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Title: Monitoring SARS-CoV-2 in wastewater.

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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

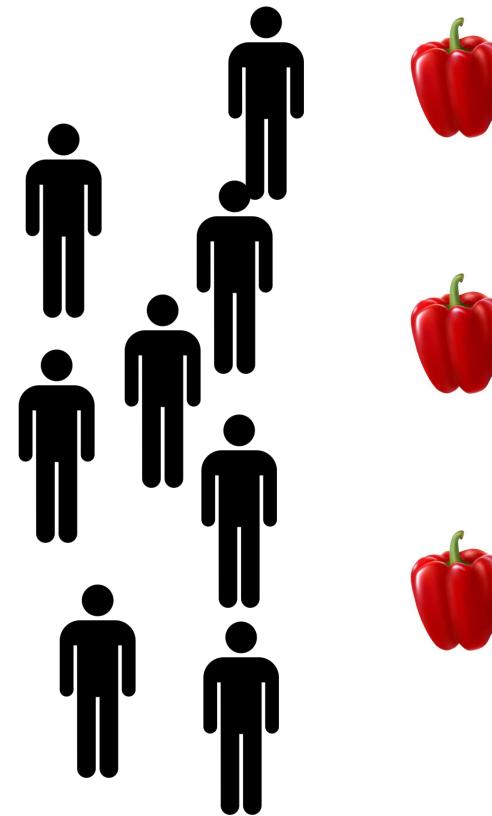
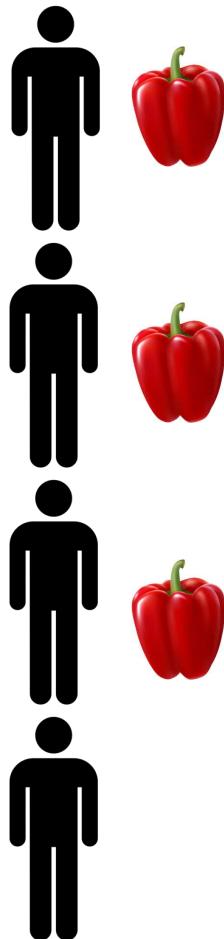


<https://www.science.org/doi/10.1126/science.abd0700>

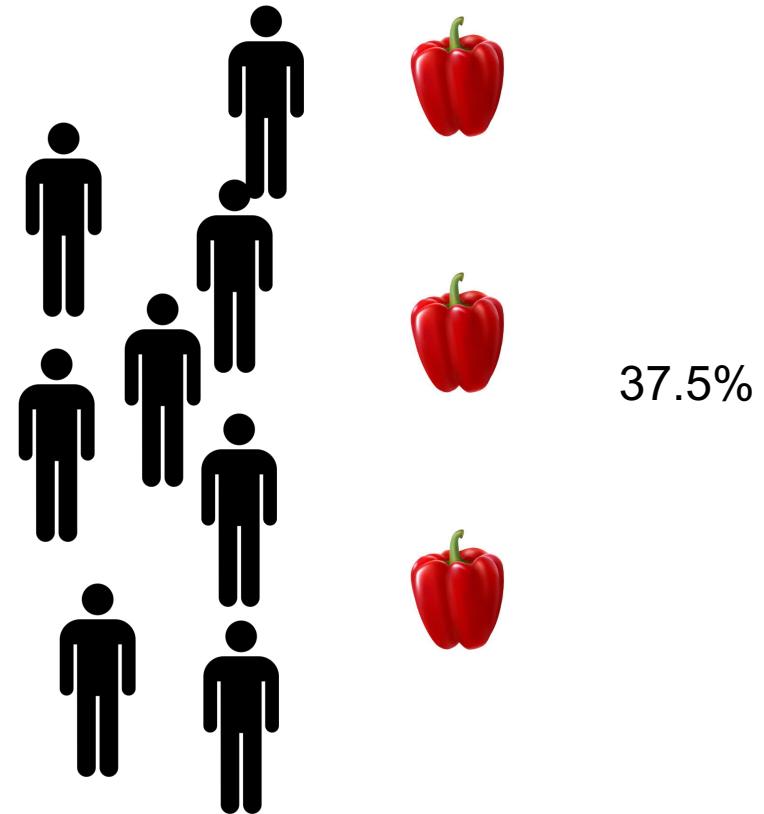
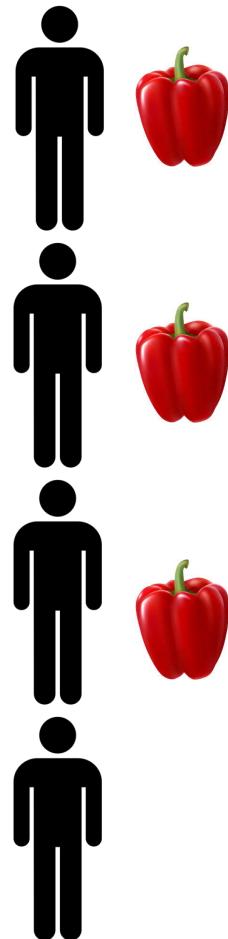
Benefits of Wastewater Testing

- 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

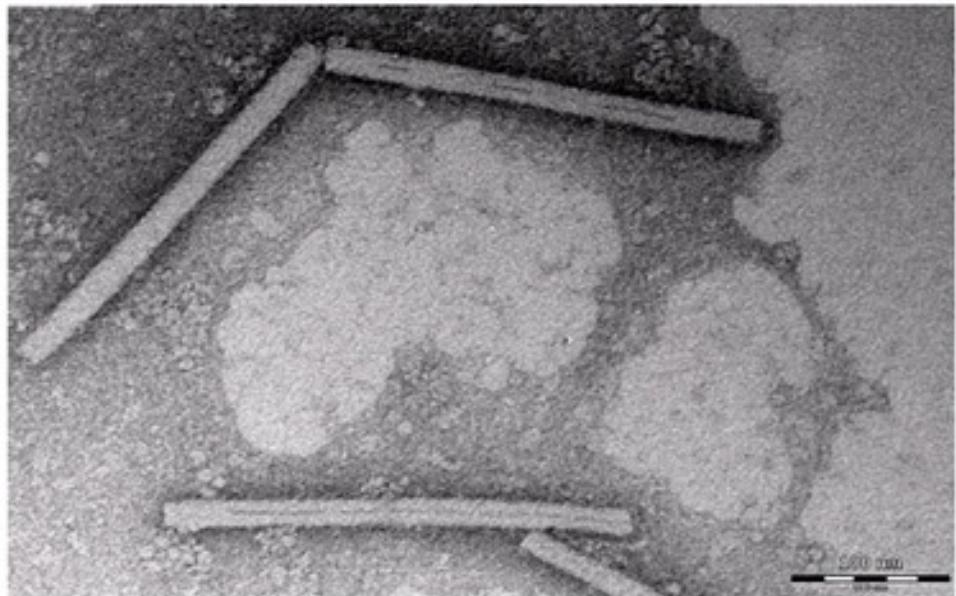


Normalizing SARS-CoV-2



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



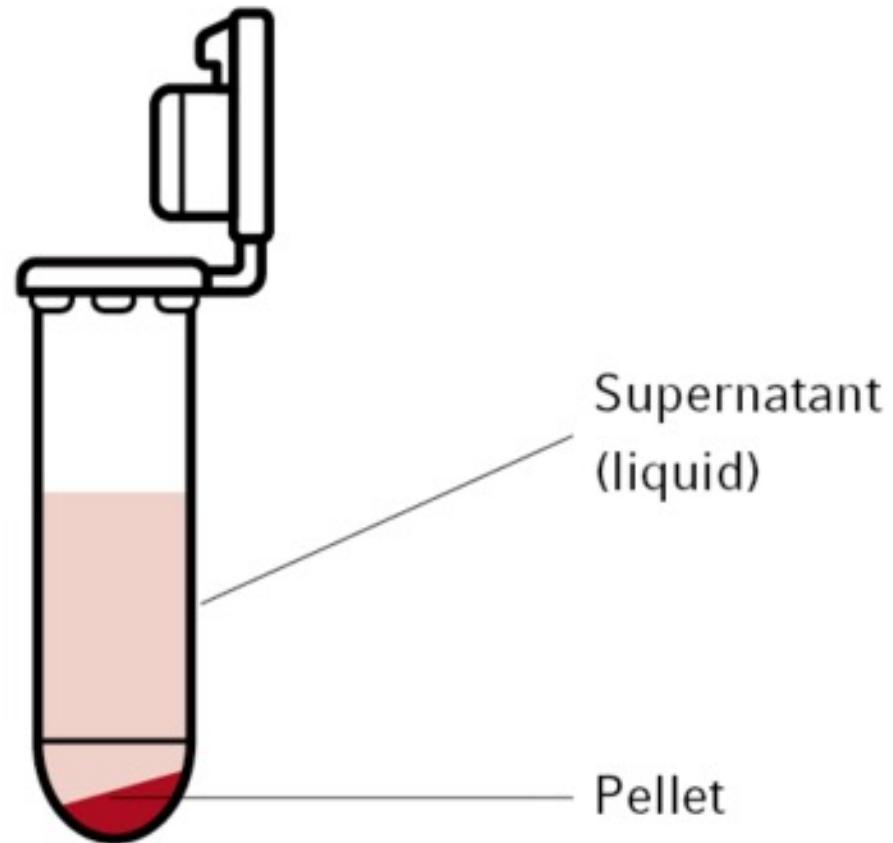
https://upload.wikimedia.org/wikipedia/commons/4/4c/PMMoV_particles.png

Methodology

- Collection
- RNA Extraction
- RT-qPCR

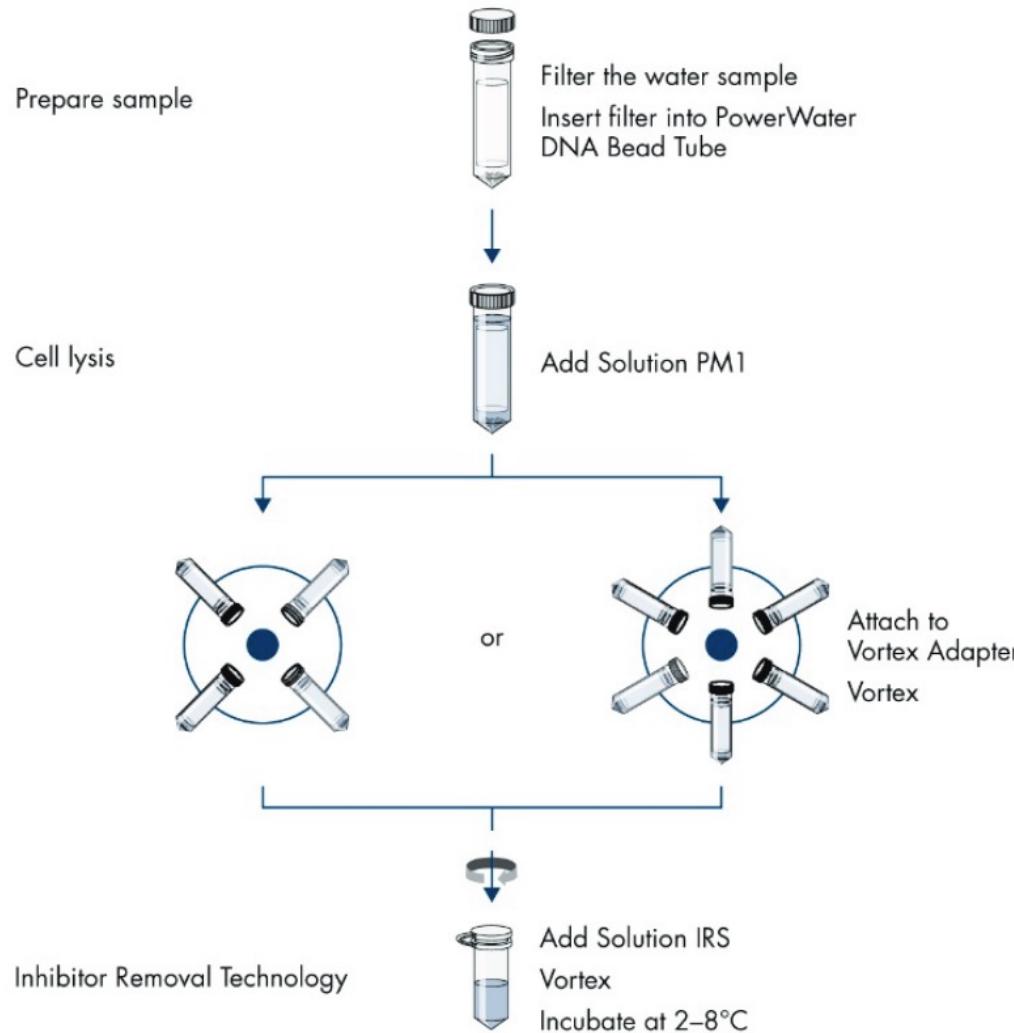
Collection

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



RNA Extraction

RNeasy PowerWater Kit Procedure



RNA Extraction

Bind DNA



Add Solution PM3 and PM4
Vortex
Load into MB RNA Spin Column



DNase I treatment



Wash with Solution PM5
Add DNase I and Solution PM6
Incubate at room temperature



Wash



Wash with Solution PM7
Wash with Solution PM5
Wash with Solution PM4



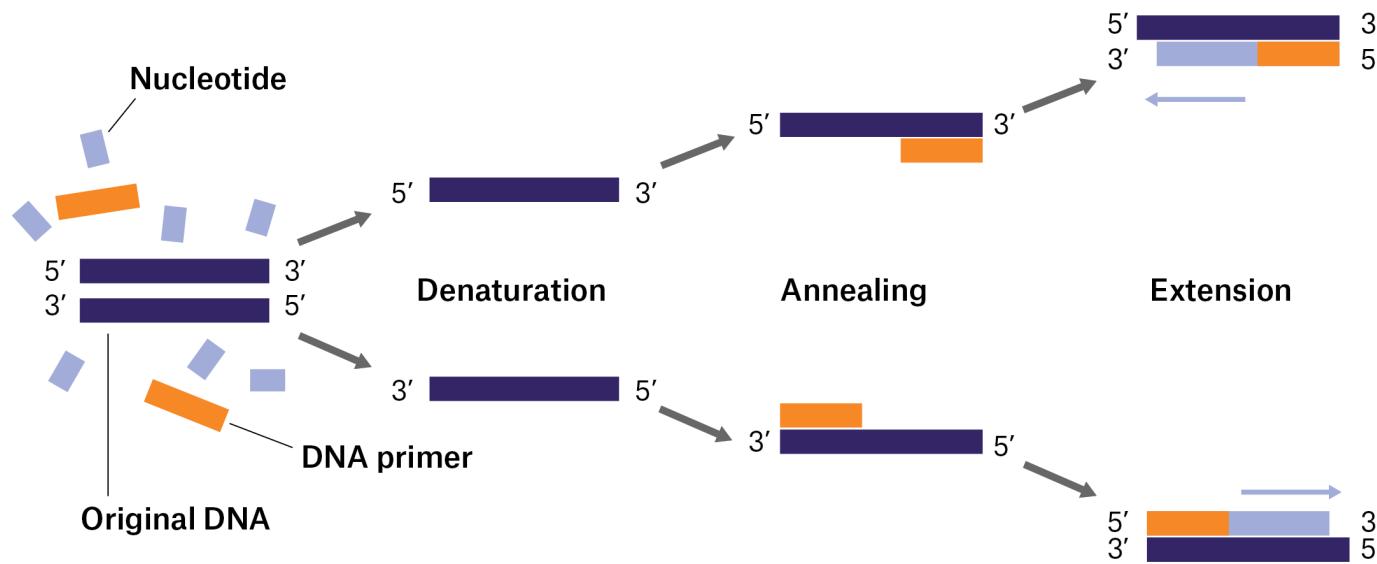
Elute



Elute with RNase-Free Water

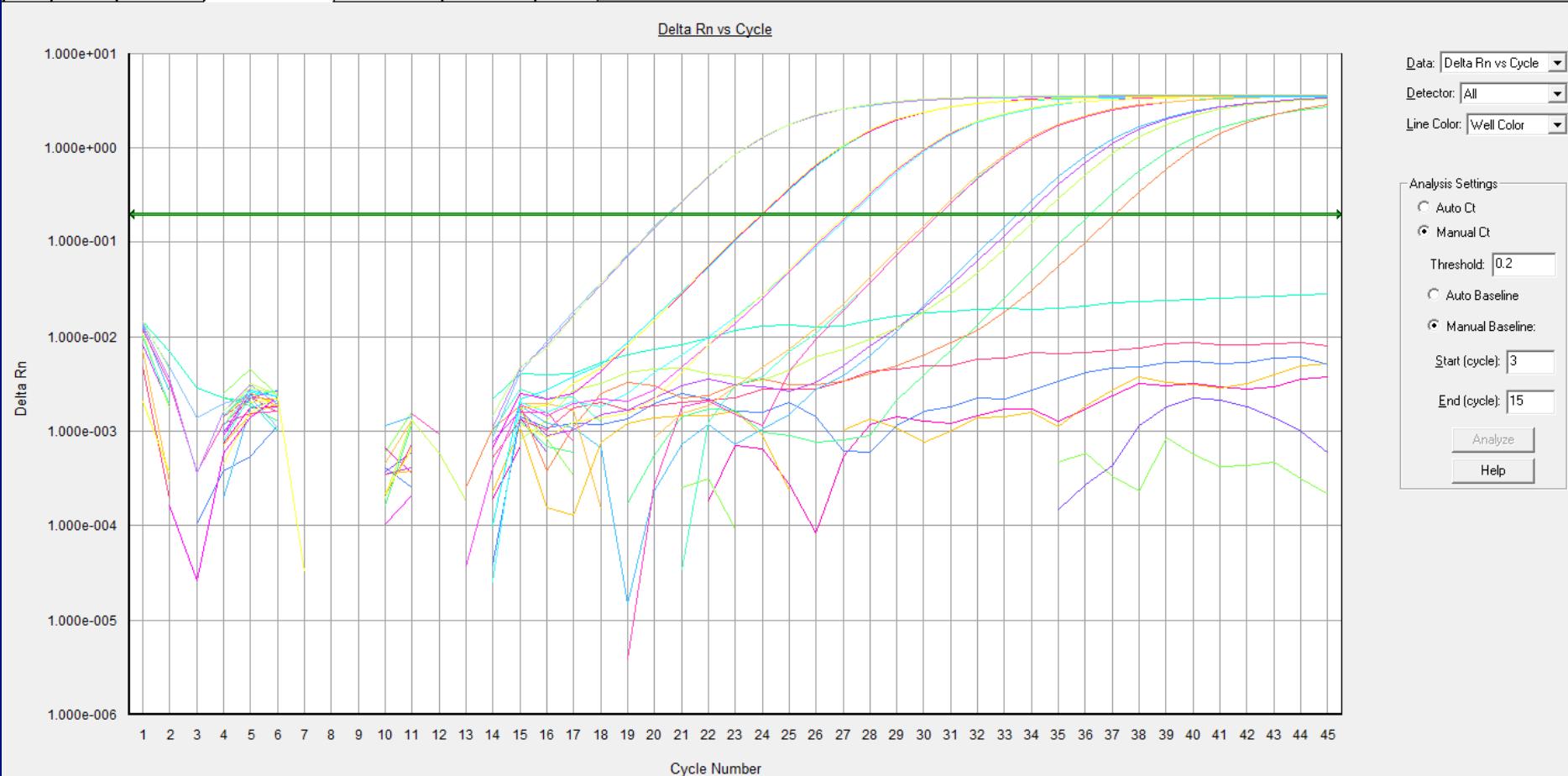
RT-qPCR

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

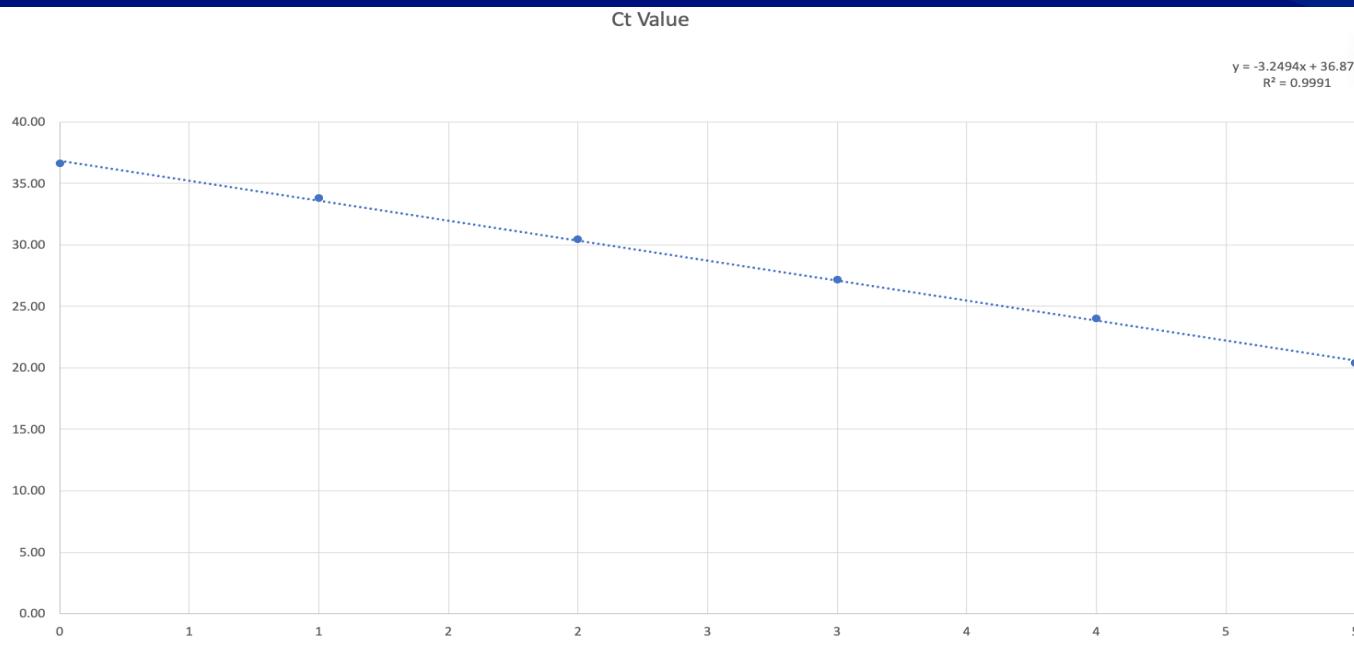


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Amplification Curve – June 9th



Standard Curve – June 9th

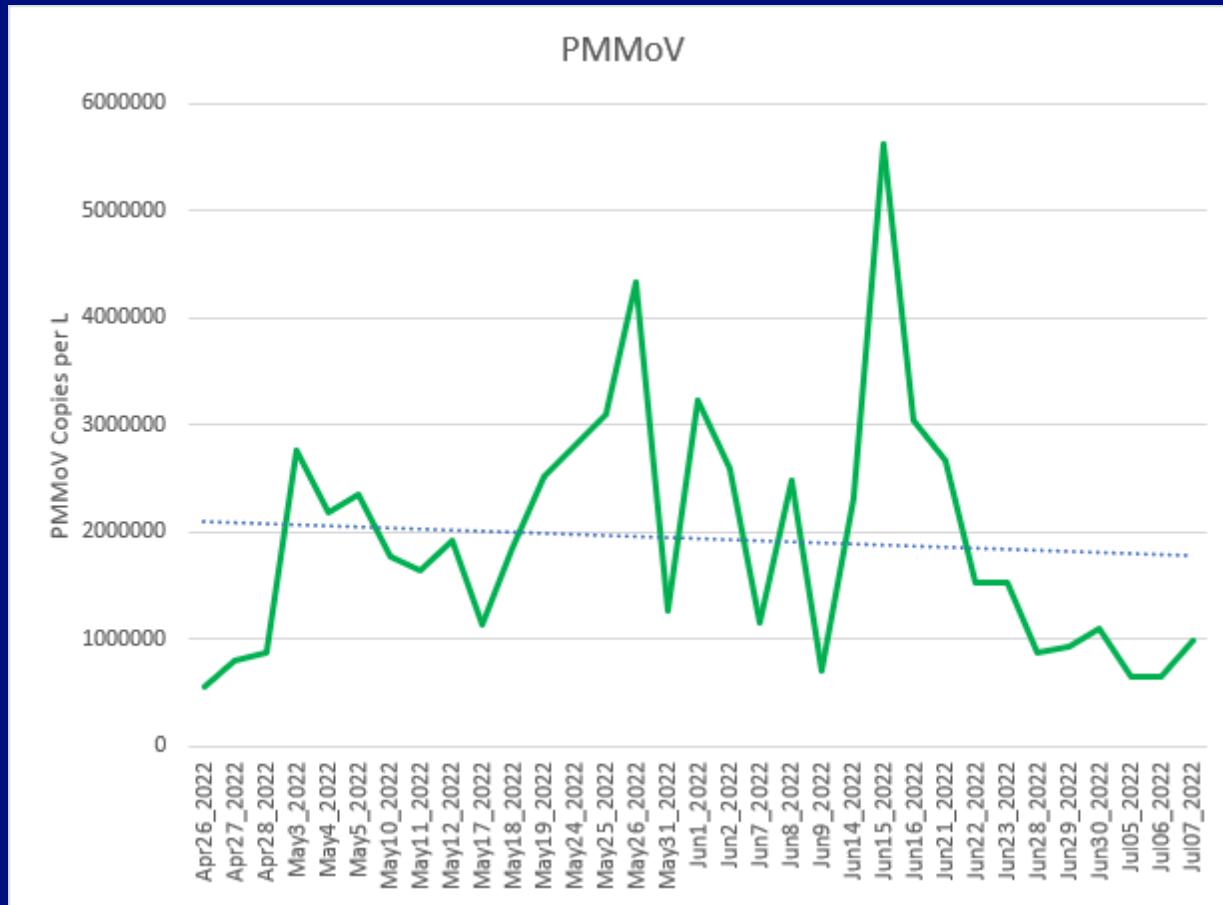


$$y = -3.2494x + 36.874$$
$$R^2 = 0.9991$$

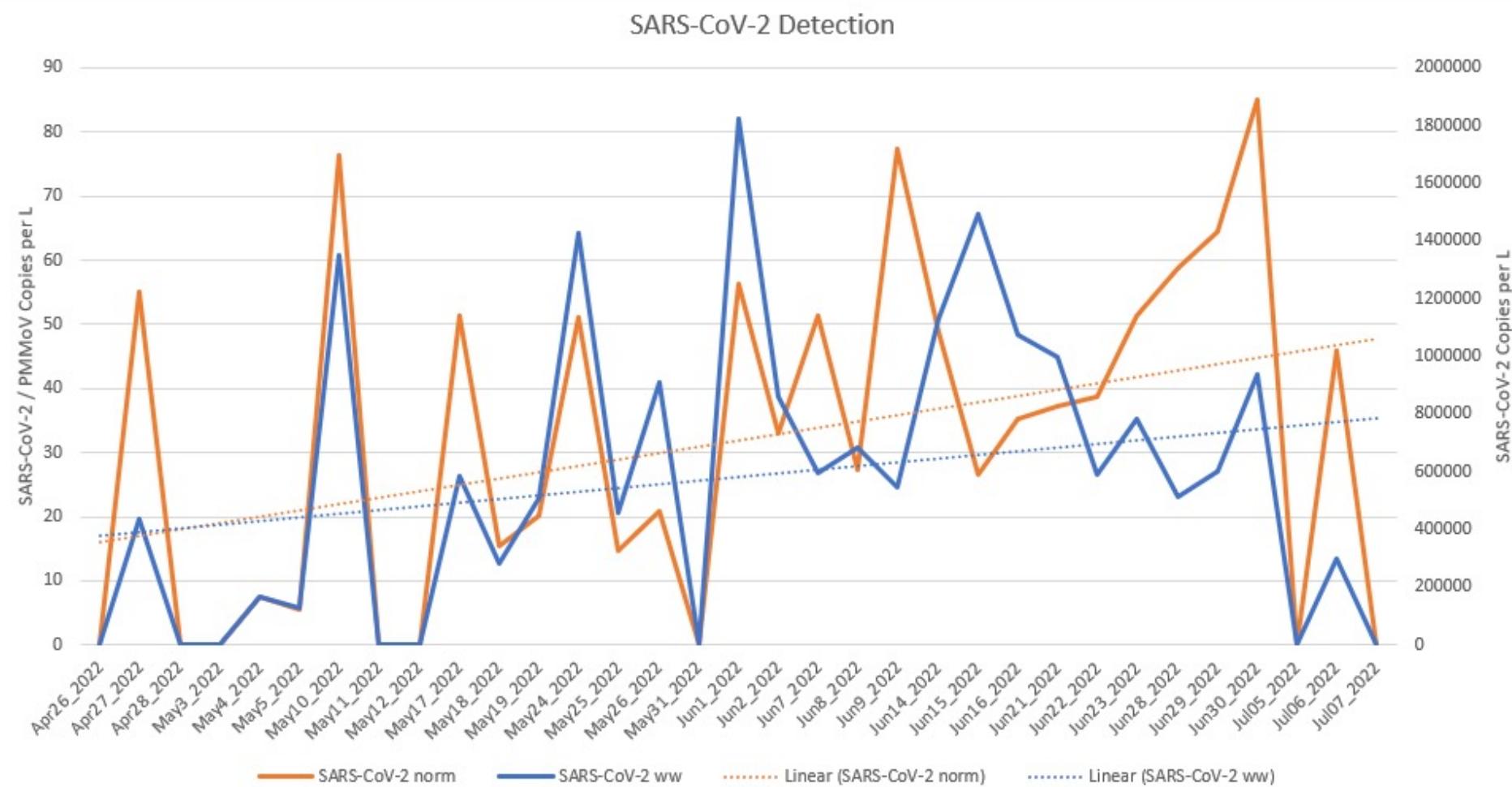
DNA Copies / uL	Log DNA Copies / uL	Ct Value
1	0	36.62
10	1	33.84
100	2	30.45
1000	3	27.17
10000	4	24.00
100000	5	20.43

$$32 = -3.2494x + 36.874$$
$$-4.874 = -3.249x$$
$$x = 10^{1.5}$$
$$x = 31.6 \text{ copies/uL}$$

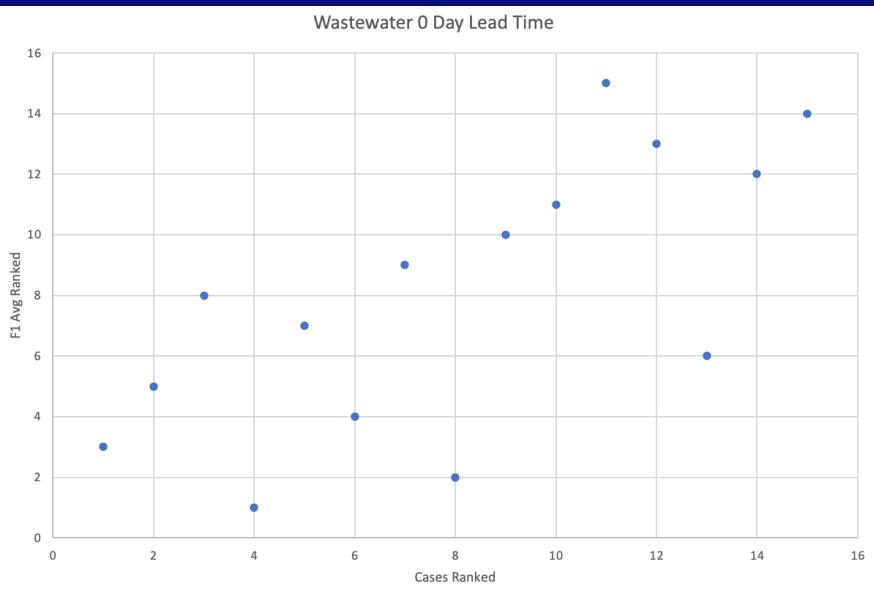
PMMoV Graph



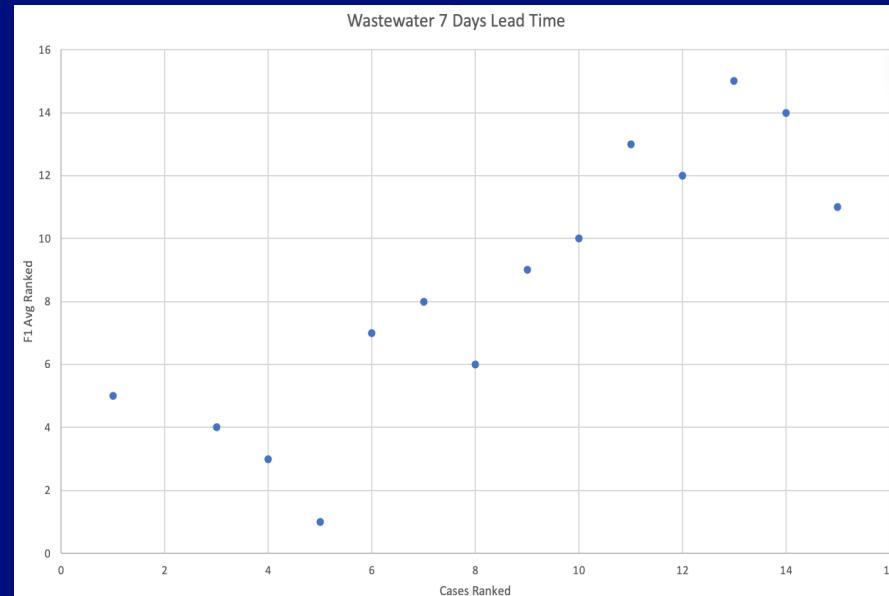
SARS-CoV-2 Detection Graph



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?