



The Importance of Cryptosporidium Surveillance

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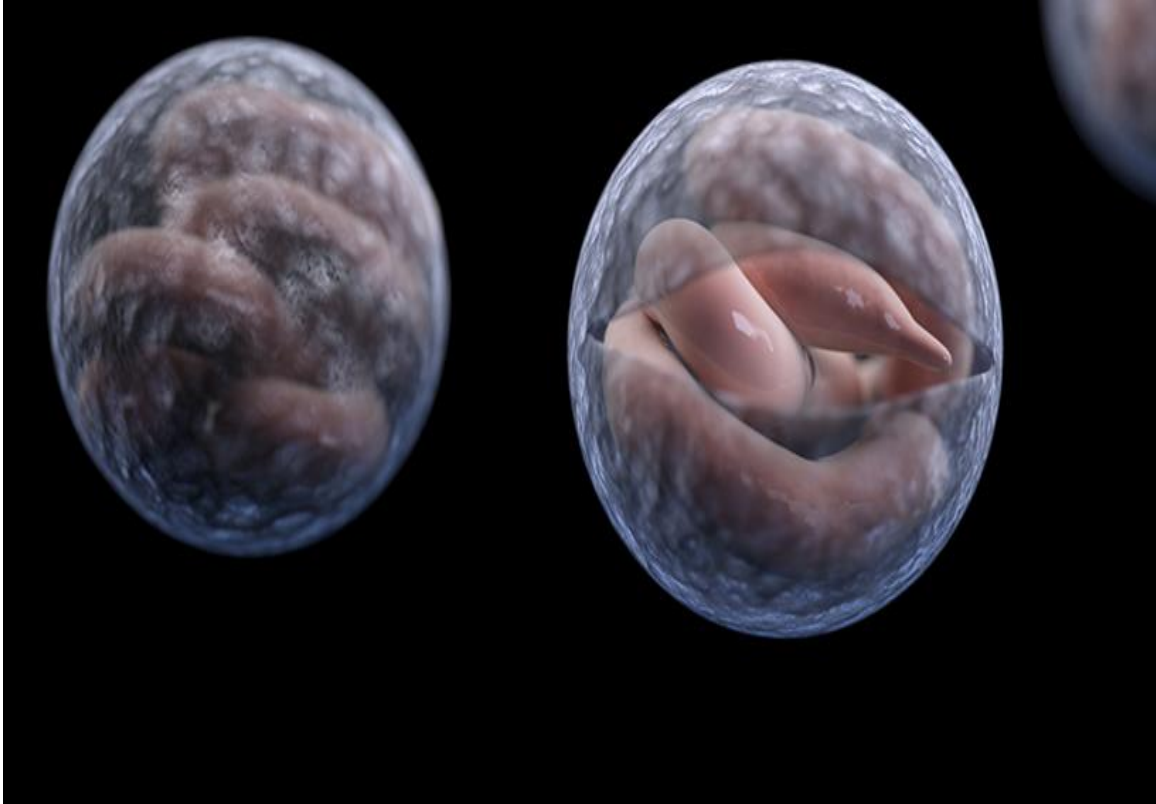
APHL-CDC Fellow

2022 WCLN Regional Meeting



Does your lab perform *Cryptosporidium* testing?

- A. Yes
- B. No



Cryptosporidium general information and diagnosis



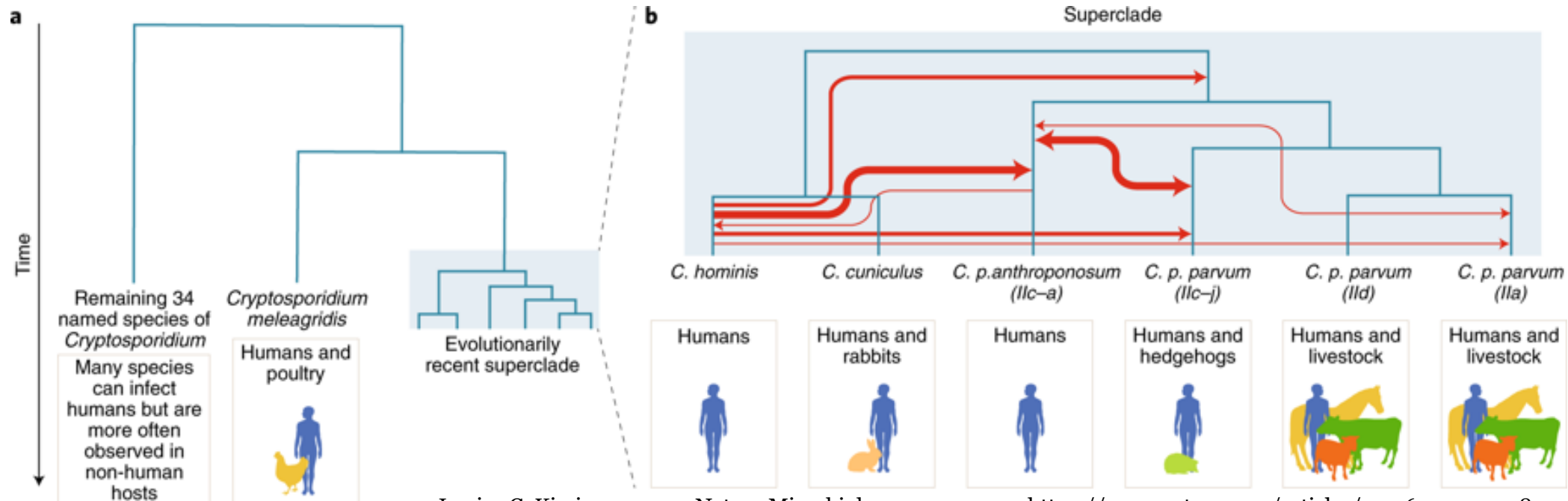
All started at a summer retreat

- 5 people have become ill one week after attending a summer family gathering at a water park in Wisconsin.
- All of them experienced diarrhea for several days. Some of them also experienced abdominal cramps, headaches, and body aches.
- Biofire GI panel conducted in a local clinical lab detected presence of *Cryptosporidium* species in patient stool and negative for any other GI pathogenic infections.



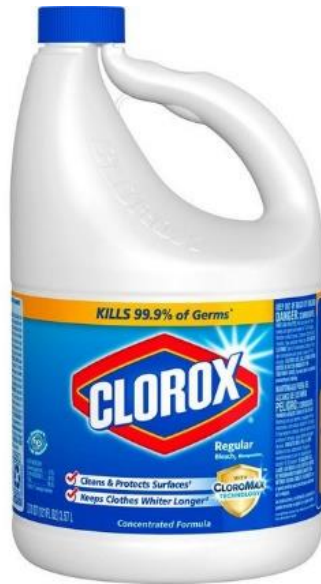
Cryptosporidium

- lives in the gut of infected humans or animals.
- ~40 species, but 2 in particular (*Cryptosporidium hominis* and *Cryptosporidium parvum*) account for most of infections.



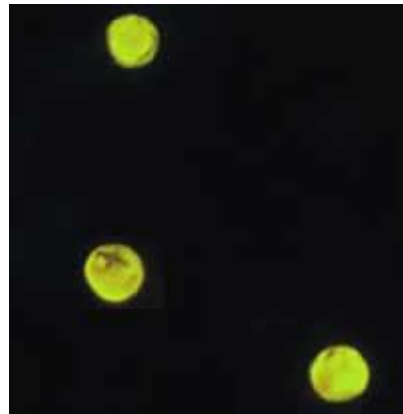
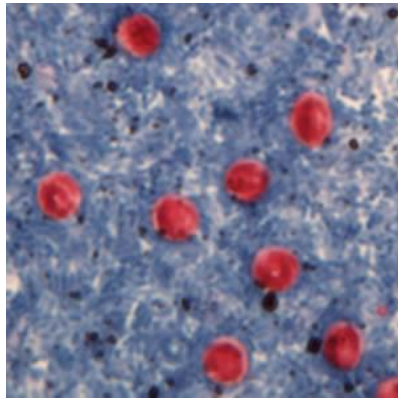
Jessica C. Kissinger. 2019. Nature Microbiology; 4: 730–731. <https://www.nature.com/articles/s41564-019-0438-1>

Cryptosporidium forms resistant oocysts



Cryptosporidium diagnosis methods

- Microscopy: Acid-fast stain, direct fluorescent antibody (DFA) assay.
- Rapid antigen cartridge test/microplate EIA tests
- PCR





What kind of test does your lab do to detect *Cryptosporidium*

- A. Microscopy
- B. Antigen-based test
- C. Molecular-based test

Risk factors

Diarrhea and swimming don't mix!

If you wouldn't do this...

...then why would you do THIS?

Don't swim or let your kids swim if sick with diarrhea!

Learn more at www.cdc.gov/healthyswimming

MMWR | SWIM HEALTHY, STAY HEALTHY

ABOUT 500 OUTBREAKS IN 15 YEARS

MOST OUTBREAKS HAPPEN

IN HOTEL POOLS AND HOT TUBS

IN THE SUMMER

STAY HEALTHY IN THE WATER

DON'T SWIM WITH DIARRHEA

CHECK THE INSPECTION SCORE

DON'T SWALLOW THE WATER

MORE THAN 27,000 PEOPLE SICK AND 8 DEATHS LINKED TO POOLS, HOT TUBS, AND WATER PLAYGROUNDS

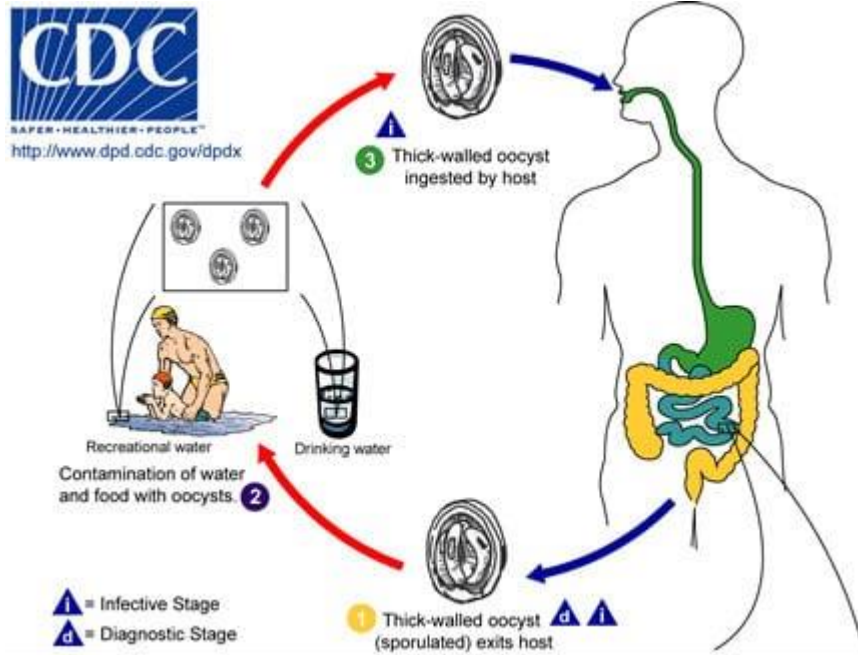
WWW.CDC.GOV

Data from 46 states and Puerto Rico from 2000-2014 as published in Hlavsa et al. MMWR, May 2016. <https://go.usa.gov/vQJ3G>

What risk factor causes the outbreak?

- There was an fecal incident reported in the kiddie pool by parents of one of the ill children.
- All of the ill children attended primary school before the family gathering.
- One of the children also attended a swimming lesson at a local YMCA.





U.S. Cryptosporidiosis Outbreaks: 2009–2017

Outbreaks of diarrhea most commonly linked to

Pools* (35%)



Don't swim
with diarrhea

Cattle (15%)



Wash hands after
touching animals

Childcare (13%)



Keep kids sick with
diarrhea home

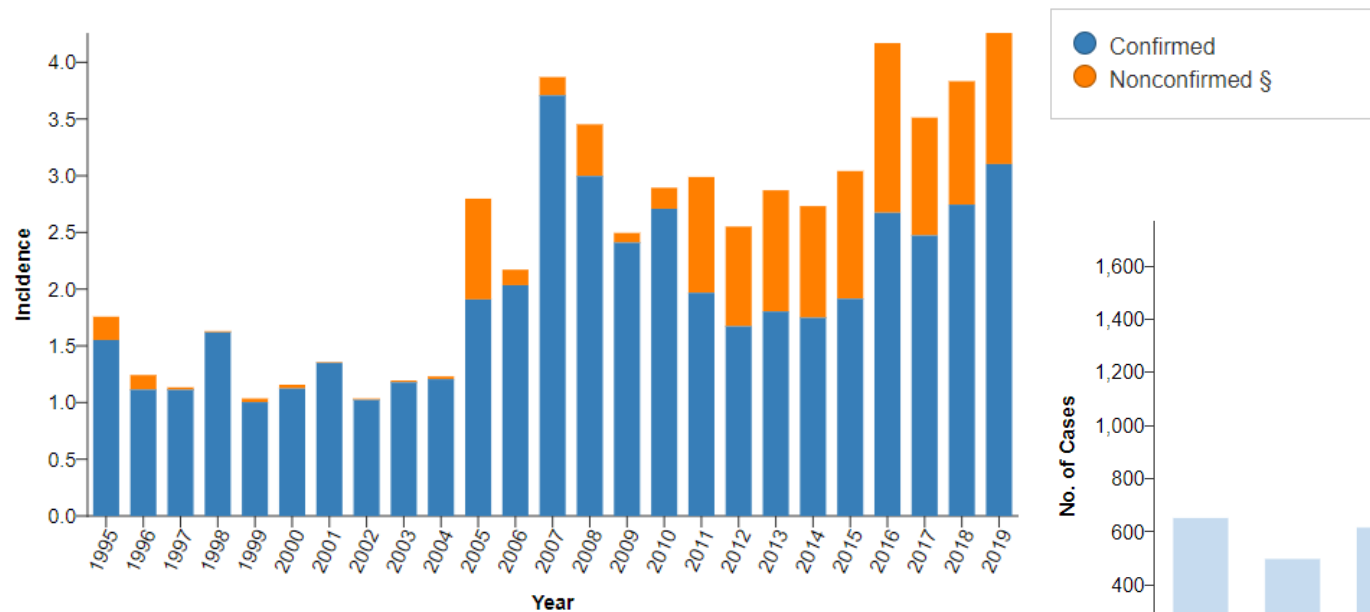
*Pools and water playgrounds
As reported in Gharpure et al. *MMWR* 2019 (bit.ly/MMWR627)

CDC MMWR

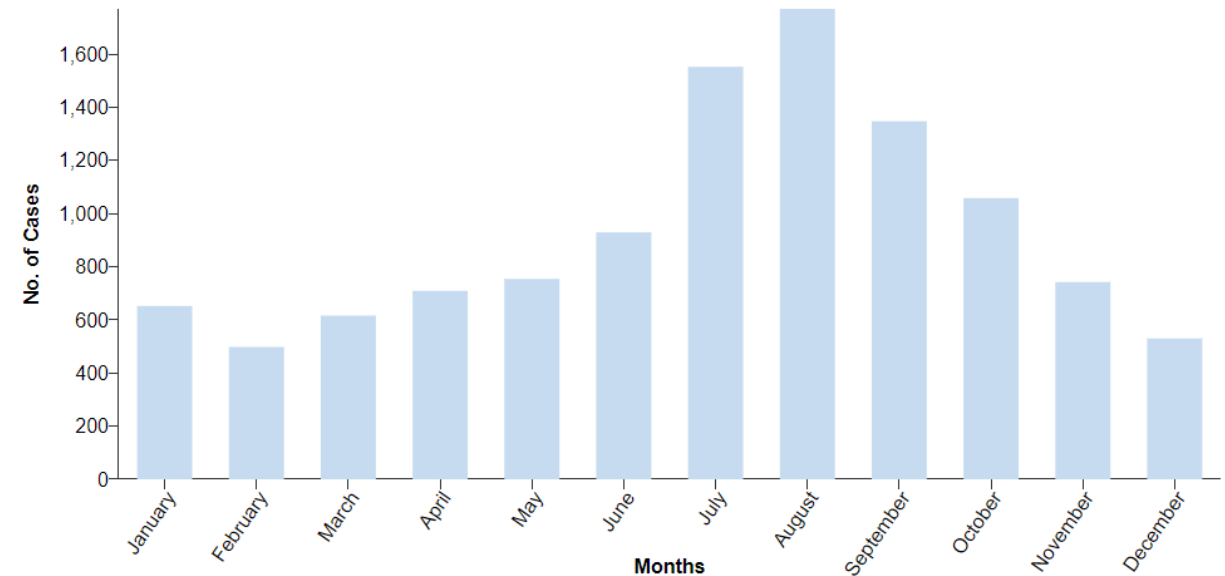


Disease Trend

Supplemental Figure 1. Incidence* of reported cryptosporidiosis cases, by year and case classification — National Notifiable Diseases Surveillance System, United States, 1995[†]–2019 (N=181,621)

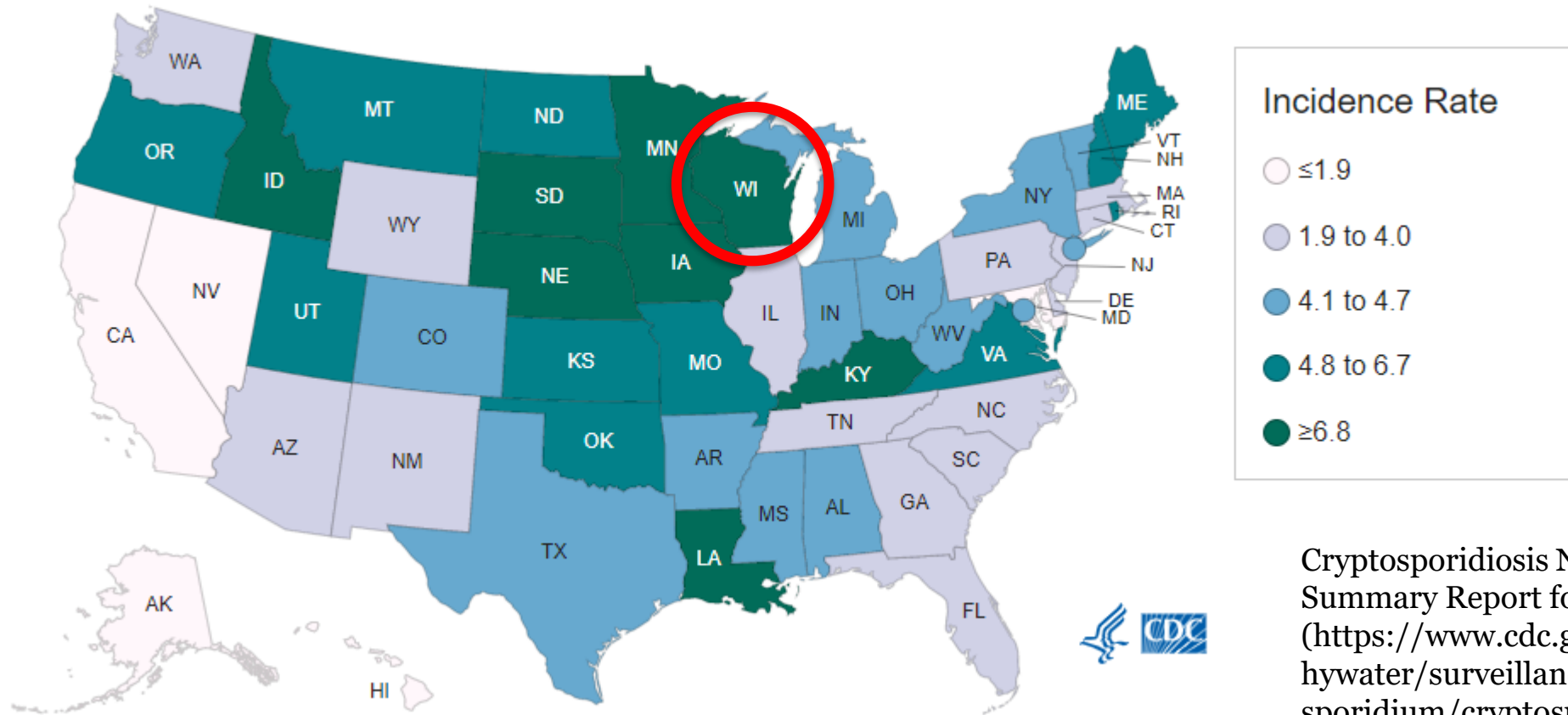


Cryptosporidiosis NNDSS
Summary Report for 2019
(<https://www.cdc.gov/healthwater/surveillance/cryptosporidium/cryptosporidium-2019.html>)





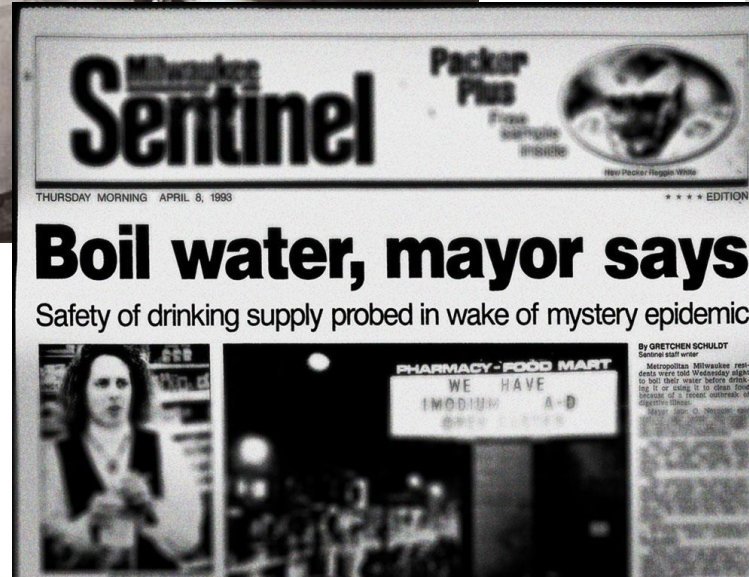
Geographic Distribution (per 100000)



Cryptosporidiosis NNDSS
Summary Report for 2019
(<https://www.cdc.gov/healthwater/surveillance/cryptosporidium/cryptosporidium-2019.html>)



1993 Waterborne Cryptosporidium Outbreak in Milwaukee



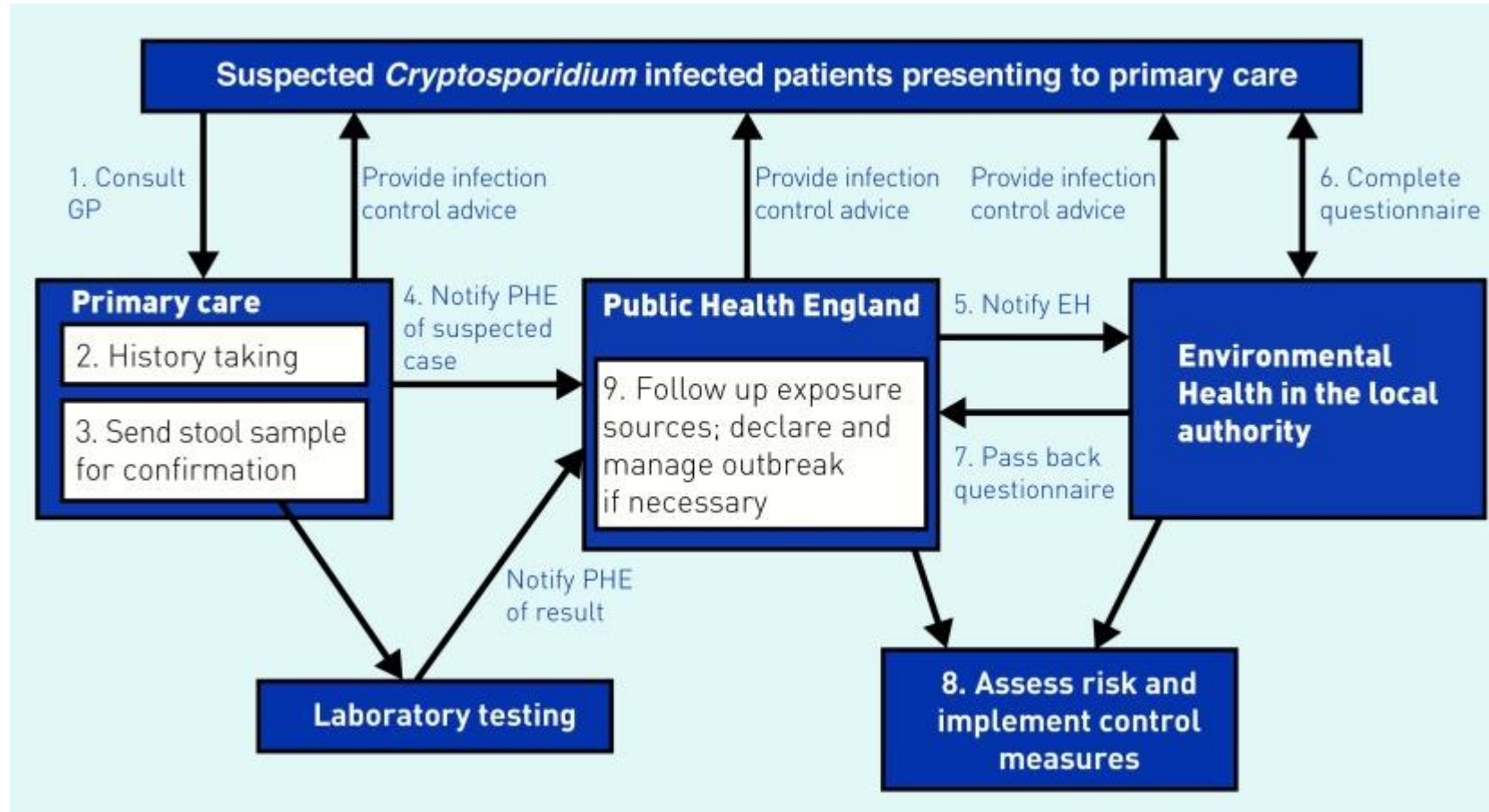
**Largest
Cryptosporidiosis
outbreak in the world.**



Identify the source of outbreaks



How do sources of outbreaks are identified



Horne S, Sibal B, Sibal N, Green HK. *Cryptosporidium* outbreaks: identification, diagnosis, and management. *Br J Gen Pract.* 2017 Sep;67(662):425-426. doi: 10.3399/bjgp17X692501. PMID: 28860302; PMCID: PMC5569737.



Genotyping

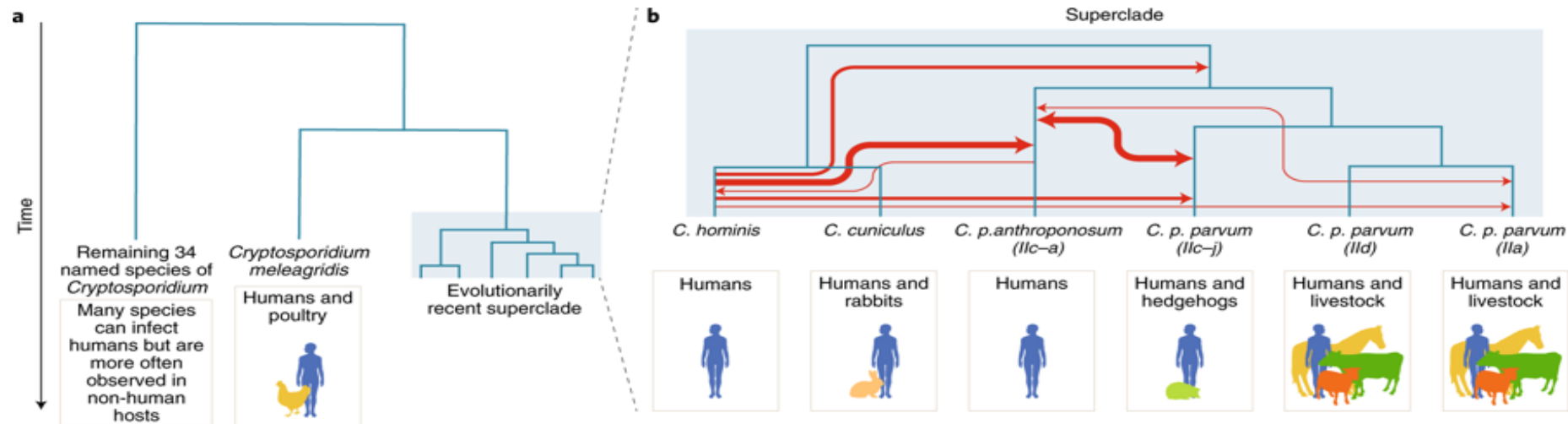
- Genotyping of *Cryptosporidium* can
 - Identify the *Cryptosporidium* species and subtypes that most commonly cause human cryptosporidiosis.
 - Identify the demographic and temporal-spatial distribution of common *Cryptosporidium* species and subtypes.
 - Identify the source of outbreaks and contamination.



CryptoNet

Identify the correct Crypto species

- CDC subtyping revealed that the infected individuals were infected with *Cryptosporidium hominis* subtype IfA12G1
- This strain was also isolated from patients during a recreational water associated *Cryptosporidium* outbreak in the North Shore Milwaukee area in the same year.





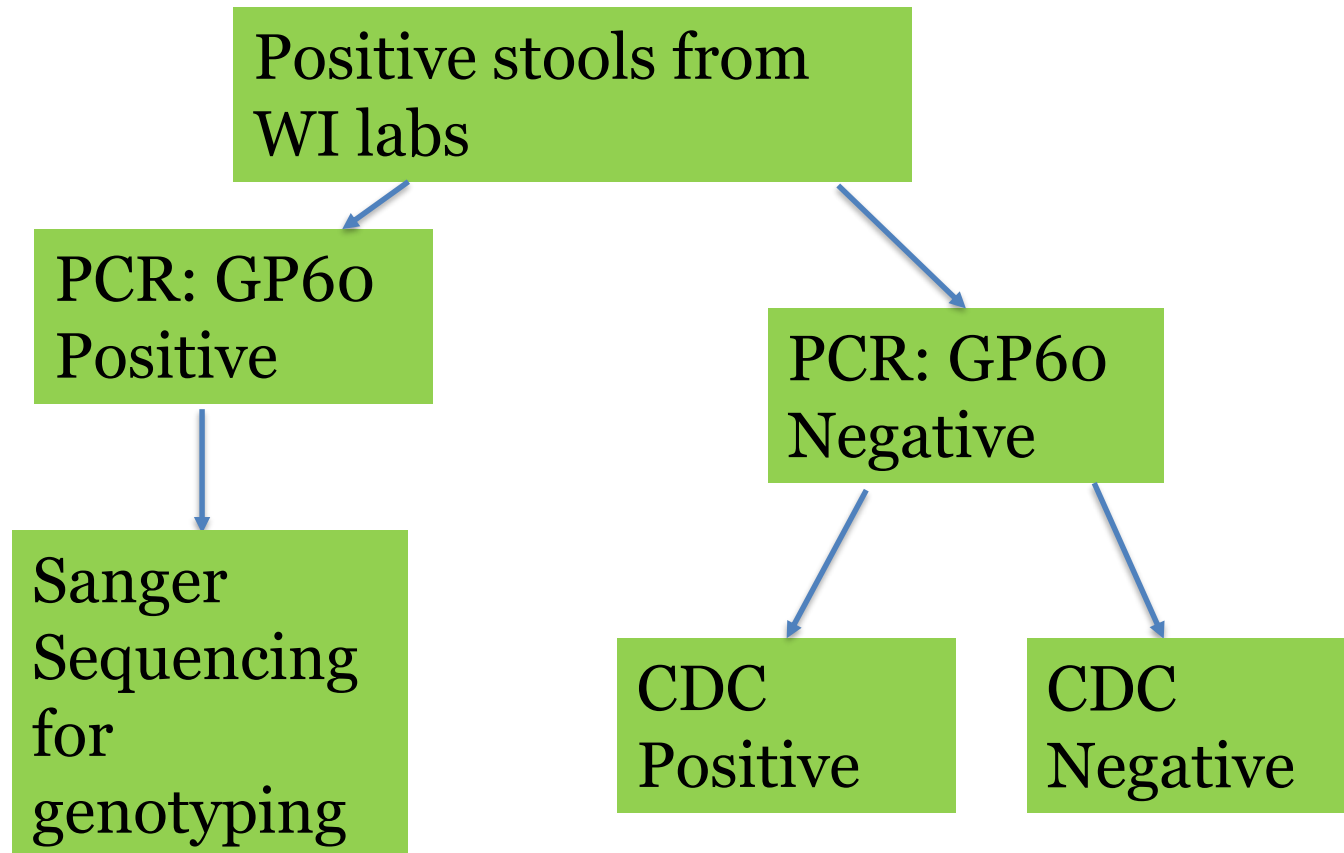
Sanger Sequencing

- Sanger sequencing is most ideal for short sequences and provides high accuracy.
- Sanger sequencing part of the gp60 gene is most commonly used for further discriminating *C. parvum* and *C. hominis* at WSLH.



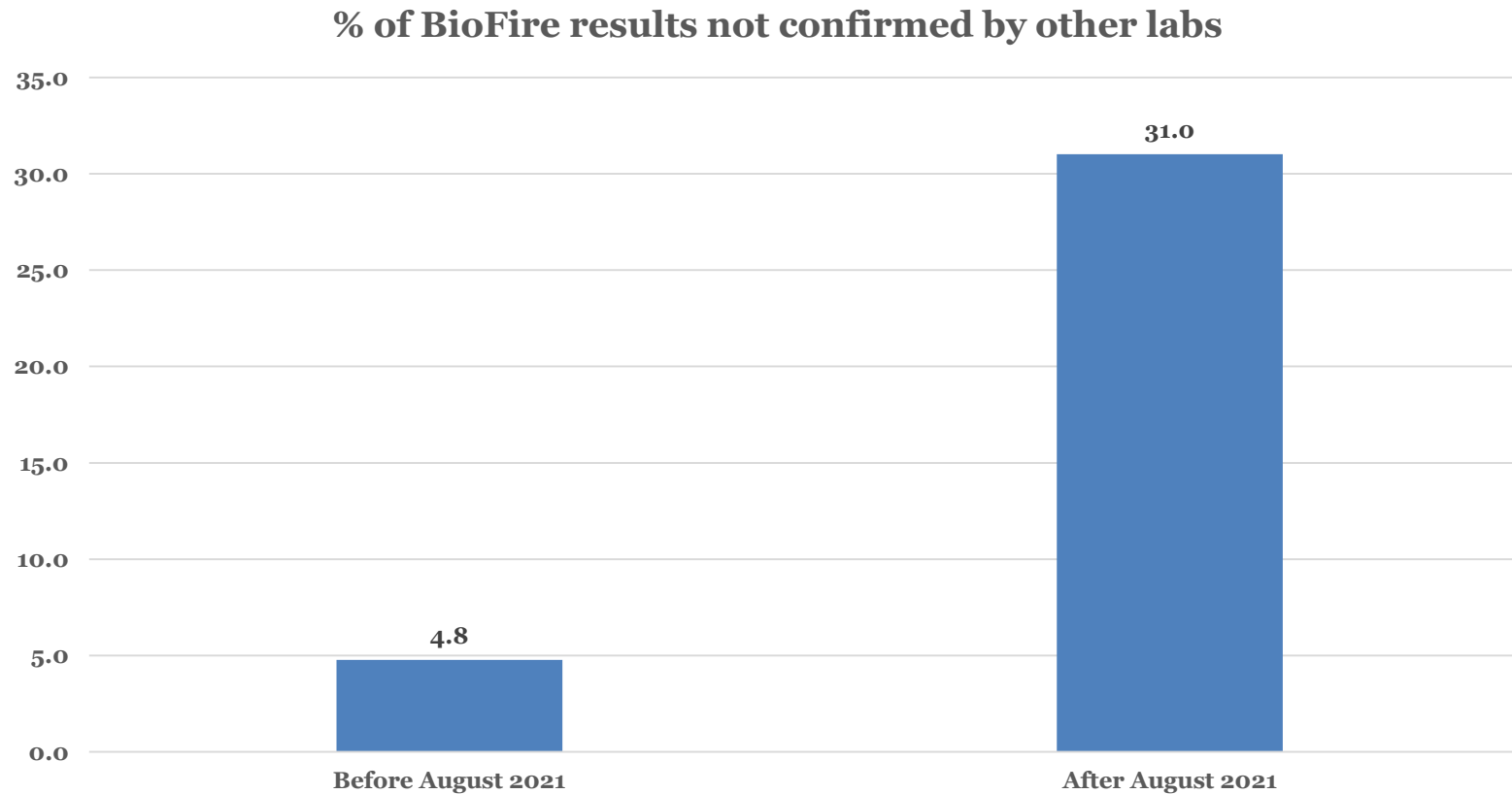


WSLH genotyping pipeline





Importance of Cryptosporidium surveillance to identify issues.





- We are working closely with CDC, CryptoNet labs, and the manufacturer.
- The Biofire false positive discovery would not be possible without the networking of state public health labs and clinical labs.
- We would like the clinical labs to engage in conversations with the manufacturer (Cary-Blair lot#/Biofire GI kit lot#/raw Biofire run files).
- We can take Biofire positive samples from labs for fee-exempt confirmatory DFA testing.

NGS Sequencing

- Next generation sequencing (NGS) is a high-throughput process that can rapidly sequence whole genomes of a species and allows us to study multiple regions of genome.
- NGS can be used to identify additional *Cryptosporidium* species and outbreak relatedness that could not be identified from Sanger sequencing.
- WSLH was chosen to CDC to validate *Cryptosporidium* NGS protocols.





Conclusion

- It is crucial for WI clinical labs to maintain good *Cryptosporidium* surveillance.
- The partnership of WSLH and WI clinical labs, and the voluntary submission of *Cryptosporidium* positive specimens for genotyping at WSLH, are a valuable asset to *Cryptosporidium* case and outbreak surveillance in Wisconsin.

Please send us more formalin-free residual stools for sequencing!





Questions?