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**Title:** Monitoring SARS-CoV-2 in wastewater.

**Author(s):** Urbatsch, Dana Deann

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# Monitoring SARS-CoV-2 in wastewater.

Dana Urbatsch B-10

# Abstract

The monitoring of SARS-CoV-2 through individual testing is subject to inaccuracies. Wastewater biosurveillance has been used around the world to detect the presence of drugs, viruses, and other chemicals present in a population. Fecal shedding of SARS-CoV-2 implies that wastewater can be analyzed in order to detect varying levels of the virus within a given community. Pepper Mild Mottle Virus is present consistently in human fecal matter, and can be used to normalize the amount of SARS-CoV-2 detected. In this experiment, the amount of SARS-CoV-2 in the wastewater sample was compared to the amount of PMMoV present. Wastewater samples were separated into three fractions: direct wastewater, a solid pellet, and filter paper extract. An RT-qPCR test was used on the extracted RNA from each fraction to detect the relative amount of SARS-CoV-2 and PMMoV copies in each sample. The amount of SARS-CoV-2 was successfully normalized to the amount of PMMoV.

# Improvements in SARS-CoV-2 Detection

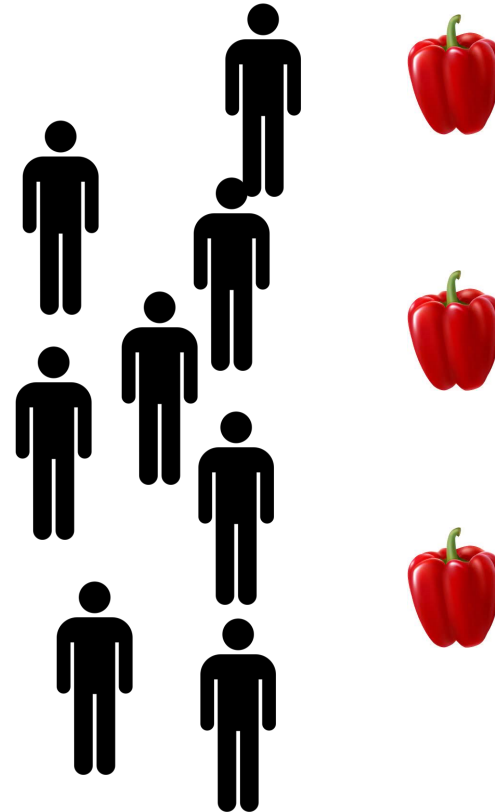
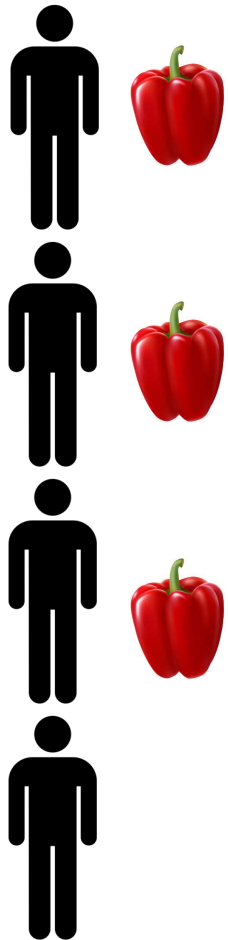
- Wastewater monitoring has a variety of applications
- People shed the virus through fecal matter
- Wastewater monitoring can show fluctuations in the population over time
- Can supplement SARS-CoV-2 tracking



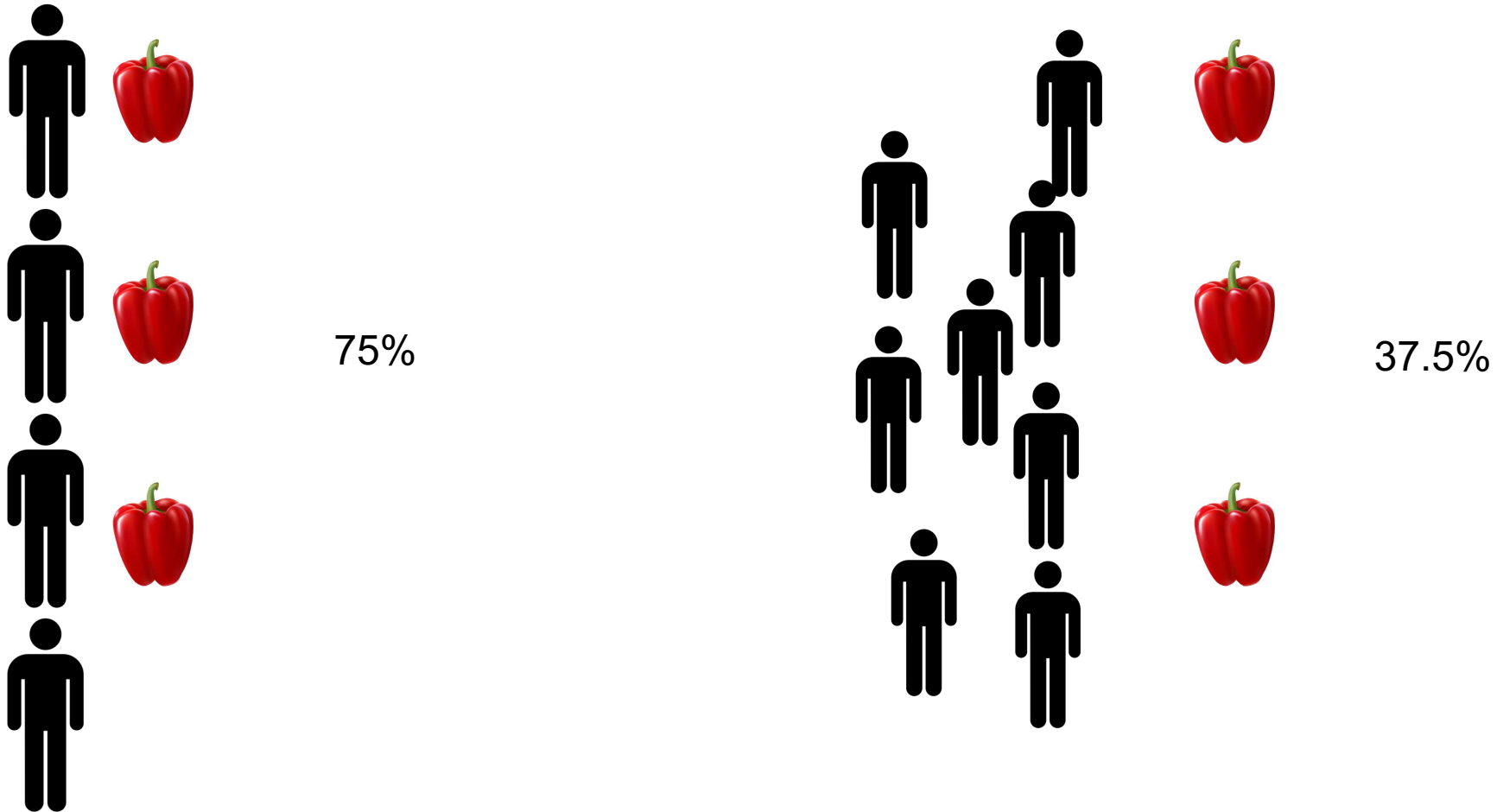
# Benefits of Wastewater Testing

- Noninvasive
- Anonymous
- Accounts for asymptomatic people
- Is not limited by shortages of tests
- Shows trends several days in advance of individual testing

# Normalizing SARS-CoV-2



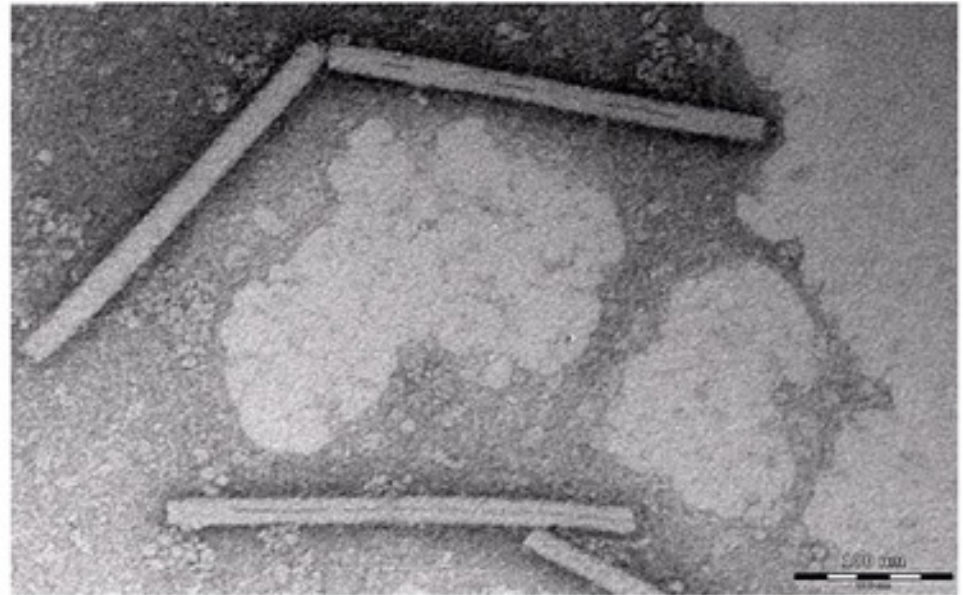
# Normalizing SARS-CoV-2





# Pepper Mild Mottle Virus (PMMoV)

- Can be consumed through peppers or products from peppers, like hot sauce
- Present in human feces, but not animal feces
- Used to show the amount of human feces



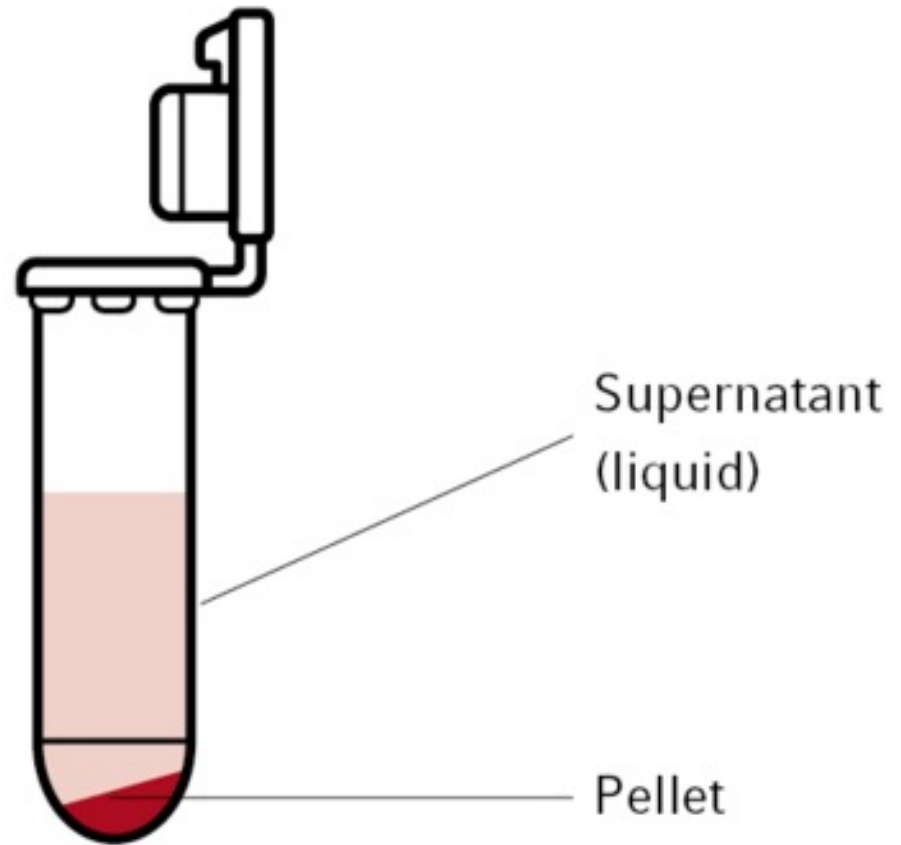
[https://upload.wikimedia.org/wikipedia/commons/4/4c/PMMoV\\_particles.png](https://upload.wikimedia.org/wikipedia/commons/4/4c/PMMoV_particles.png)

# Methodology

- Collection
- RNA Extraction
- RT-qPCR

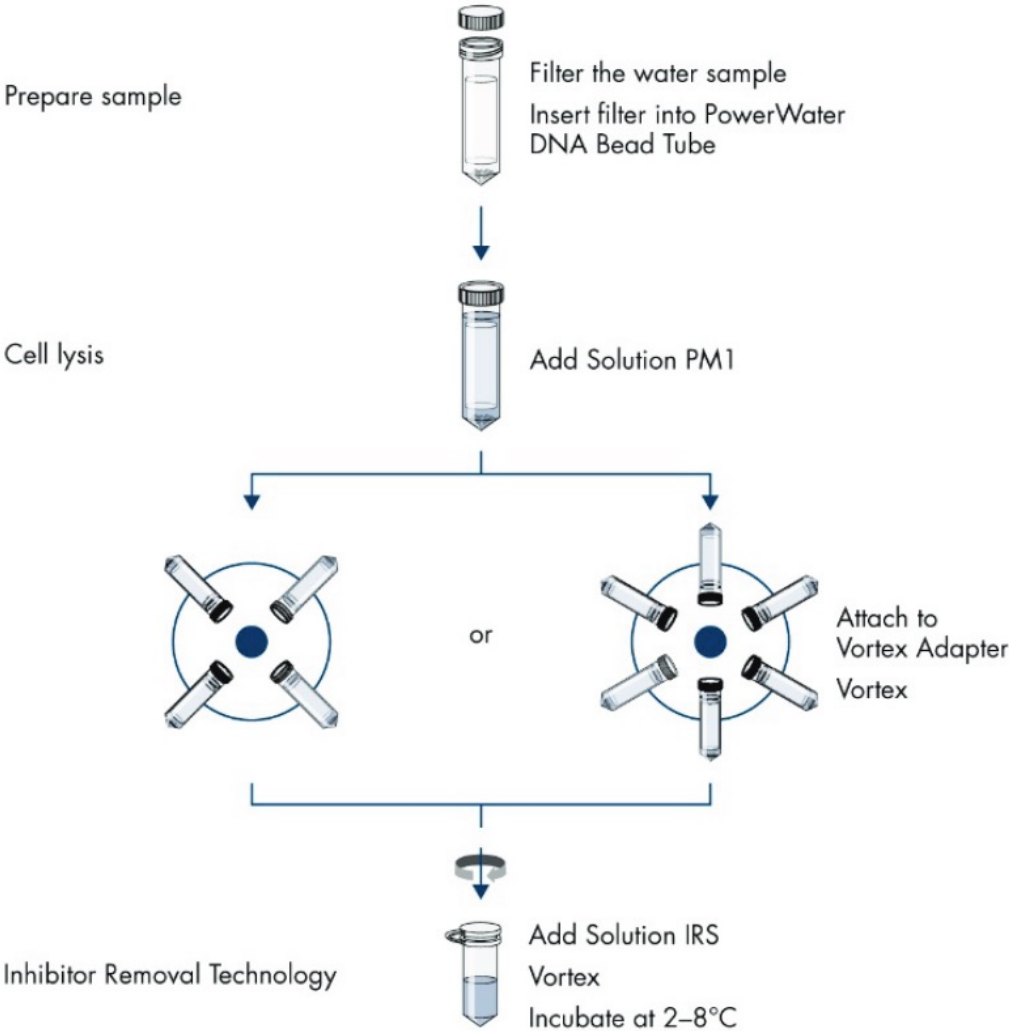
# Collection

- LANL Wastewater
- Collection days
- Fractions 1, 2, and 3

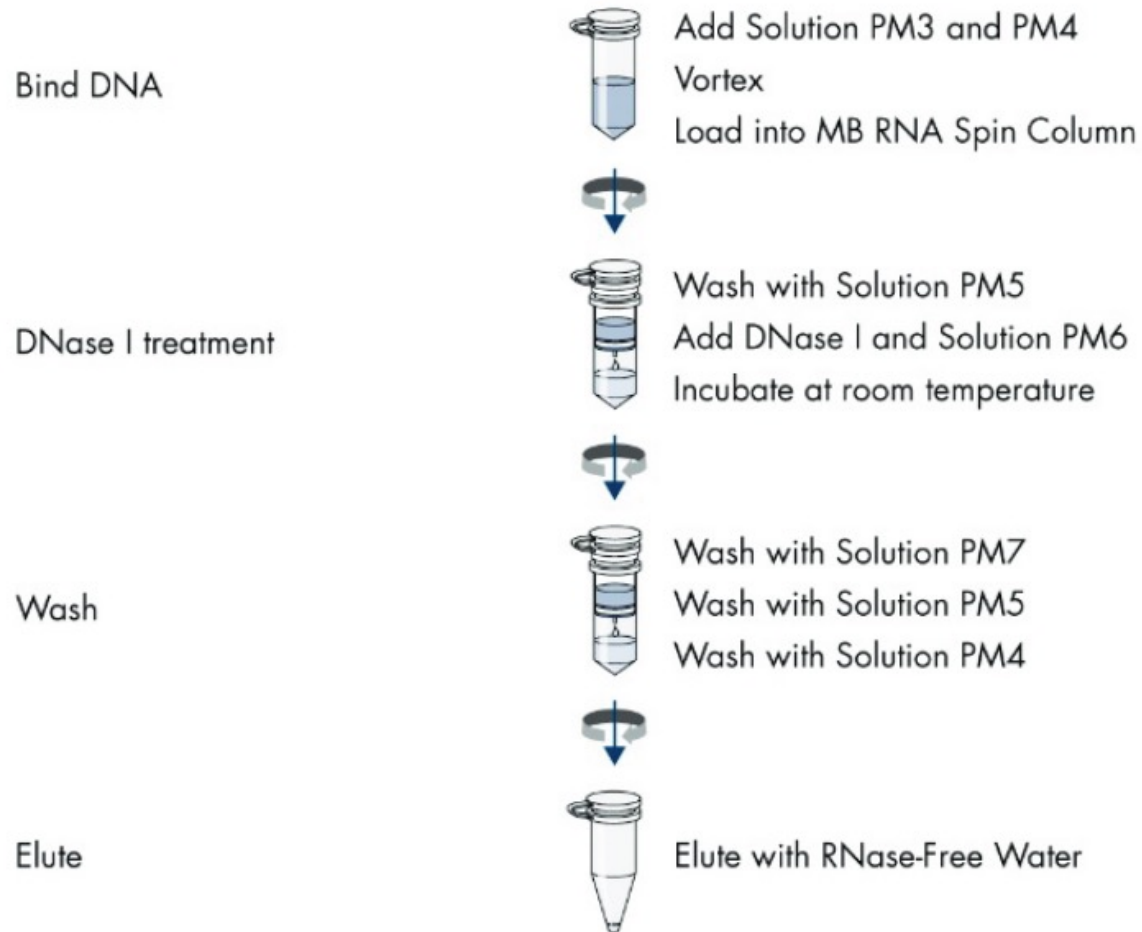


# RNA Extraction

## RNeasy PowerWater Kit Procedure

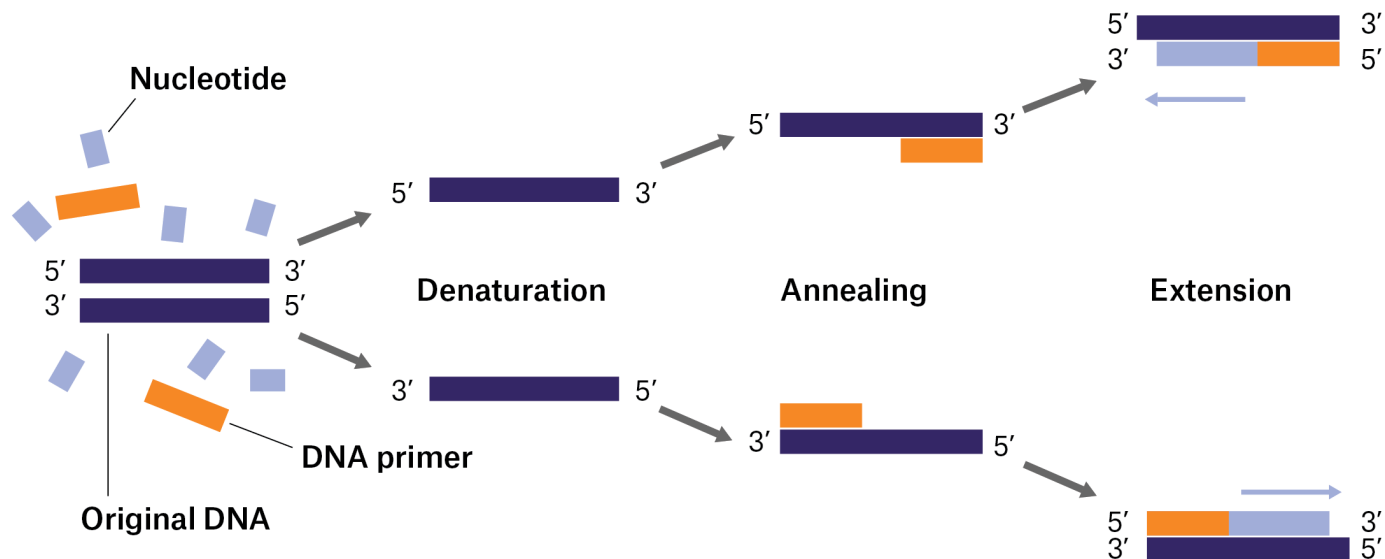


# RNA Extraction



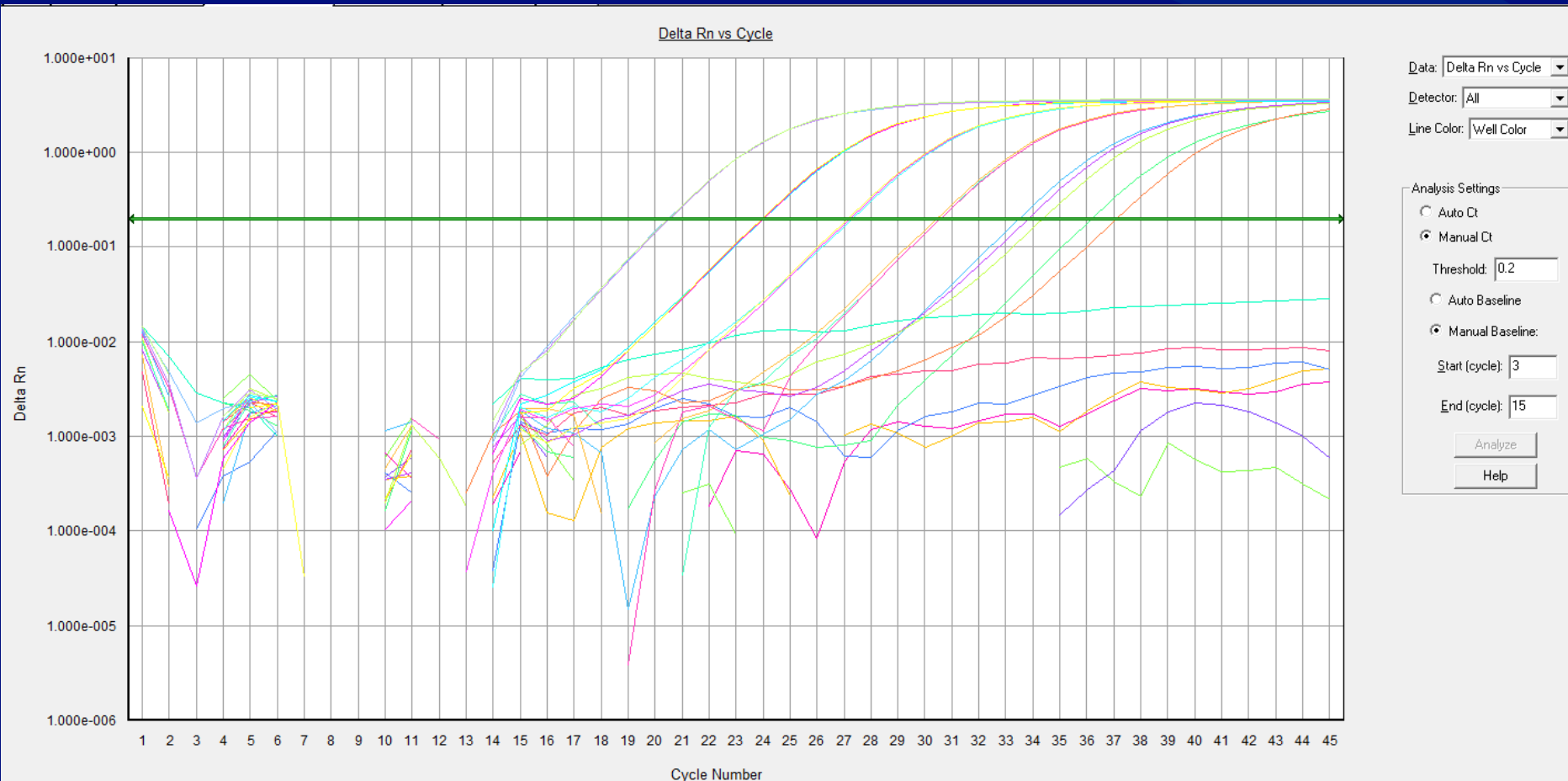
# RT-qPCR

- Reverse Transcription Quantitative Polymerase Chain Reaction
- Three Steps: Denaturing, Annealing, Extension
- TaqMan assay vs SYBR Green



©GA International

# Amplification Curve – June 9th



# Standard Curve – June 9th

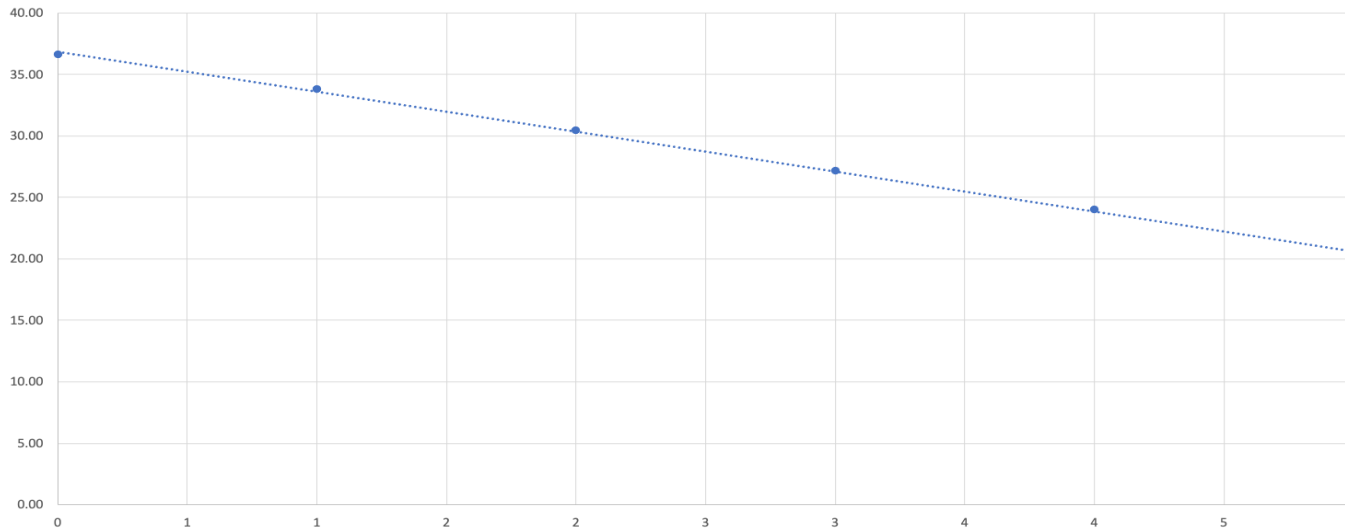
Ct Value

$$y = -3.2494x + 36.874$$

$$R^2 = 0.9991$$

$$y = -3.2494x + 36.874$$

$$R^2 = 0.9991$$



$$32 = -3.2494x + 36.874$$

$$-4.874 = -3.249x$$

$$x = 10^{1.5}$$

$$x = 31.6 \text{ copies/uL}$$

DNA Copies / uL

Log DNA Copies / uL

Ct Value

1

0

36.62

10

1

33.84

100

2

30.45

1000

3

27.17

10000

4

24.00

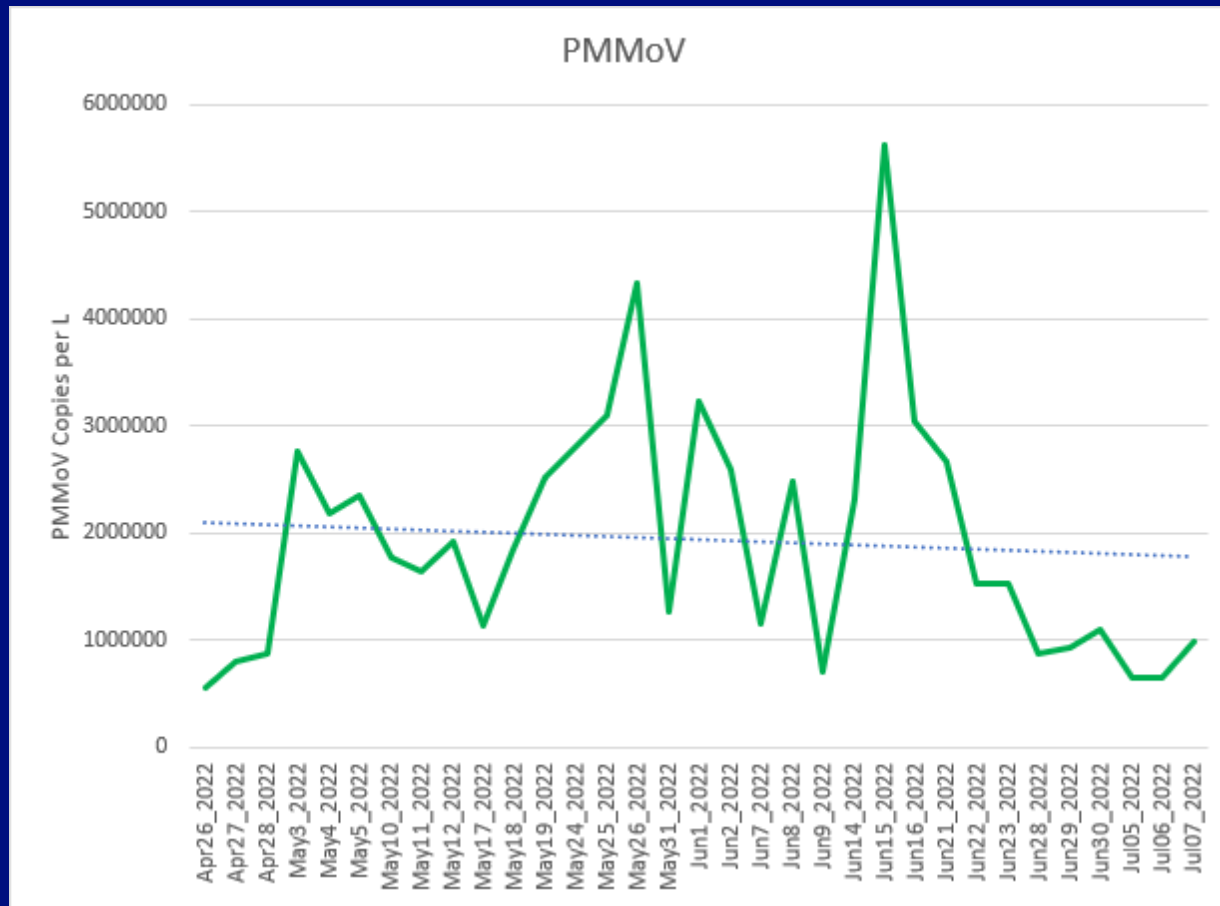
100000

5

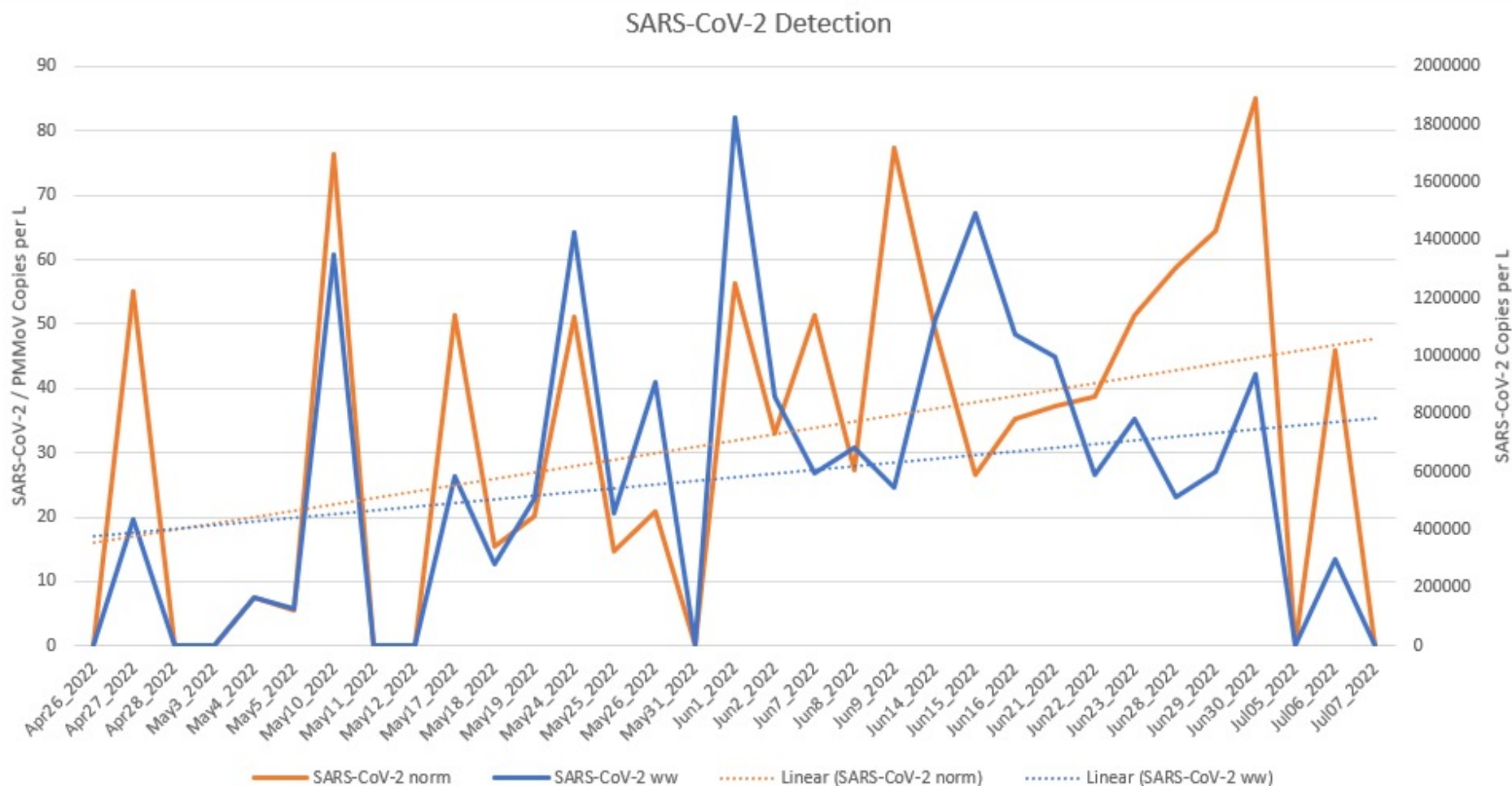
20.43



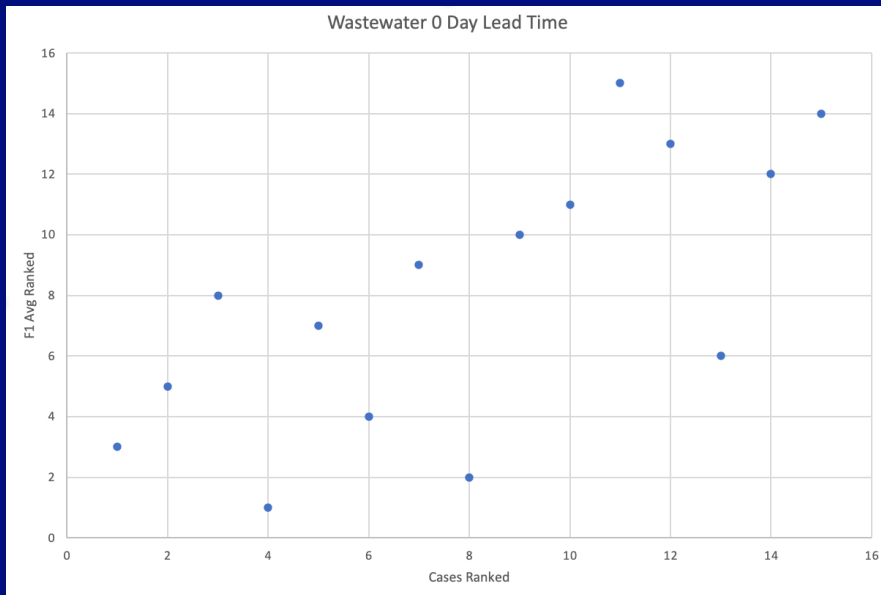
# PMMoV Graph



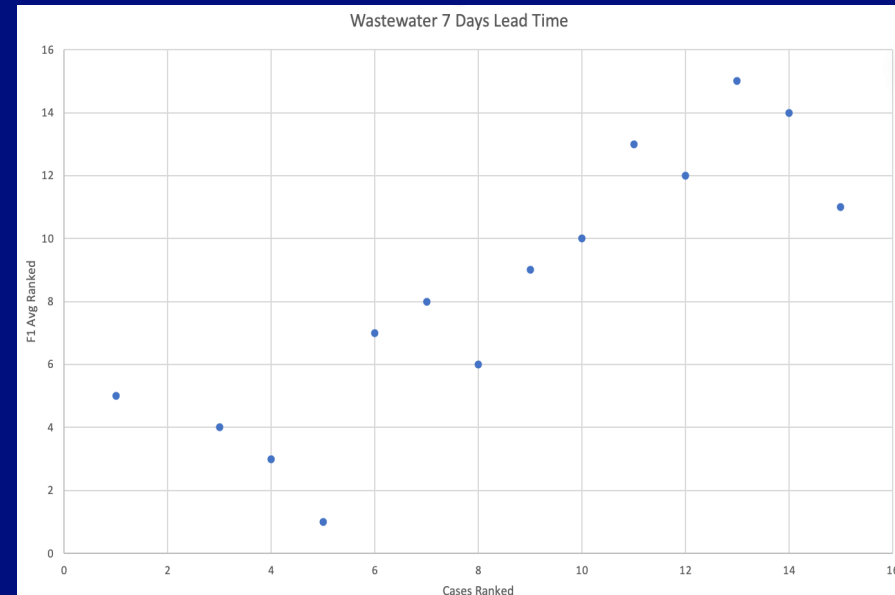
# SARS-CoV-2 Detection Graph



# Wastewater Detection and Individual Testing Correlation



Spearman Correlation Wastewater 0 Day  
Lead Time= 0.7



Spearman Correlation Wastewater 7 Day  
Lead Time= 0.86745

# Summary and Future Applications

- Wastewater monitoring
  - Other applications
  - Important for SARS-CoV-2 monitoring
  - Could show earlier trends
  - Widespread implementation
- Normalization using PMMoV to ensure more accurate monitoring

## Acknowledgements

- Andy Hatch and Shawn Starkenburg
- Department of Energy

**Thank You!**

**Questions?**