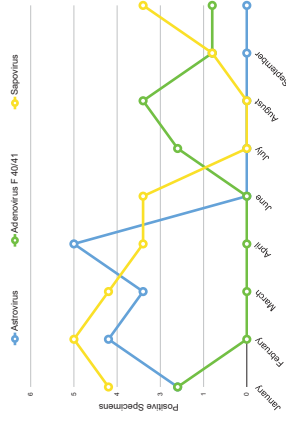
## Patient Demographics

Tests Run	Positive Tests*	% Positivity
Inpatient	703	23%
Outpatient	877	29%
Reference	636	31%
Public Health	39	67%

\*Includes all analytes except *C. difficile*

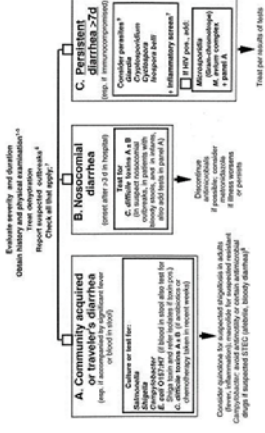
- Testing was distributed between inpatients, outpatients, reference clients (clinics, small hospitals throughout Western Iowa/Eastern Nebraska)
- Some testing was also run for public health purposes when concerns of an institutional outbreak were present

## Prevalence of under diagnosed pathogens



**Practical Guidelines for the Management of Infectious Diarrhea**  
 Richard C. Guerrant, Thomas Van SOEST, Ted S. Stoner, Nathan M. Theelman, Laurence Slutsker, Robert V. Tauxe, Thomas Hennessy, Patricia M. Griffin, Herbert DuPont, R. Bradley Sack. Clin Infect Dis (2011) 52 (8): 331-351. DOI:https://doi.org/10.1093/cid/cir114

Figure 1



## Implementation - Treatment Guidance

Table 1. Etiology and Treatment Recommendations<sup>a, b, c, d, e, f, g, h, i, j, k, l, m, n, o, p, q, r, s, t, u, v, w, x, y, z</sup>

Pathogen	Common Presentation	Commonly Implicated Sources and Seasonality	Treatment Recommendations	Antibiotic (if indicated)
<b>Bacteria</b>				
<i>Campylobacter</i>	Fever, abdominal cramps, and diarrhea within 4-8 hours; fecal leukocytes and erythrocytes	Fresh water, poultry, milk and dairy products	Most patients recover without antimicrobial therapy. Antibiotics have been shown to reduce symptoms duration by 1-3 days and are useful in immunocompromised patients. Consider antimotility agents for control of diarrhea and risk factors for complications (eg, pregnant women).	Azithromycin 500 mg daily × 3 days Fluoroquinolone × 3 days <sup>h</sup> Immunocompromised patients may require prolonged therapy (7-14 days)
<i>Shigella</i> ( <i>Shiga</i> A/B)	More than 3 watery, lab findings may include blood and mucus	Recent antibiotic use, especially broad spectrum agents	Test not reported on panel. If CD is suspected, order the C. difficile toxin assay.	Microcystin 500 mg TID × 10-14 days Vancomycin 125 mg QID × 10-14 days
<i>Yersinia enterocolitica</i>	Fever, abdominal cramps, and diarrhea within 4-8 hours	Fresh water, seafood, institutional travel	Most patients recover without antimicrobial therapy. Consider antimotility agents in severe diarrhea, extremity of age, and immunocompromised	Fluoroquinolone × 3 days <sup>h</sup> TMP/SMX DS BID × 3 days
<b>Viruses</b>				
<i>Adenovirus</i> F 40/41	Vomiting and non-bloody diarrhea within 30-51 hours	Children < 2 yr, day care	No therapy available. Treat symptomatically.	Antibiotics not indicated
<i>Rotavirus</i>	Vomiting and diarrhea within 1-3 days	Children < 5 yr, day care, school, swimming pools, children, household, day care		
<i>Norovirus</i>	Vomiting and diarrhea within 1-2 days	Peak season - winter		
<i>Enterovirus A</i>	Vomiting and diarrhea within 1-2 days	Peak season - winter		
<i>Sapovirus</i>	Vomiting and diarrhea within 1-2 days	Children		

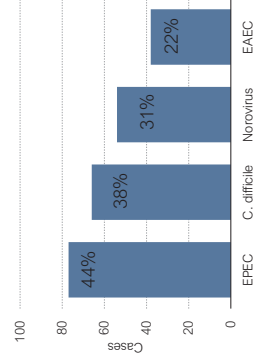
## GIP and Traditional Methods How Do They Compare

	2014	2015
Number of Tests	3417*	2255
Total Positive (%)	169 (4.9%)	370 (16.4%)
<i>Salmonella</i>	16	30
<i>Shigella</i> /EIEC	3	15
STEC	8	32
<i>Campylobacter</i>	18	51
Norovirus	115	197
<i>Giardia</i>	7	17
<i>Cryptosporidium</i>	2	28

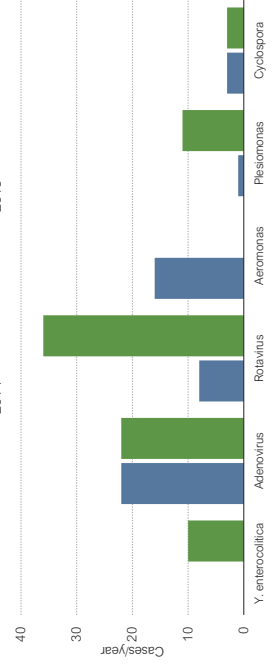
\*Combined Stool culture, Giardia/Crypto EIA, Norovirus PCR

## The high incidence of coinfection

- 19% (176/910) of positive panels had multiple targets detected
- 2 targets detected in 15.9% of positive specimens
- 3 or more targets detected in 3.3% of positive specimens

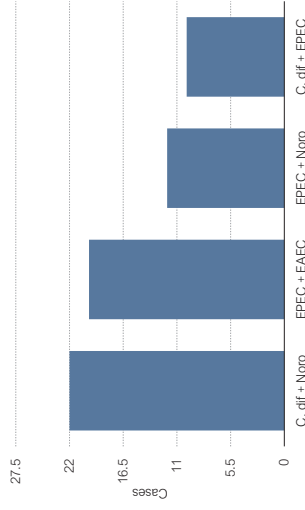


## Increased Detection with New Methods



- Increased detection of all targets on the panel compared to traditional methods with the exception of *Cyclospora* and *Adenovirus*
- The absence of *Aeromonas* on the panel led to decreased detection
- Limitation: difficult to compare year to year due to seasonality, outbreaks

## The high incidence of coinfection

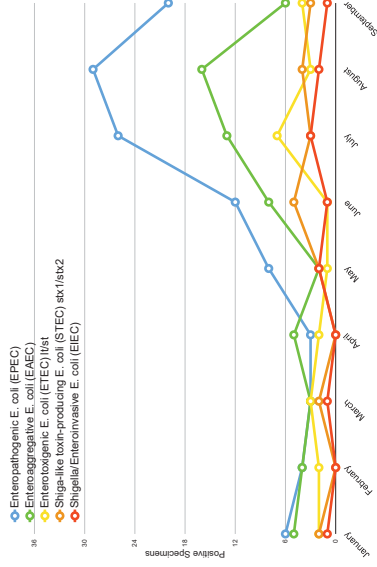


- The most common coinfection observed was *C. difficile* and Norovirus
- The presence of diarrheagenic *E. coli* in many coinfections occurred primarily in the summer months

## Diarrheagenic groups of *E. coli*

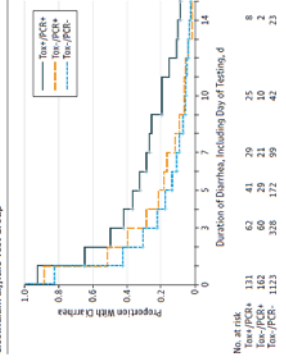
- Enterotoxigenic *E. coli* (ETEC)
- Enteroinvasive *E. coli* (EIEC)
- Enteropathogenic *E. coli* (EPEC)
- Enteroaggregative *E. coli* (EAEC)
- Diffusely adherent *E. coli* (DAEC)
- Enterohemorrhagic *E. coli* (EHEC)
  - *Escherichia coli* O157:H7

## Seasonality of Diarrheagenic *E. coli*



## Why not a solely molecular approach?

Figure 2. Kaplan-Meier Curves of Time to Resolution of Diarrhea by *Clostridium difficile* Test Group

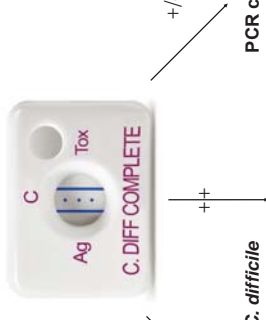


- Symptom duration and CDI complications for toxin - and PCR + patients resembles patients with no detectable *C. difficile*.

## How do you interpret this result?

- EPEC
- ETEC
- EAEC
- *Salmonella*
- STEC

## *C. difficile* Testing



### Negative for toxigenic *C. difficile*

### Positive for toxigenic *C. difficile*

- Molecular testing cannot distinguish between asymptomatic colonization and symptomatic infection

## Effect of CIDT on Public Health

- Rapid detection, increased sensitivity, and ability to detect more pathogens
- Public Health Laboratories require organisms in culture for confirmed diagnosis
- Loss of subtyping and other follow up testing (e.g. PFGE)
- Shift to whole genome sequencing will still require an isolate
- Local outbreaks may be easier to detect; tracking necessary for widespread outbreaks
- PulseNet data that is used to search for cluster identification and confirmation of outbreaks
- Loss of antimicrobial susceptibility testing



Figure 1. Estimates show that most cases for laboratory diagnosis will be reported as part of routine disease surveillance.

## The trouble with *Clostridium difficile*

- *Clostridium difficile* was the only target on the panel not routinely reported
- *C. difficile* diagnosis is typically done by two different methodologies
  - PCR to detect microbial DNA
  - Enzyme immunoassay to detect presence of toxin and antigen
- Reviewed literature showing the association of morbidity and mortality with toxin production. Overall, presence of toxin associated with increased morbidity and mortality, not presence of DNA. Toxin negative, PCR positive patients difficult to assess.
- Opted to maintain our traditional *C. difficile* testing algorithm
- When *C. difficile* was detected and no order was placed for *C. difficile* testing the provider was verbally notified of the result

## Comparison of *C. difficile* diagnosis using multiple platforms

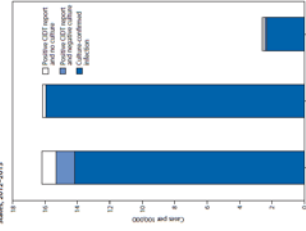
GIP Result	Antigen	Toxin	<i>C. diff</i> Confirm	Total
+	+	+	N/A	88
+	+	-	+	102
+	+	-	-	16
+	-	-	N/A	47
+	no additional testing ordered			86

- Most common target amplified on the GI panel 15% (342/2255)
- Only 35% of GIP + samples that were tested by the traditional algorithm were toxin positive
- As has been documented, molecular testing has increased sensitivity compared to EIA
- Will continue to evaluate the most appropriate way to test and report *C. difficile*



## Public Health and the GI Panel

FIGURE 1. Incidence of culture-confirmed bacterial infections and gastroenteritis by selected pathogens – FoodNet, United States, 2012–2013



Consultation with the Nebraska Public Health Laboratory resulted in an appropriate algorithm for follow up testing when necessary

As other institutions in the state implement CIDT, Nebraska Medicine is performing recovery cultures of appropriate pathogens

- Salmonella
- STEC
- Yersinia enterocolitica
- Vibrio
- Shigella and Campylobacter are recovered on the patient account for appropriate susceptibility testing

MMWR / March 13, 2015 / Vol. 64 / No. 10

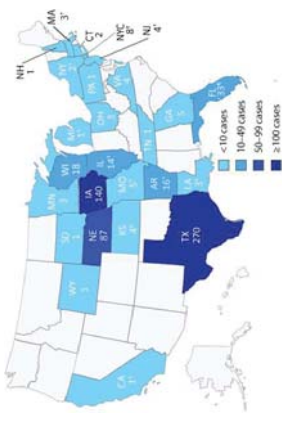
## Isolate Recovery from the GIP

- State actively monitoring sites where CIDT may be adopted
- Recommend follow up culture for confirmation or forwarding specimen to Nebraska Medicine for culture
- Prioritize Salmonella and STEC culture

Organism	Detected	Recovered (%)
Campylobacter	51	35 (69)
Shigella/EIEC	15	10 (67)
Salmonella	30	23 (77)
STEC	22	9 (41)
STEC O157	5	3 (60)
Vibrio	3	2 (67)
Y. enterocolitica	10	3 (30)

### Case #1: An unexpected outbreak

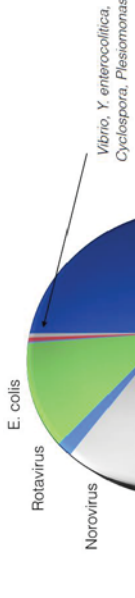
- Day 1 of GI Panel testing for FDA studies
- Cyclosporiasis Outbreak
- Multiple outbreaks affecting 25 states with 631 confirmed cases



### Case #2: A complex case of gastroenteritis

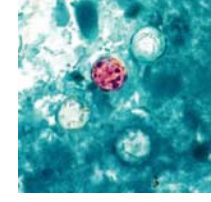
- 71 year old female transferred to Nebraska Medicine due to a pelvic fracture
- Also presented with acute kidney injury
- Reported a 2 week history of diarrhea
- GI Panel was ordered
  - Vibrio cholera, Plesiomonas, EPEC detected
- Blood cultures were drawn
  - Positive for Edwardsiella - also recovered from stool
- Subsequent blood culture was positive for Bacteroides

Do we need a multiplex panel for gastrointestinal pathogens?



Is there value in assays detecting rarely seen pathogens?

## Cyclospora cayentanensis



- Common in tropical and subtropical regions
- Spread through the ingestion of contaminated food or water.
  - First identified as a foodborne pathogen in the mid 1990s
  - Not passed person to person
- In the absence of treatment, symptoms may last up to a month and relapses are common
- Traditional diagnosis through modified acid fast staining
- Ability to link C. cayentanensis cases is still developing

TABLE 1. Ordering practices related to the stool specimens positive for Cyclospora using the BioFire FilmArray GI panel<sup>a</sup>

Time of specimen collection in relation to outbreak <sup>b</sup>	No. of specimens for which Cyclospora testing was or was not ordered by provider	
	Ordered	Not ordered
Before (before 28 June)	0	3
Early (between 28 June and 12 July)	3	5
After outbreak established (after 12 July)	8	0

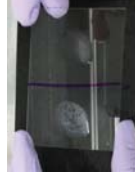
<sup>a</sup> For investigational use only.  
<sup>b</sup> The Centers for Disease Control and Prevention became aware of two domestically acquired, laboratory-confirmed cases of Cyclospora infection on 28 June (1).

- Loss of proficiency at parasite exam at many clinical laboratories lead to a backlog of specimens and delayed diagnosis
- Between 2014 and 2015 the incidence of Cyclospora was the same between traditional methods and GIP testing
- Another test site showed a 40% increase in diagnosis between the culture and GIP for a Shigella outbreak

J. Clin. Microbiol. 2015 Nov 5;53(11):3909.

## Vibrio cholera

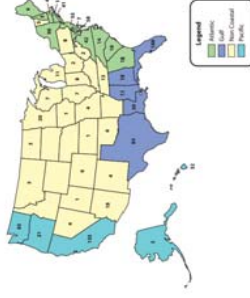
- Toxin PCR was performed at UNMC and subsequently at the CDC
  - The isolate was found to be nontoxicogenic
- Serotyping O1 and O139 was performed at NPHL and the CDC
  - O141 and O75 are found in the US and produce mild gastroenteritis
- The isolate has a unique PFGE pattern in their limited database (1200 isolates)
- In 2014, we had 5 orders for Vibrio with 0 recovered, in 2015 we detected 3 specimens positive for Vibrio



## Patient History

- Patient had recently returned from a trip in South Dakota
  - Suspicion for Vibrio was low
- Following multiple inquiries and recovery of V. cholera in stool culture the patient's husband recalled eating raw oysters in a local establishment
- Epidemiological investigation determined that the oysters were Blue Point oysters that likely originated in the Chesapeake Bay area

Figure 1. Number of cases of Vibrio infections (including Vibrio vulnificus, V. vulnificus O1 and O139, by state, 2013-2014. U.S. Bureau of Census.



At the end of February 2017 this sleepy little town got sick.



### Beaver City Gastrointestinal Illness Survey

The Nebraska Department of Health and Human Services (NDHHS), and South...

Case #3  
21 people in Beaver City, NE with gastroenteritis was picked up by CITD

- The first few patients were seen at a local clinic and samples sent to the near by critical access hospital. They do have a full microbiology but stools are primarily done by BioFire. Then 10 other patients in a course of two days.
- All twenty one people had *Campylobacter jejuna*.
- Southwest Public Health and the Department of Health and Human Service of Nebraska set up a survey to get to the bottom of this.
- A contaminated well on a farm that leached into the groundwater was suspected to be the cause.
- According to FoodNet, the percentage of *Campylobacter* diarrheal illnesses diagnosed only by CIDTs increased from 13% in 2012-2014 to 24% in 2016.

Evaluation of the BioFire FilmArray® Gastrointestinal Panel in a Midwestern Academic Hospital

- Studied published in December of 2016 in the European Journal of Clinical Microbiology Infectious Disease.
- Total of 2257 stools collected from January to December of 2015 were tested using the GIP.
- GIP detected one pathogen in 911(40%).
- Coinfections were detected in 176 (7.8%).
- The most frequently detected pathogens were *C. difficile* (15.2%), norovirus (8.9%), enteropathogenic *Escherichia coli* (7.1%), enteroaggregative *E. coli* (3.4%), *Campylobacter spp.* (2.3%) and sapovirus (2.0%).

## Conclusions

- Implementation of culture independent testing leads to increased detection of many causes of gastroenteritis and improved turnaround time
- Further studies needed to evaluate improvements in patient care, infection control practices, and appropriate antibiotic treatment
- Increased awareness of previously untested or under ordered pathogens
- Abundant coinfections may indicate important areas for further investigation
- Potential to rapidly diagnose outbreak associated infections
- Comprehensive testing will hopefully decrease unnecessary testing and provide patient and provider with a diagnosis
- Changes in the workforce might necessitate testing that requires less hands on time and less technical specialty

Recovery of bacteria from the Midwestern Academic Hospital for P/206public Health investigations varied

- 77% for *Salmonella*
- 30% for *Yersinia enterocolitica*.
- For stools positive for *C. difficile* on GIP that were tested by EIA, only 42.7% (88/206) were found to be producing the toxin.