

Monitoring of Human Pathogens and Source Identifiers for Discharges Across the United States: QMRA from Source to Bathing Site

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**Fecal Source Identification and Associated Risk
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Quantification of Pathogens and Sources of Microbial Indicators for QMRA in Recreational Waters

WERF Report PATH2R08

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Type And Geographical Area Of Monitoring Sites



7-month campaign, 3+1 distinct regions, 67 samples

The WERF “Toolkit”

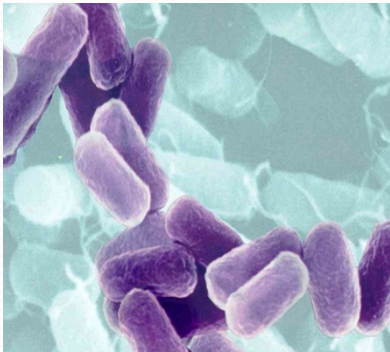
BACTERIA

V. cholerae

Campylobacter

Salmonella

V. parahaemolyticus

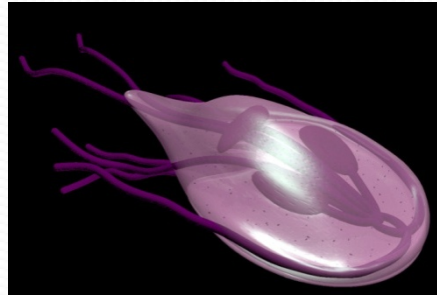


PROTOZOA

Giardia

Cryptosporidium

Toxoplasma



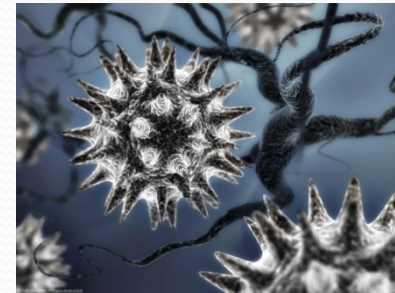
VIRUSES

Adenovirus

Enterovirus

Norovirus

Rotavirus



INDICATORS / SOURCE IDENTIFIERS

E. coli

Enterococcus

Bacteroidales (BacHum, BacCow, BacCan, BacUni)

Bird-associated assay (*Catellibacillus*)

The WERF “Toolkit”

RED = Priority Pathogens

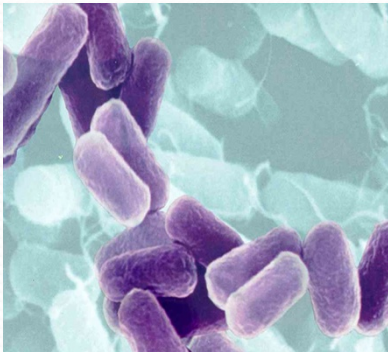
BACTERIA

V. cholerae

Campylobacter

Salmonella

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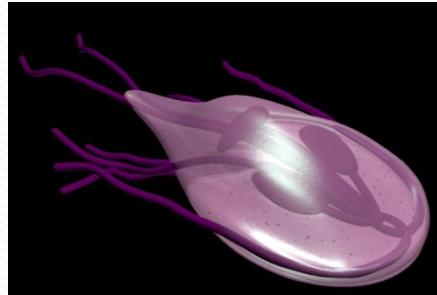


PROTOZOA

Giardia

Cryptosporidium

Toxoplasma



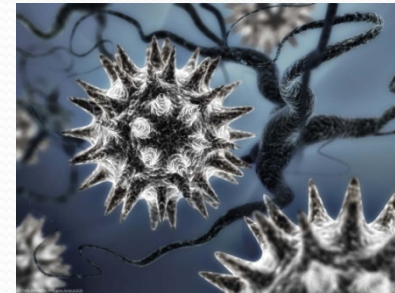
VIRUSES

Adenovirus

Enterovirus

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Rotavirus



INDICATORS / SOURCE IDENTIFIERS

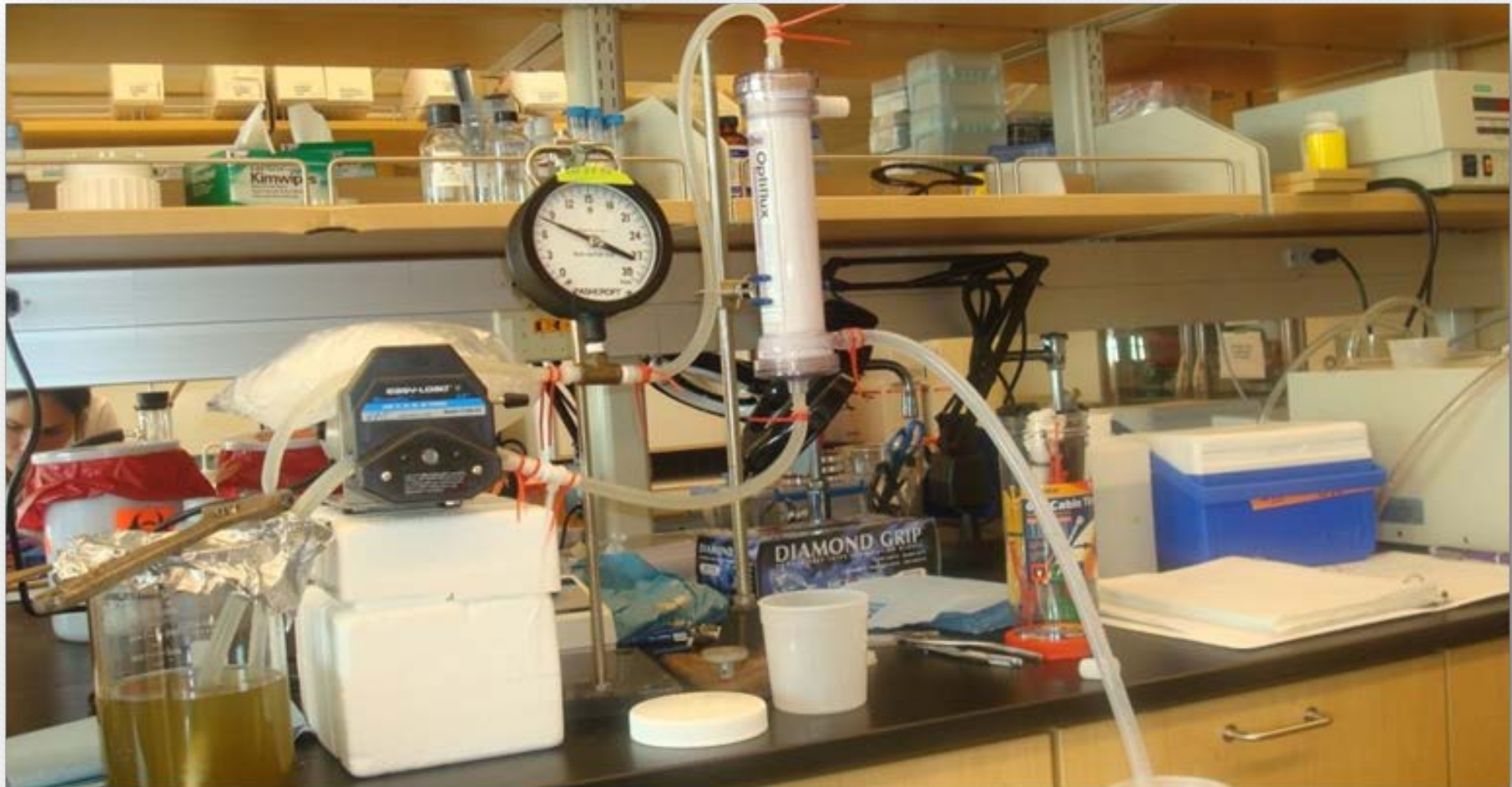
E. coli

Enterococcus

Bacteroidales (BacHum, BacCow, BacCan, BacUni)

Bird-specific assay (*Catellibacillus*)

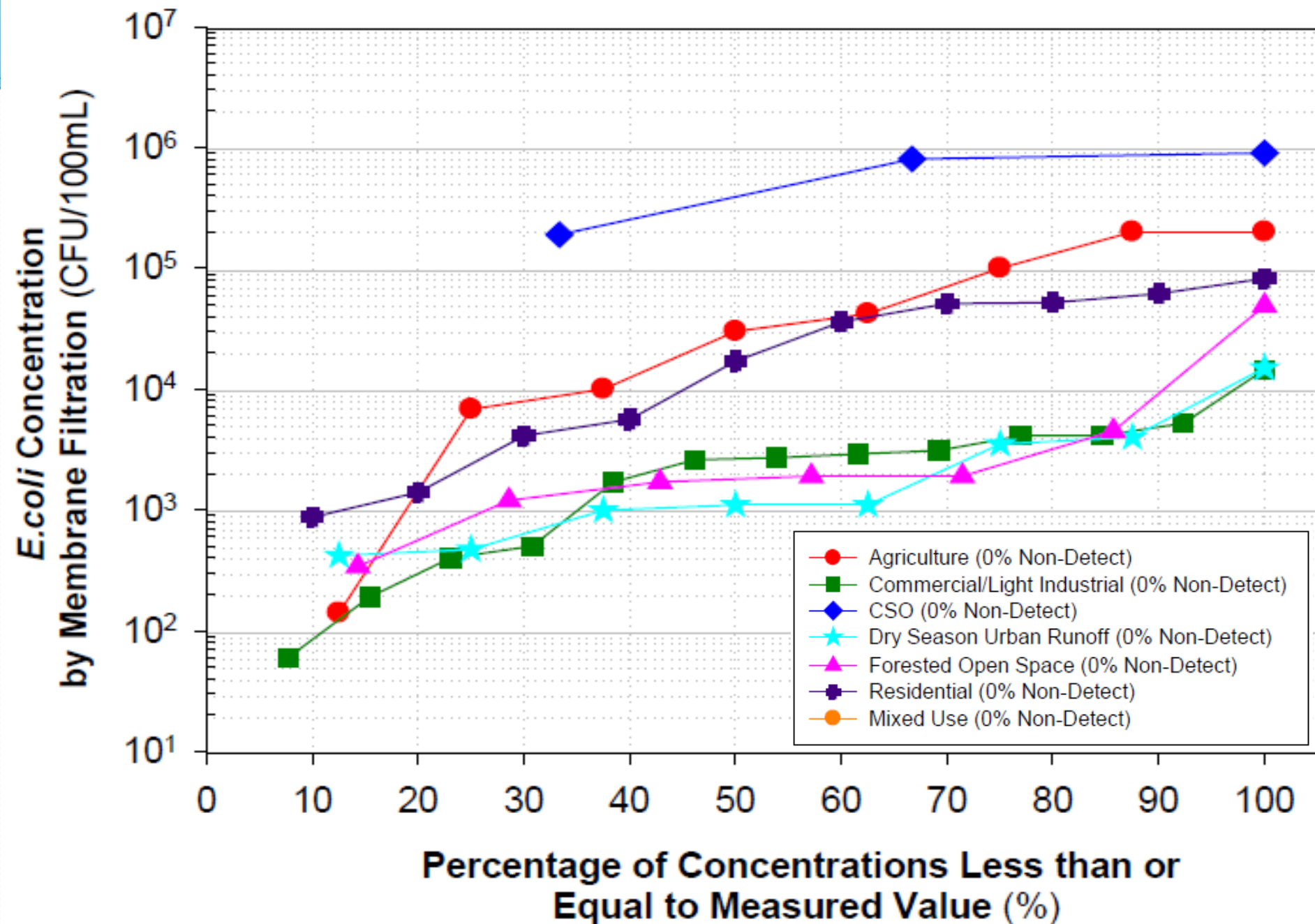
Experimental Set Up Using Fresenius Hollow-Fiber Ultrafiltration (HFF) Single-Use Cartridges



Monitoring Results: Pathogens

- Norovirus, Rotavirus, and Adenovirus 40/41 were more prevalent than *Cryptosporidium*, *Giardia* and pathogenic bacteria in stormwater or natural runoff.
- Norovirus GI and GII were reasonable predictors of other pathogens like Adenovirus 40/41, Adenovirus C, *Giardia* and *Cryptosporidium*.

E.coli Concentrations by Membrane Filtration



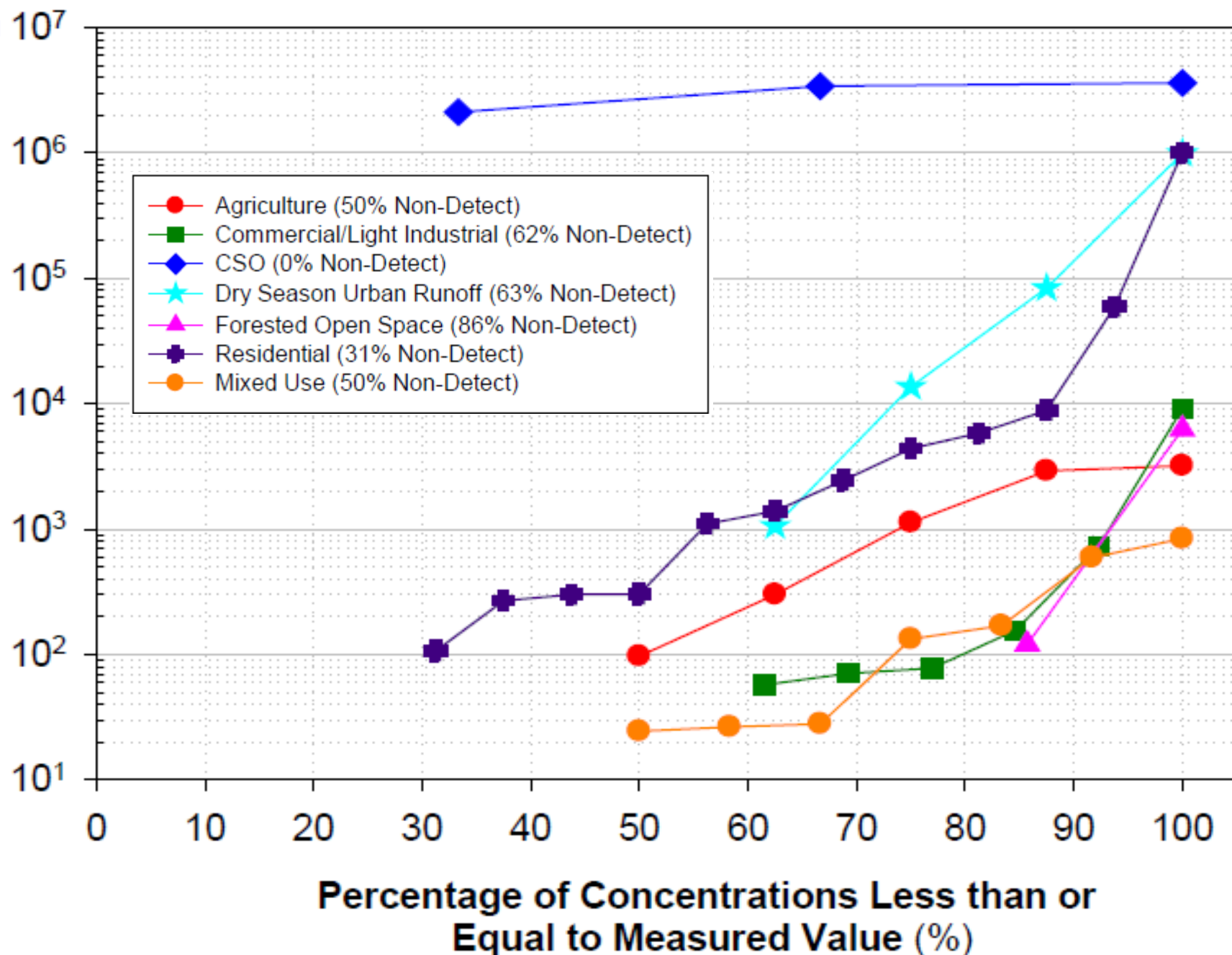
Bac

Hum

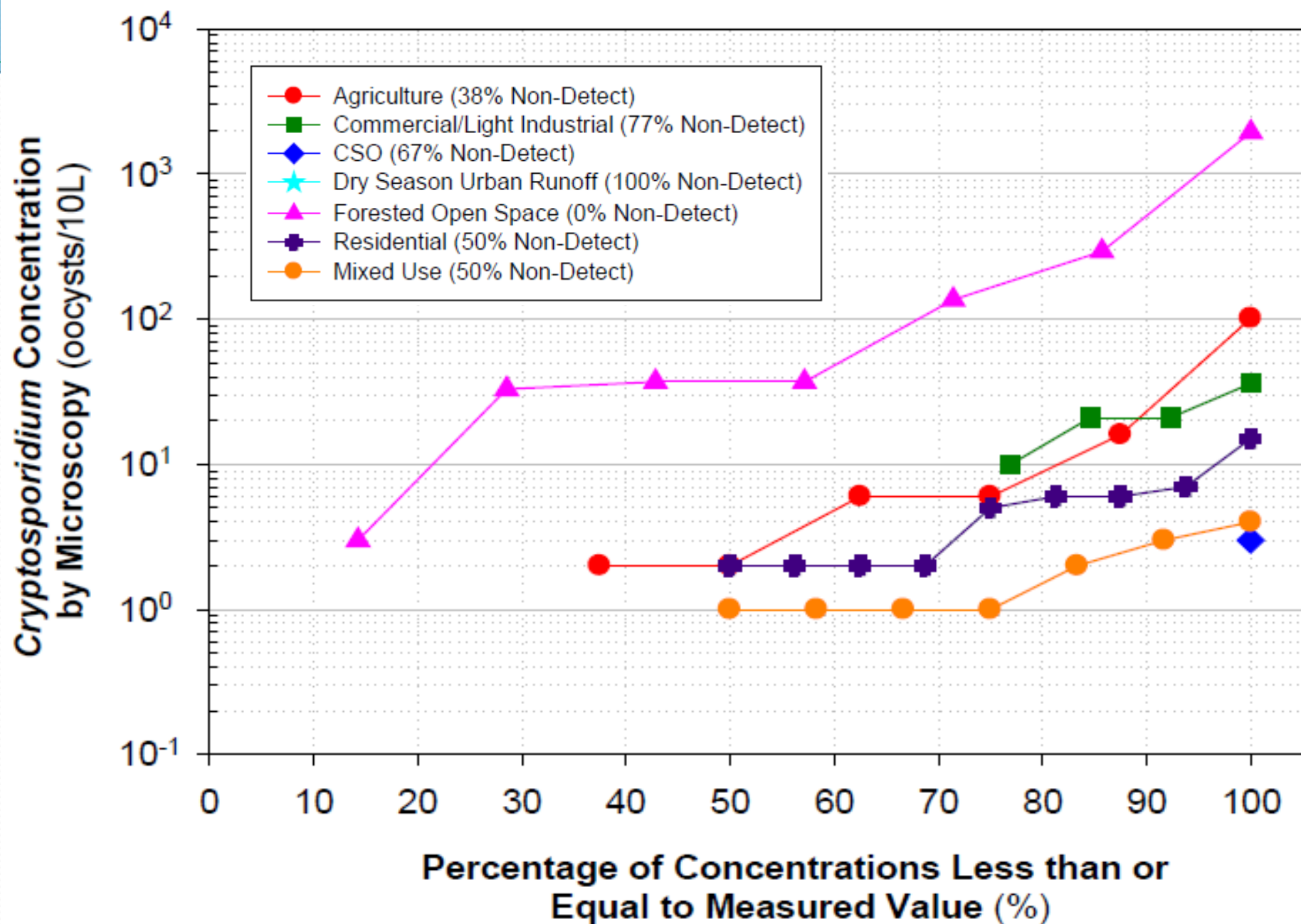
Human *Bacteroidales* Concentrations by qPCR

Human *Bacteroidales*
Concentration by qPCR (gc/mL)

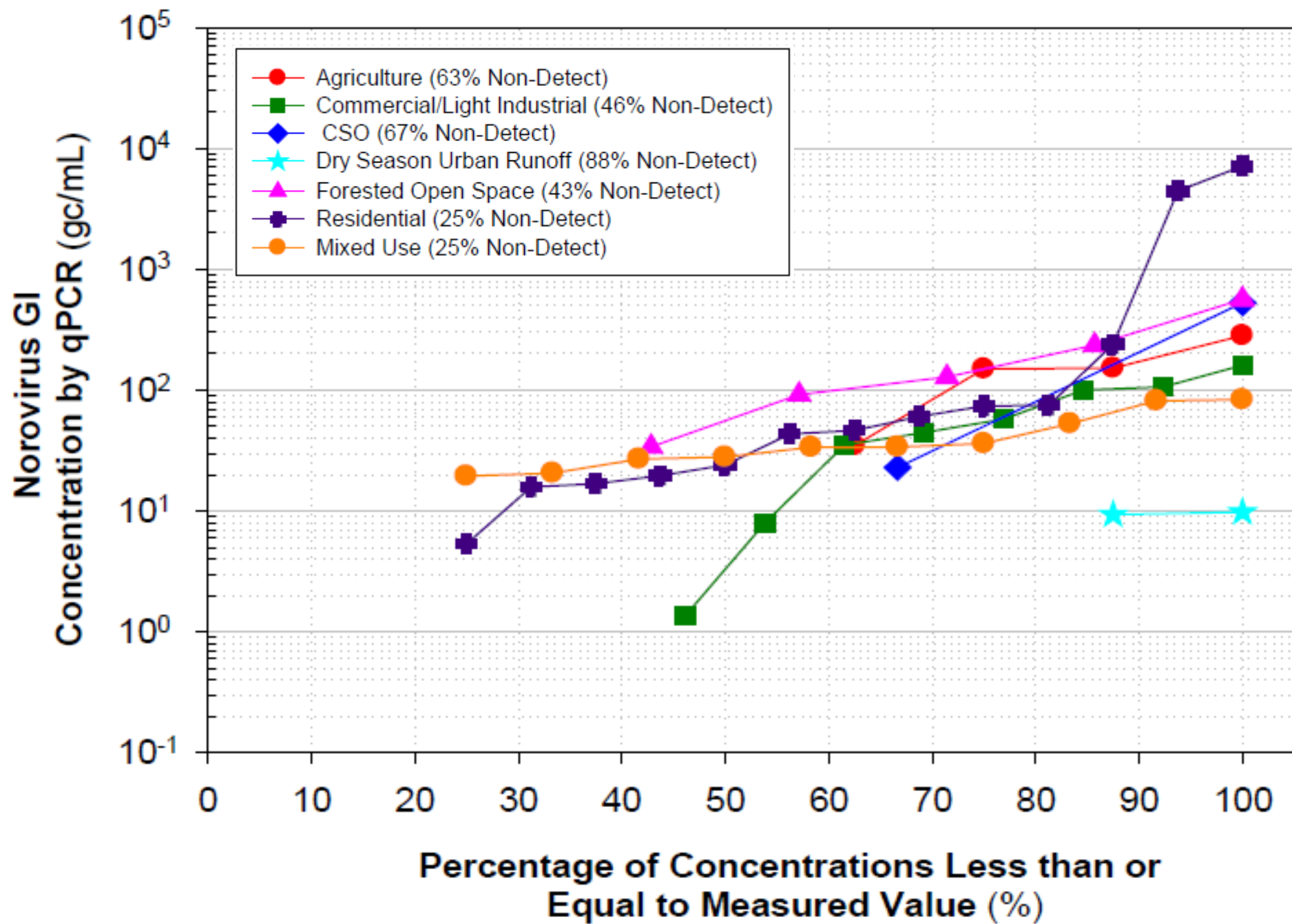
- Agriculture (50% Non-Detect)
- Commercial/Light Industrial (62% Non-Detect)
- ◆ CSO (0% Non-Detect)
- ★ Dry Season Urban Runoff (63% Non-Detect)
- ▲ Forested Open Space (86% Non-Detect)
- ◆ Residential (31% Non-Detect)
- Mixed Use (50% Non-Detect)



Cryptosporidium Concentrations by Microscopy

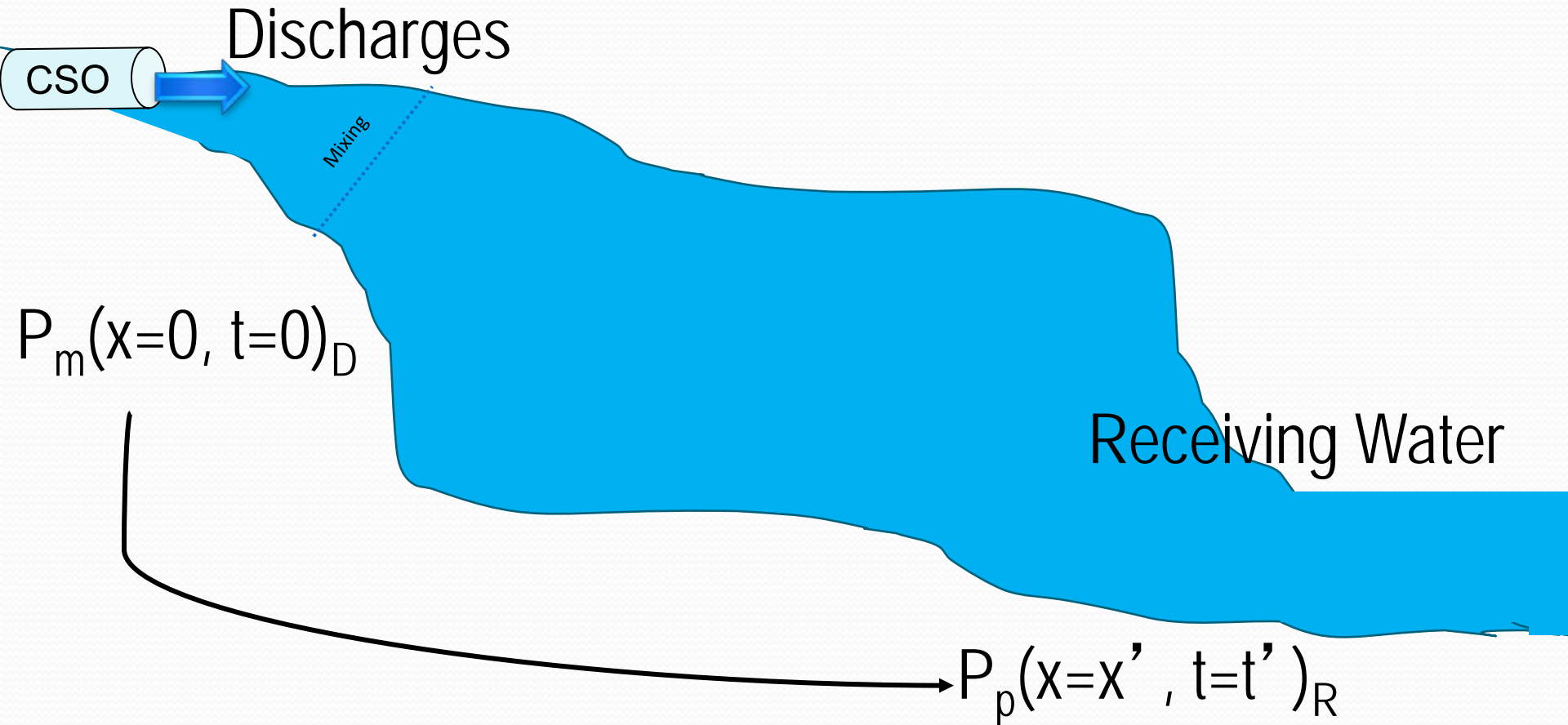


Norovirus GI Concentrations by qPCR



Step 1: Discharges \longrightarrow Receiving Water

$$P_m(x=0, t=0)_D \longrightarrow P_p(x=x', t=t')_R$$



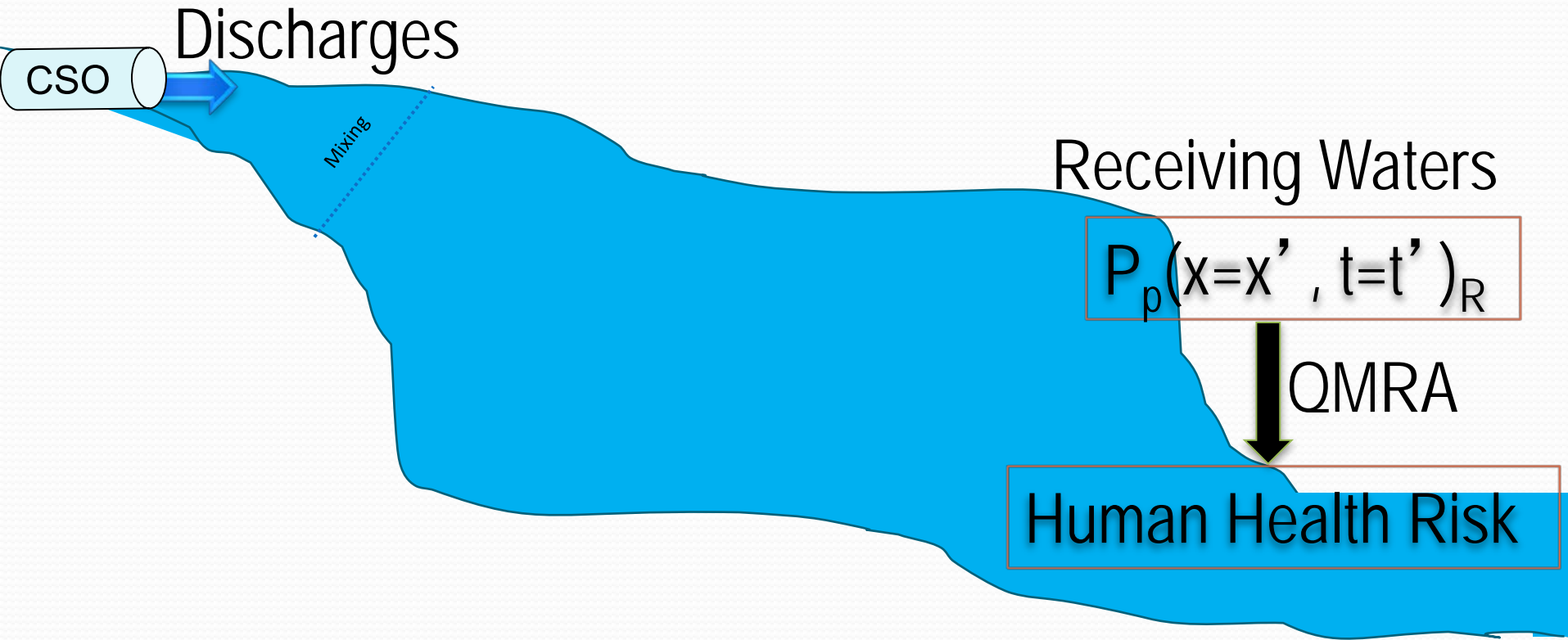
P_p = Predicted Pathogen Conc.

P_m = Measured Pathogen Conc.

x = Distance, t = Time, D = Discharge, R = Receiving Water

Step 2: Prediction of Health Risk

$P_p(x=x', t=t')_R \xrightarrow{\text{QMRA}}$ Human Health Risk

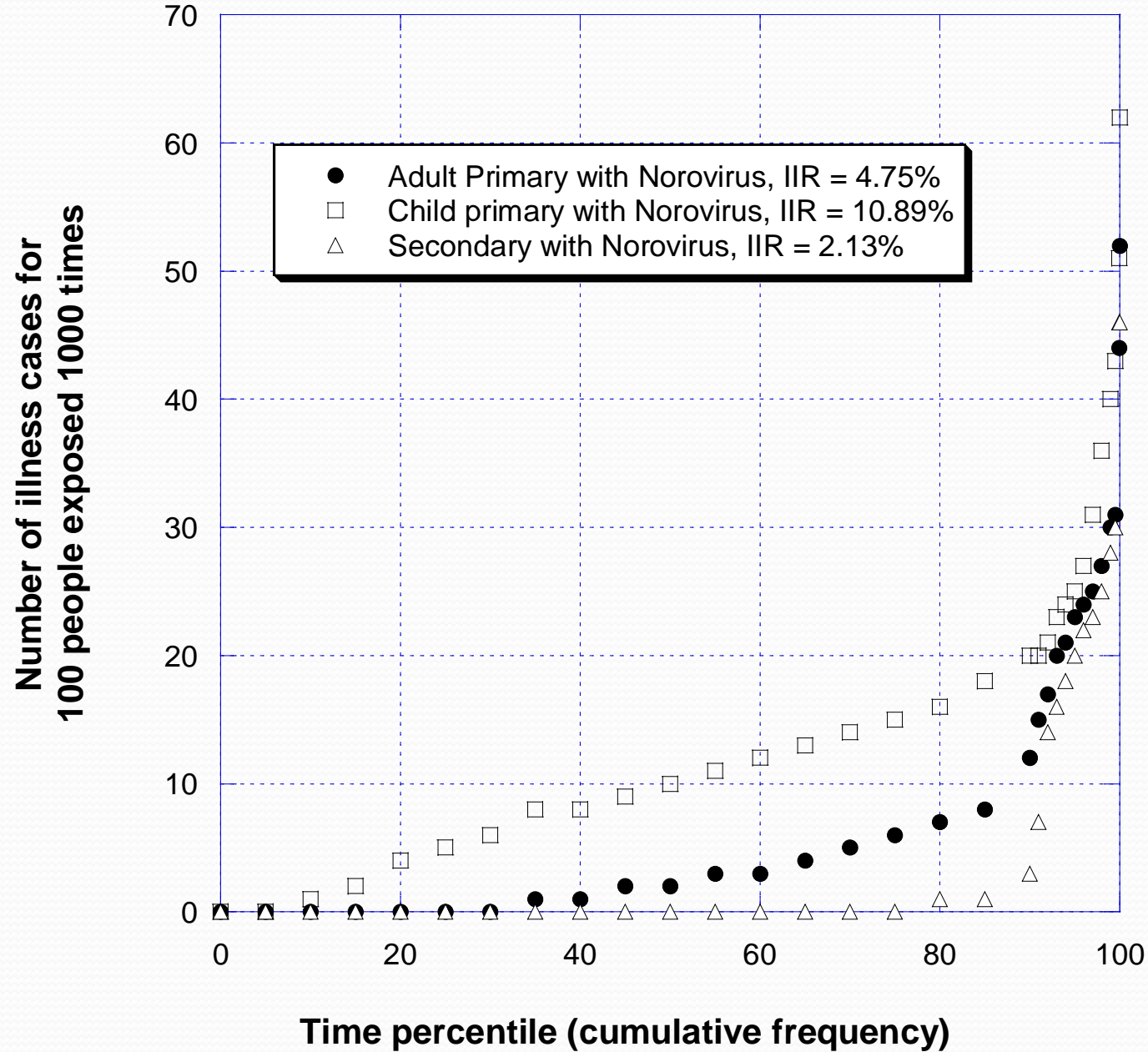


P_p = Predicted Pathogen Conc.; P_m = Measured Pathogen Conc.
 x = Distance; t = Time; D = Discharge; R = Receiving Water

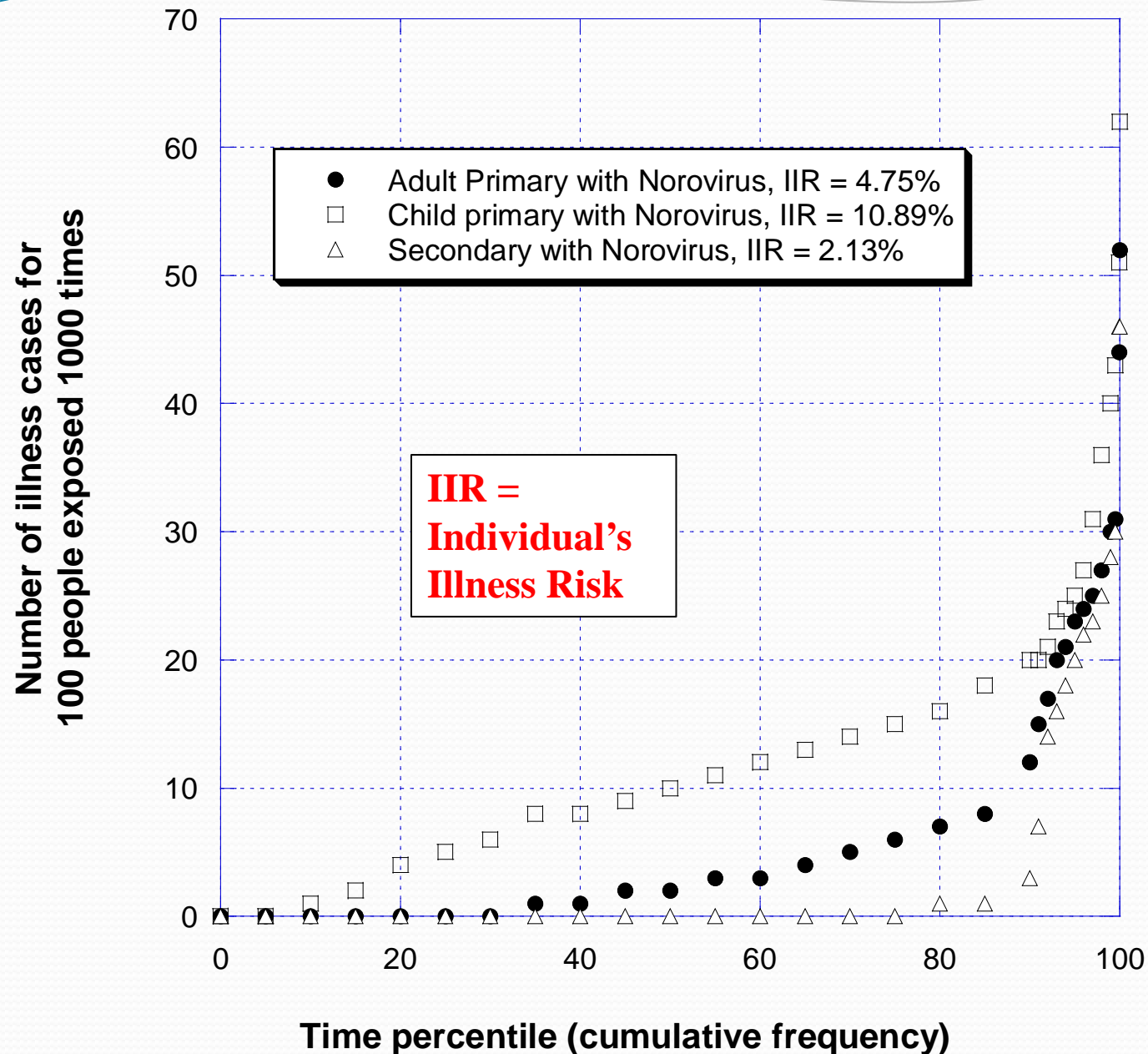
Example: Application of Discharge-based QMRA (Simplified)

- **Assumption #1 - Relative contribution of discharges-of-concern:** Four discharges were evenly “composited”, meaning a 25% by-volume contribution for each of the following: Residential stormwater, Commercial stormwater, Agricultural stormwater, and Forested Open Space stormwater.
- **Assumption #2 - Rates of dilution of discharges by other inputs:** A dilution rate of 30:1 was applied to discharges-of-concern.
- **Assumption #3 - Effects of dry versus wet conditions:** The assessed condition was wet weather and dry weather is not considered.
- **Assumption #4 - Travel times and pathogen decay rates:** While pathogen decay/inactivation is likely an important consideration for most recreational sites, this example assumes zero decay.

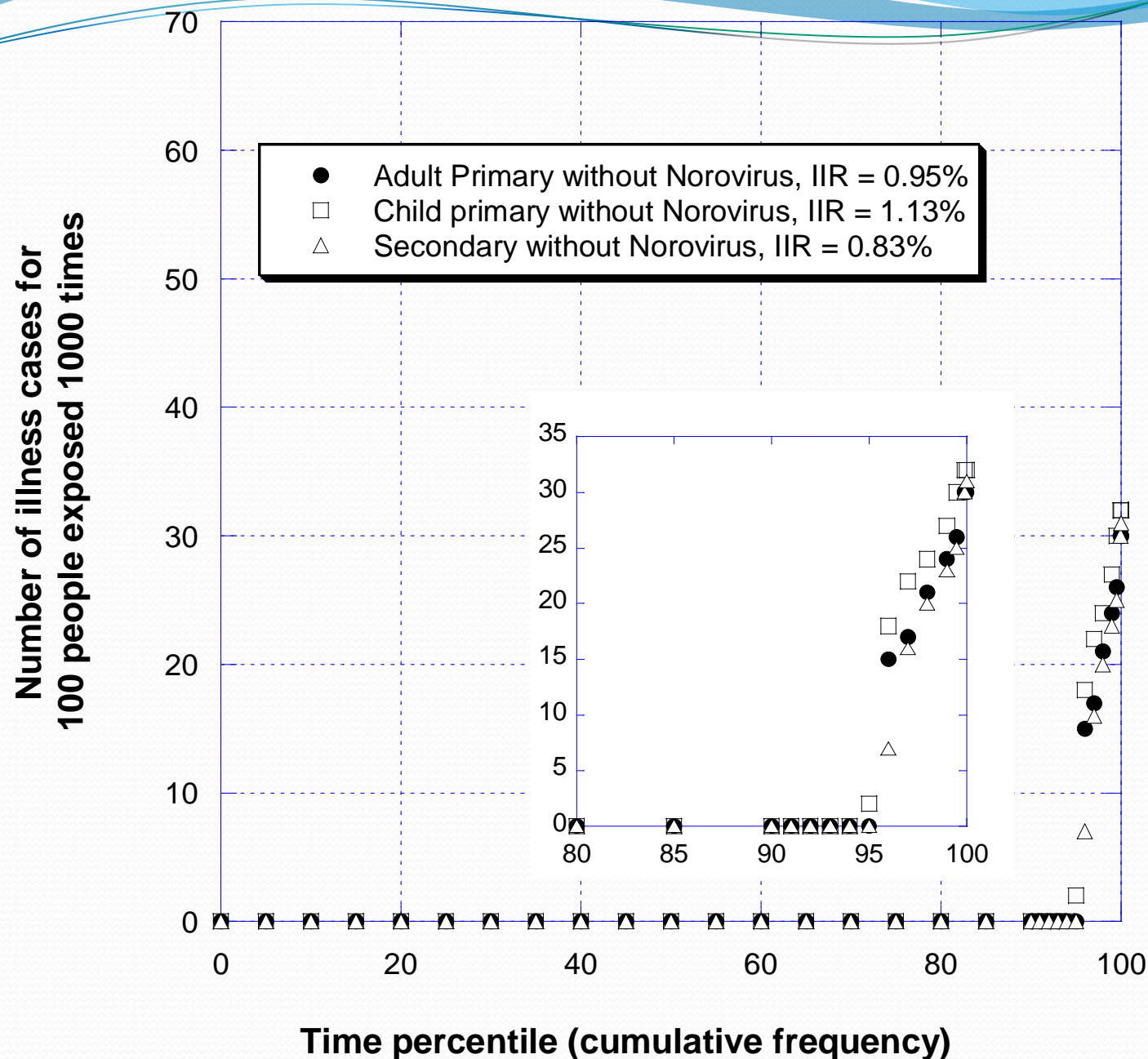
Hypothetical: Recreational Risks at an Impacted Site



Hypothetical: Recreational Risks at an Impacted Site



Risk with Norovirus Excluded from the Calculations



Conclusions

- **Norovirus** is the most dominant health risk.
- **Rotavirus** generally induced the second-highest incidence of risk among the tested pathogens.
- Can investigate relative risks from different **discharges** and for different densities and loadings
- Can use this QMRA approach for **beaches** if good knowledge of main discharges

Students and Collaborators

<u>Wuertz lab</u> Dr. Alexander Schriewer Dan Wang Arti Kundu Asma Rizva	<u>UC Vet Med (Miller lab)</u> Aiko Adell Nadira Chouicha Ann Melli Prof. Barbara Byrne Prof. Pat Conrad
<u>AMEC Earth & Environmental</u> Candice Owen	<u>NIWA, New Zealand</u> Dr. Rebecca Stott

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