



# Virginia Wastewater Surveillance Program: Community of Practice Meeting

WWS Team

VDH | Office of Environmental Health Services

June 29, 2022



# Agenda

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- ❑ **Updates & Funding Opportunities**

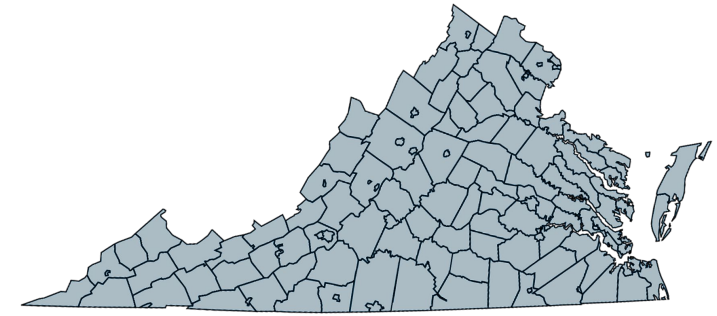
- ❑ **Topic(s) of Interest:**

- ❑ Tracking wastewater to understand infectious disease epidemiology

- ❑ **Open Discussion**

# Programmatic Updates

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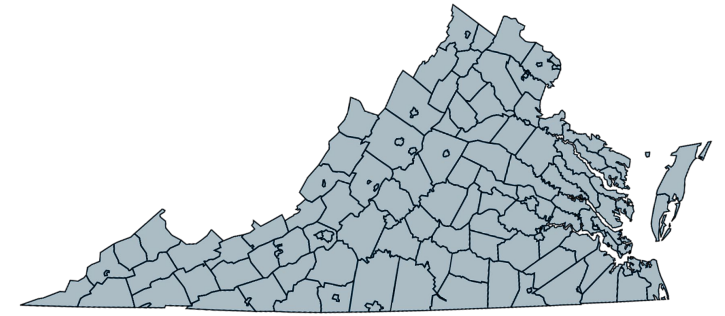


Weekly **SARS-CoV-2** monitoring at influent to **25 wastewater treatment plants** statewide started on **September 13: 41 weeks!**

- ❖ Weekly results sharing with:
  - Utilities
  - Health Department Partners
  - Environmental Health Managers
  - DCIPHER
- ❖ Updated Internal Working Dashboard Now!

# Funding Opportunities/ Updates

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## ❖ **BP4 Proposal to CDC**

- Next Funding Cycle-announcement around the corner

## ❖ **CDC-Biobot Commercial Sampling**

- Multiple sites enrolled from Virginia
- Must not be currently involved in CDC NWSS
- Sampling: twice/week

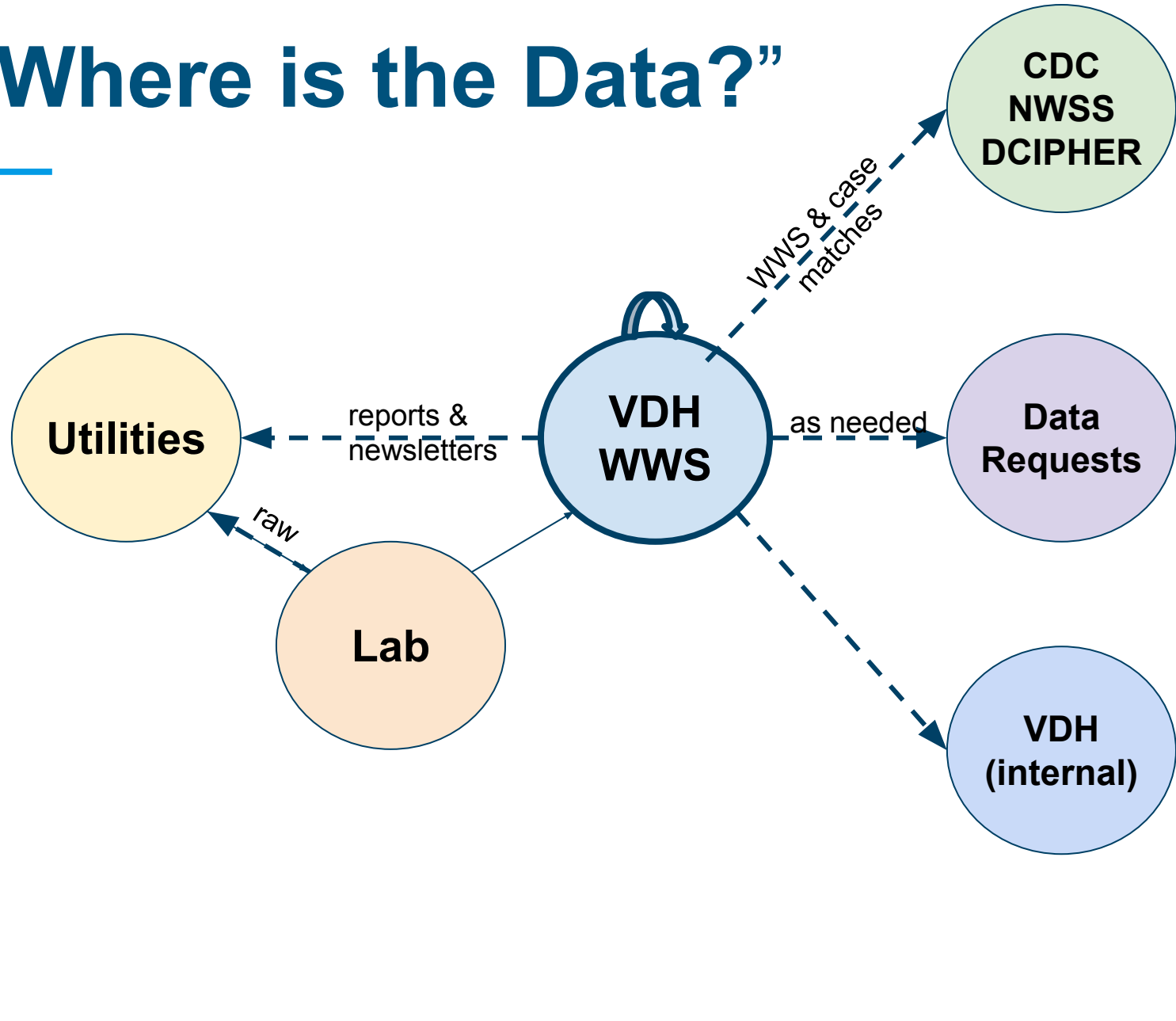
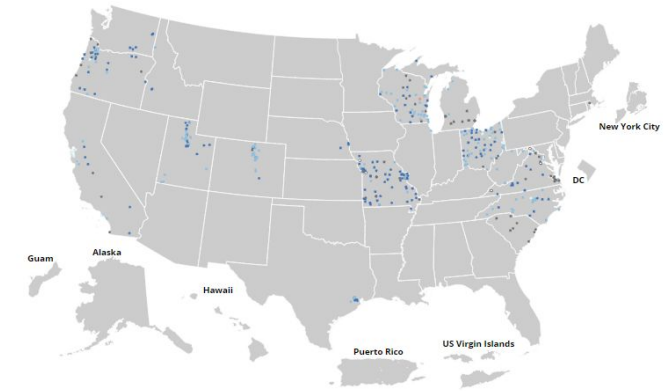
## ❖ **WEF Autosampler Program-Still accepting applications**

## ❖ **Localized Projects**

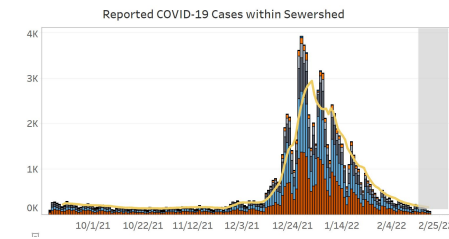
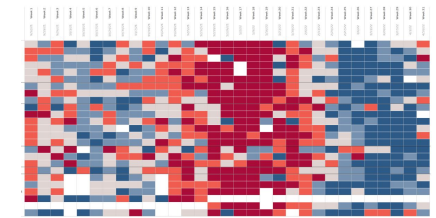
- Started 8 new sites in SW Virginia

# “Where is the Data?”

## CDC COVID-19 Data Tracker

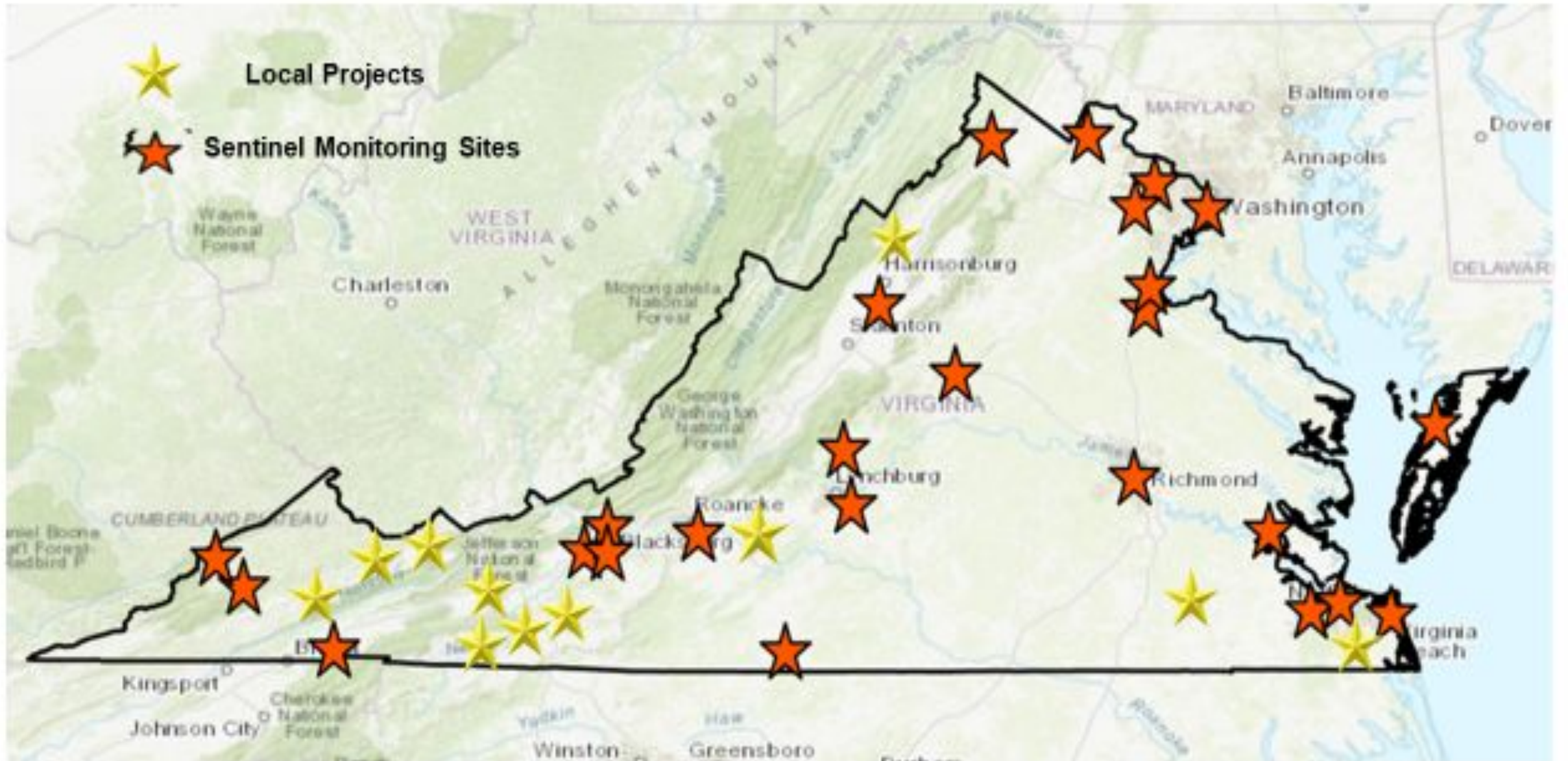


Ad hoc requests for:  
researchers, concerned citizens, etc.

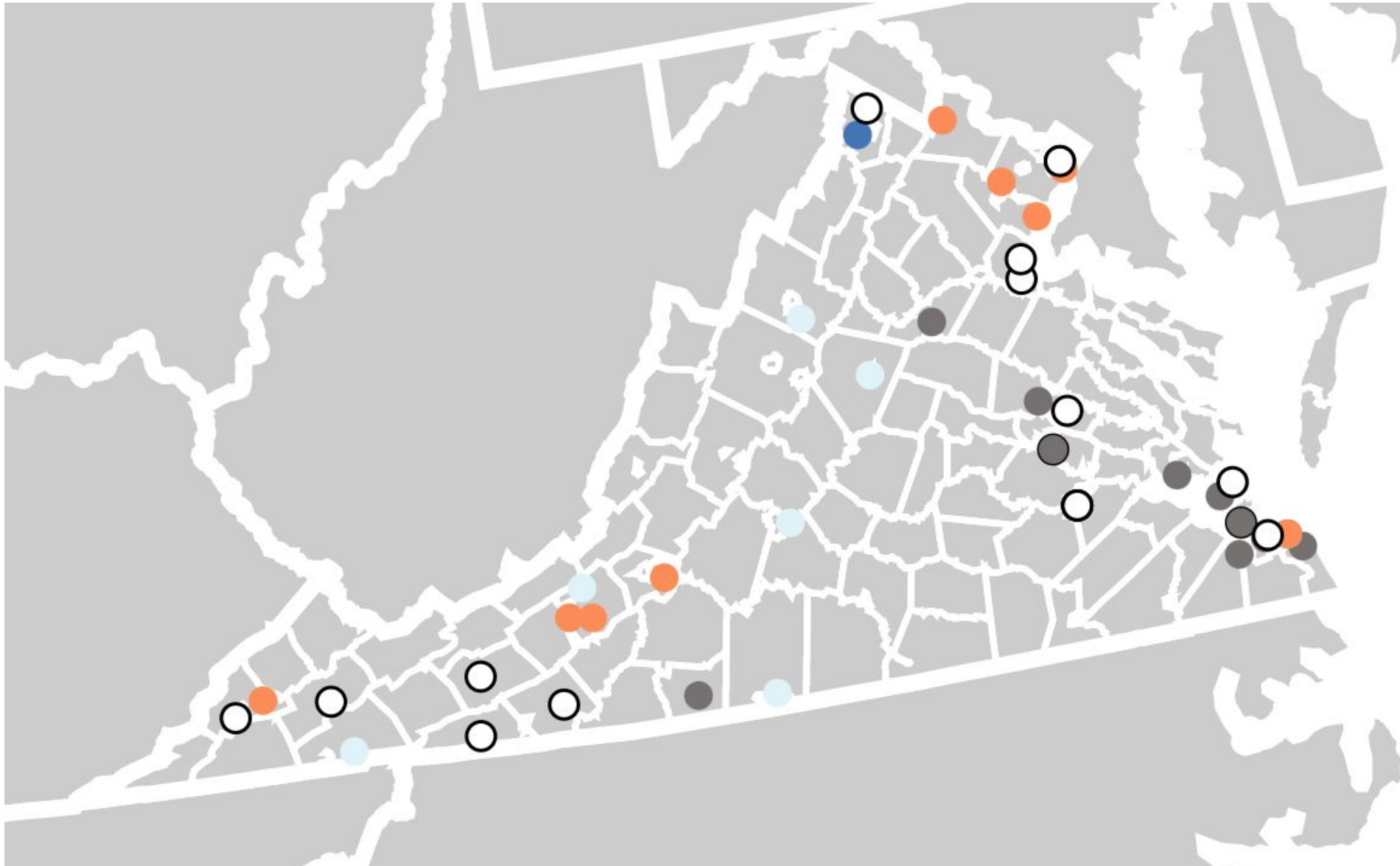


(In progress)

# Sentinel Monitoring Sites and Localized Projects



# COVID Data Tracker



○ CDC Commercial Contract Site

# Targeted Tracking Wastewater to Understand Infectious Disease Epidemiology

Dr. Amy J. Mathers MD, D(ABMM)

Associate Professor of Medicine and Pathology  
University of Virginia School of Medicine Division  
of Infectious Diseases & International Health  
Medical Director Antimicrobial Stewardship  
Associate Director of Clinical Microbiology





# Tracking wastewater to understand infectious disease epidemiology

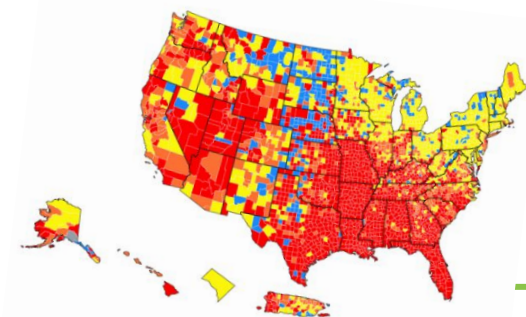
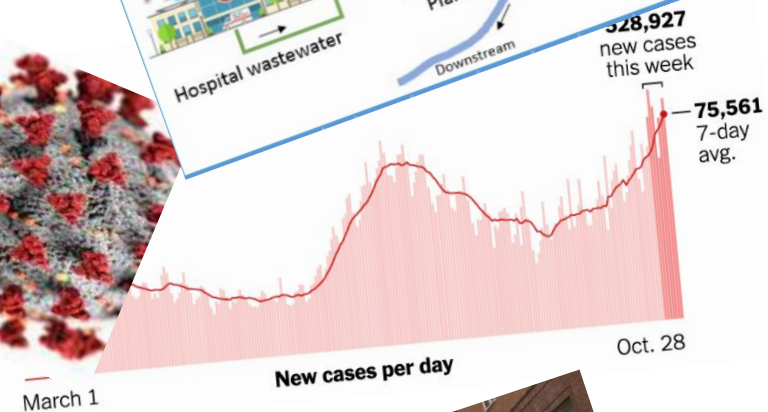
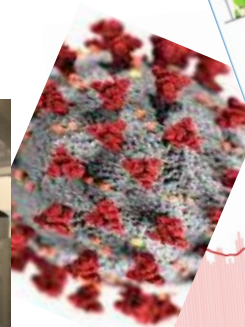
## Virginia WWS Community of Practice Discussion

Amy Mathers, MD, D(ABMM)

Associate Professor of Medicine and Pathology

University of Virginia

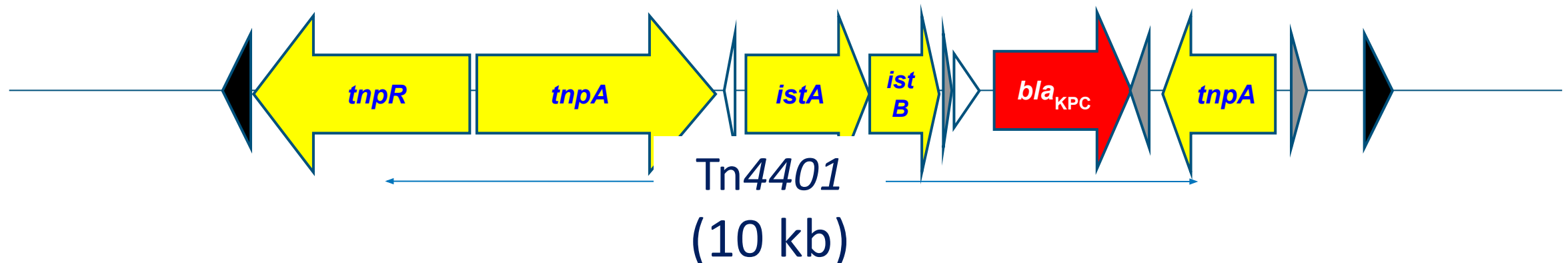
June 29<sup>th</sup>, 2022



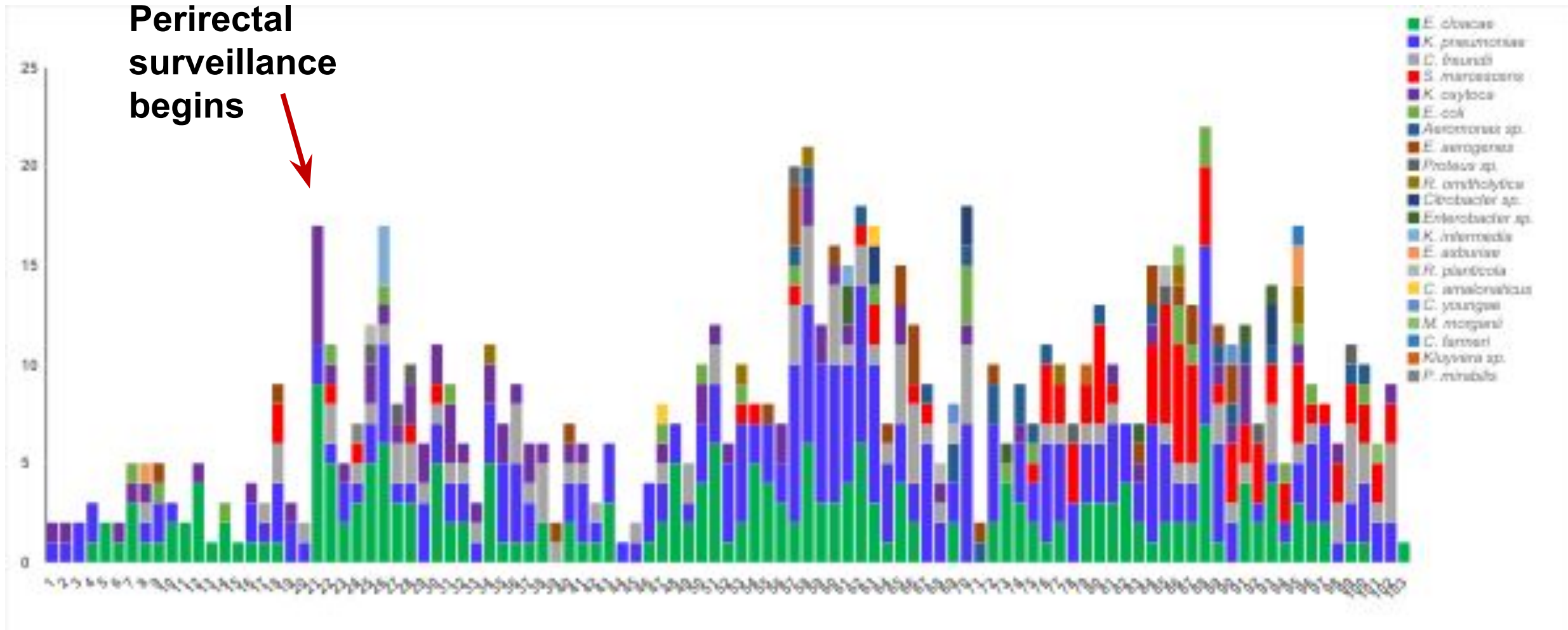
# *Klebsiella pneumoniae* carbapenemase

## (KPC)

- Found primarily in *Enterobacterales* (majority of initial reports seen in *K. pneumoniae*)
- Ambler Class A serine  $\beta$ -lactamase
- Hydrolyzes all  $\beta$ -lactams; penicillins, extended spectrum cephalosporins, aztreonam and carbapenems
- Is inhibited by avibactam, vaborbactam, relabactam and other novel  $\beta$ -lactamase inhibitors
- Gene is contained on a mobile piece of DNA which bacteria share

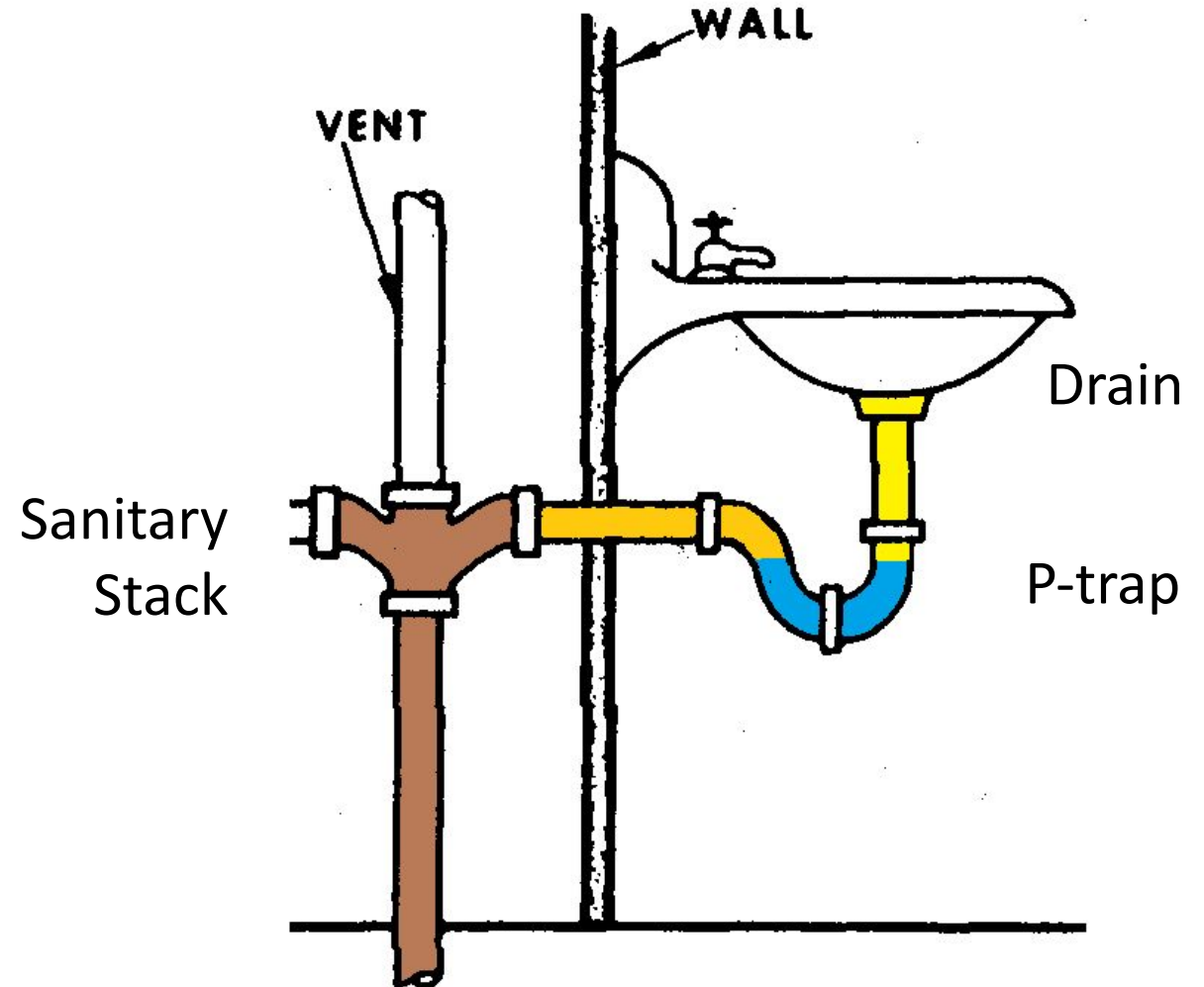


# Incidence of KPC-producing Bacteria by Species in our institution

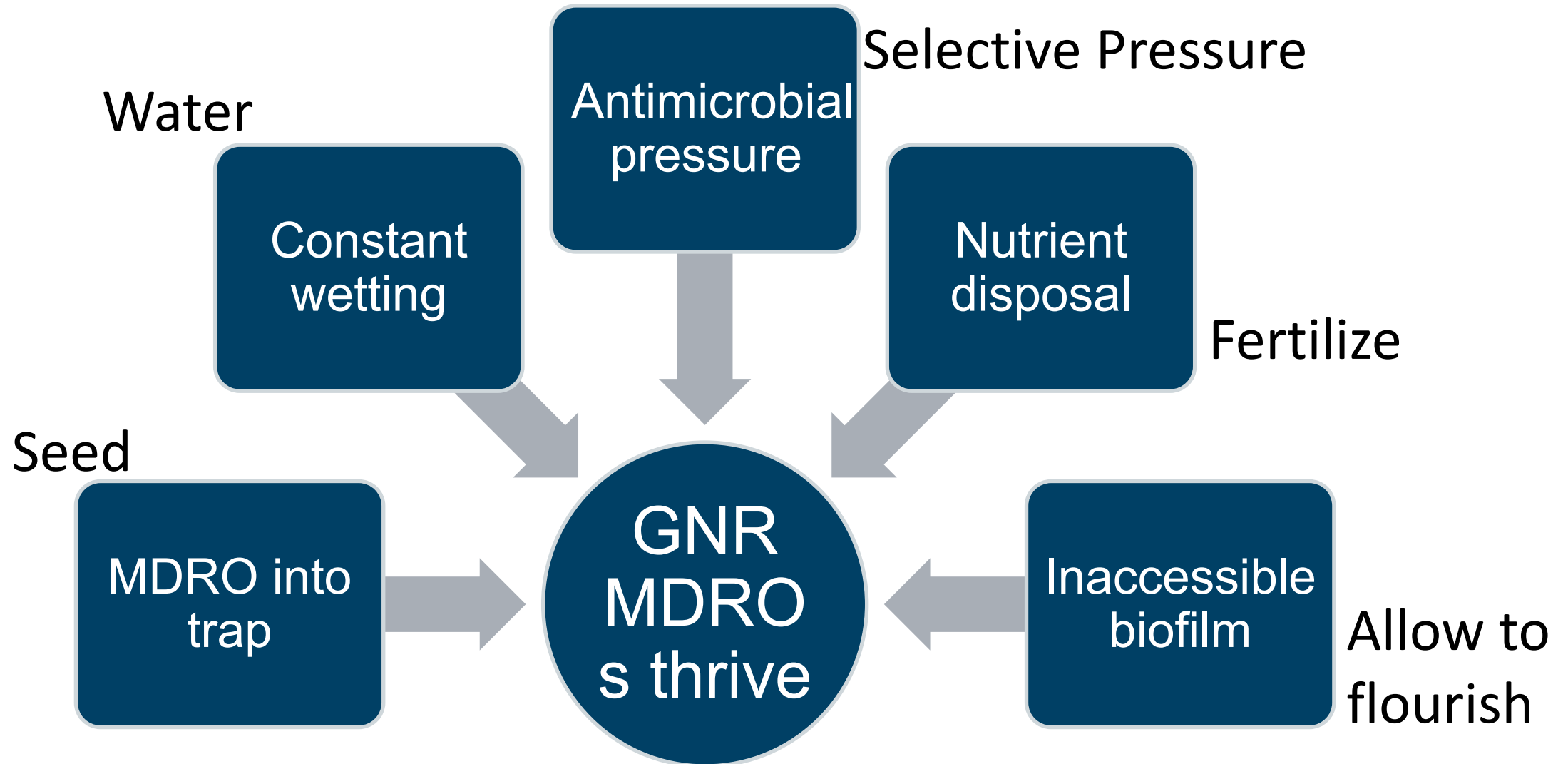


# Consequential genes of drug resistance may make this issue much more notable and high risk

- Gaps remain in our knowledge of nosocomial transmission of ESBL/carbapenemase producing Enterobacteriaceae
- Increasing evidence that some patient acquisition may occur from colonized sink drains
- Sorting out transmission chains is not easy even with a clonal outbreak but especially across species with mobile genetic elements involved



# Ideal niche for antibiotic resistant bacteria to evolve and flourish

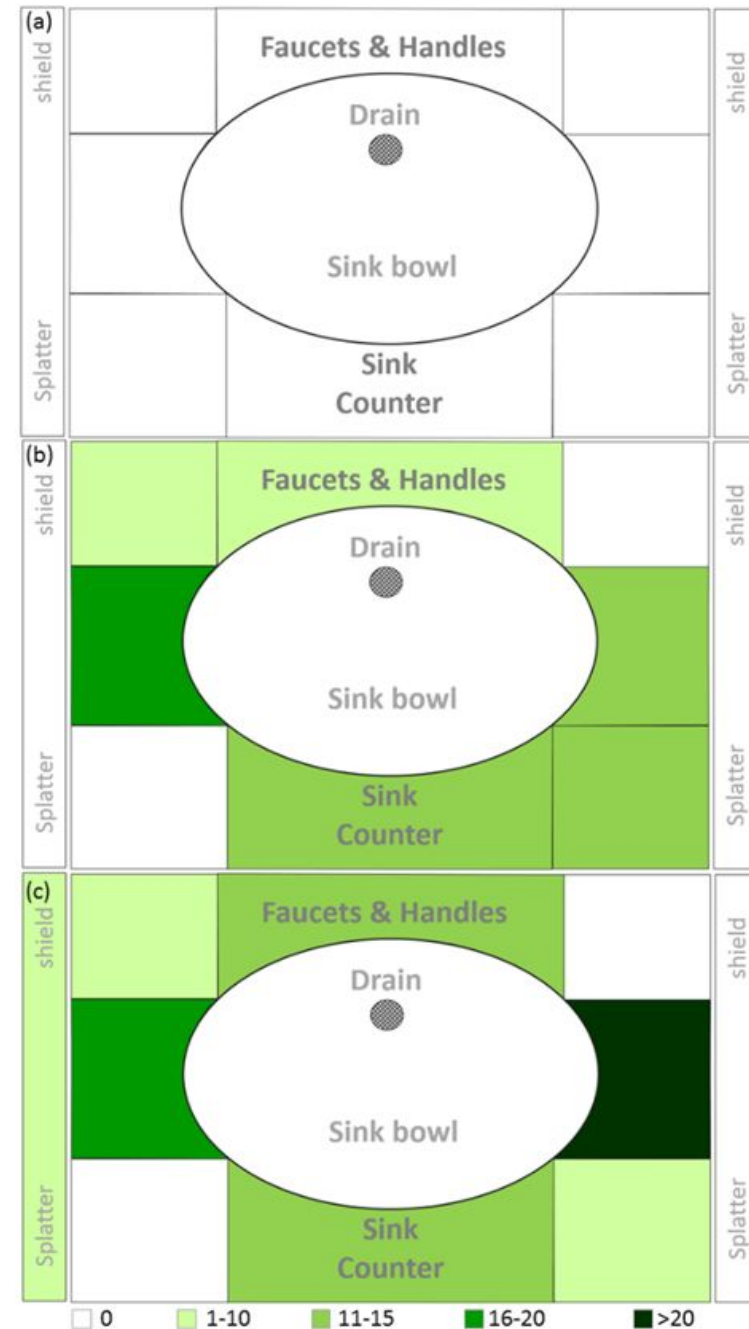
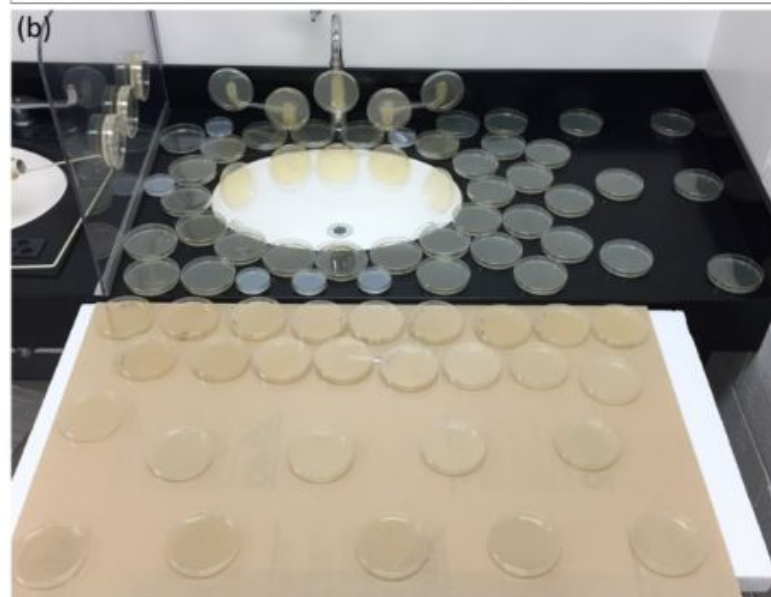
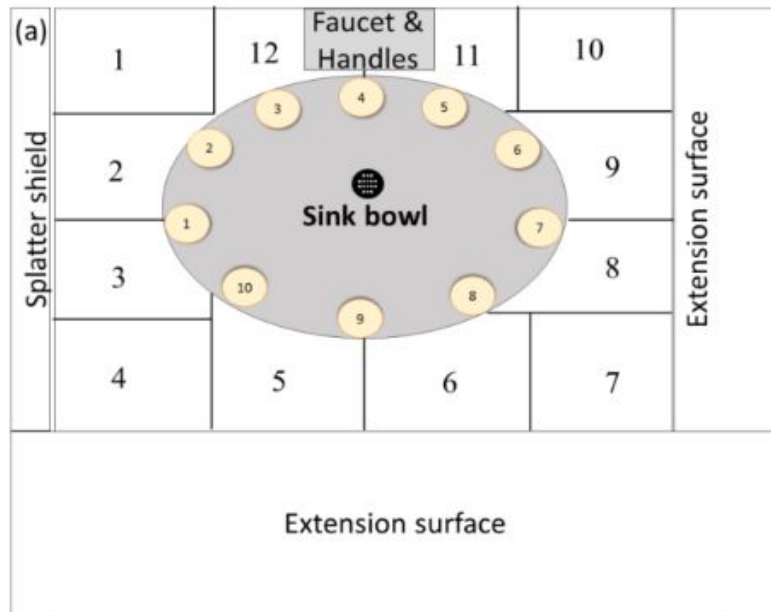


# Sink Lab(s)



- Working with CDC
- Aim to understand microbial dynamics in controlled setting
- Looks to develop interventions
- Requests from standards, industry and other hospitals

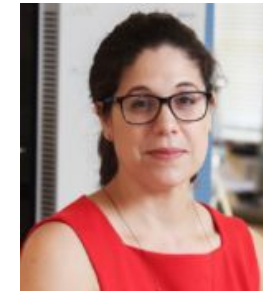
When colonizing the drain or sink bowl *GFP-E.coli* dispersed onto the sink and surrounding counter top when hit with water



# What happens to the KPCO when they leave the hospital?



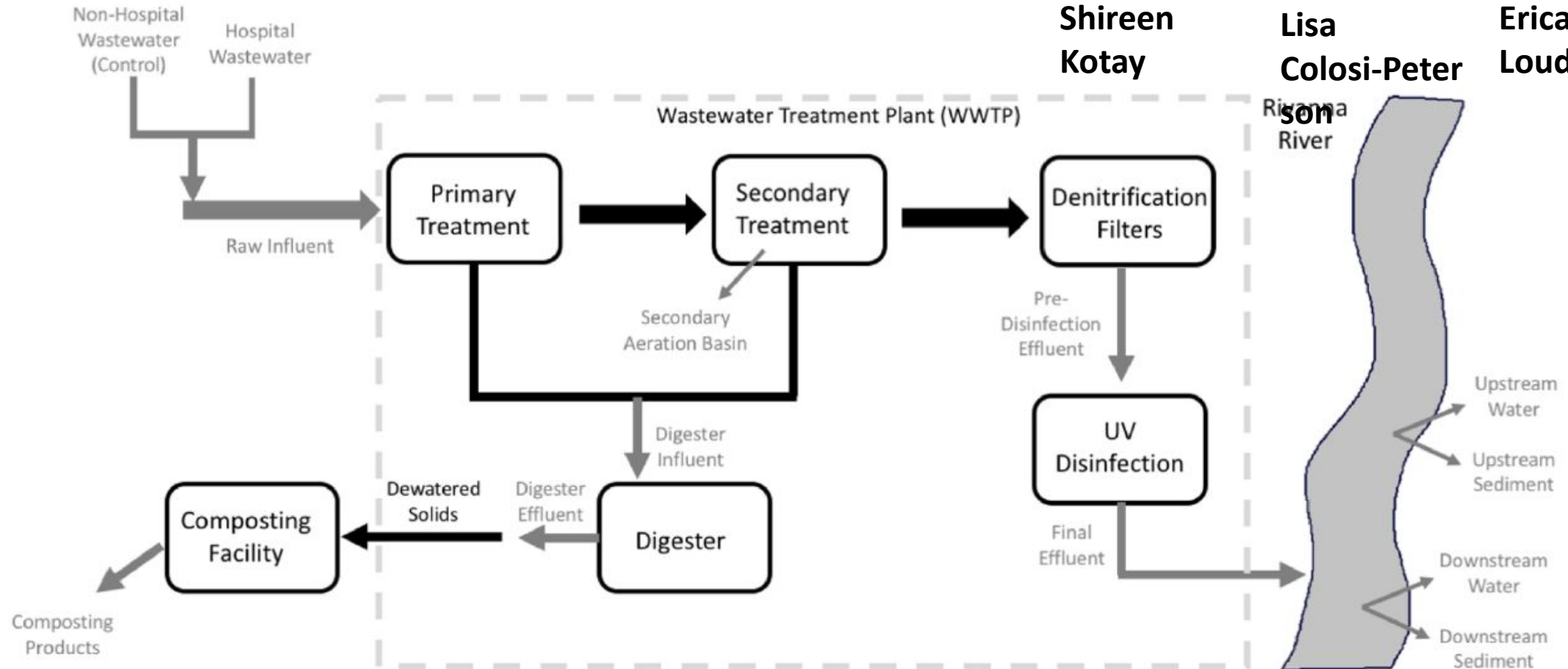
**Shireen  
Kotay**



**Lisa  
Colosi-Peter**



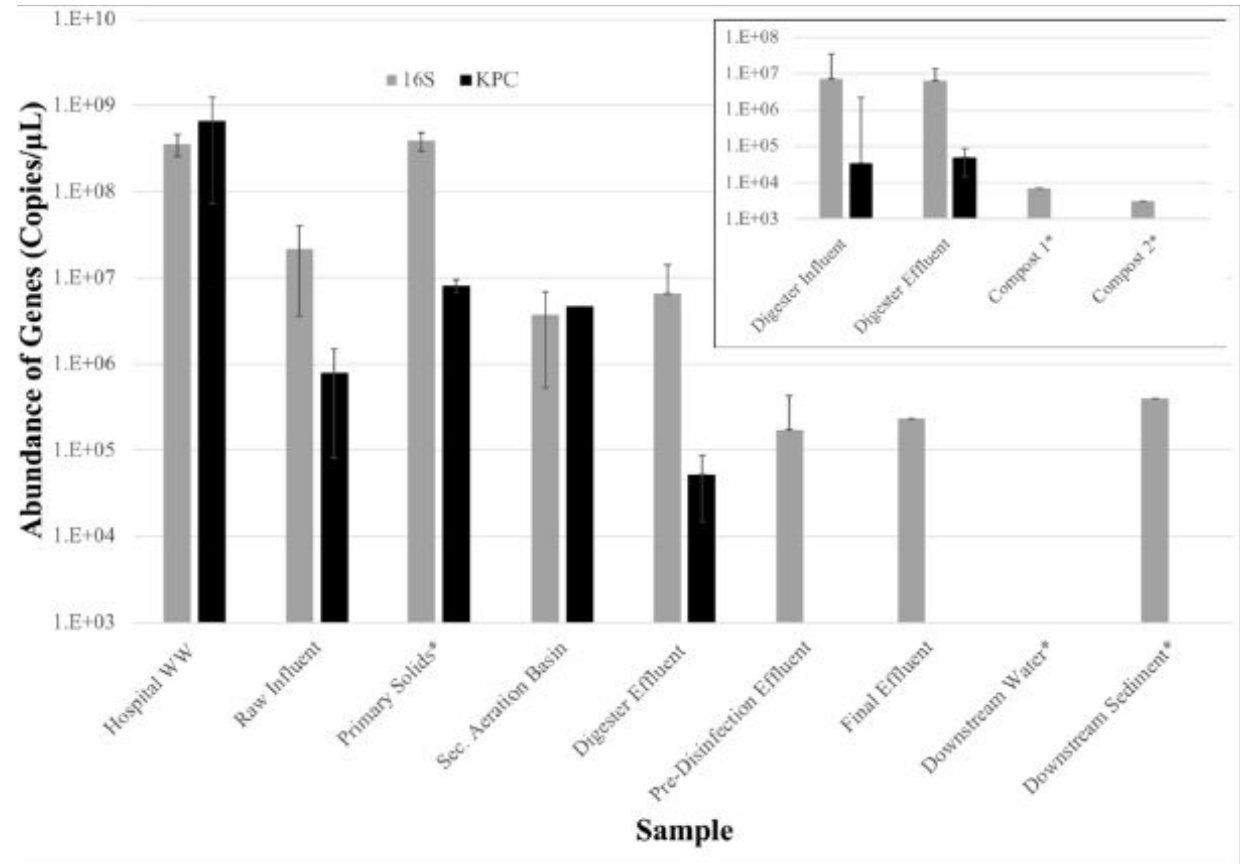
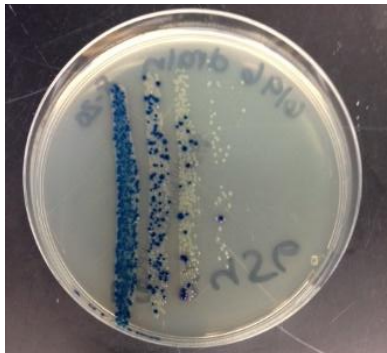
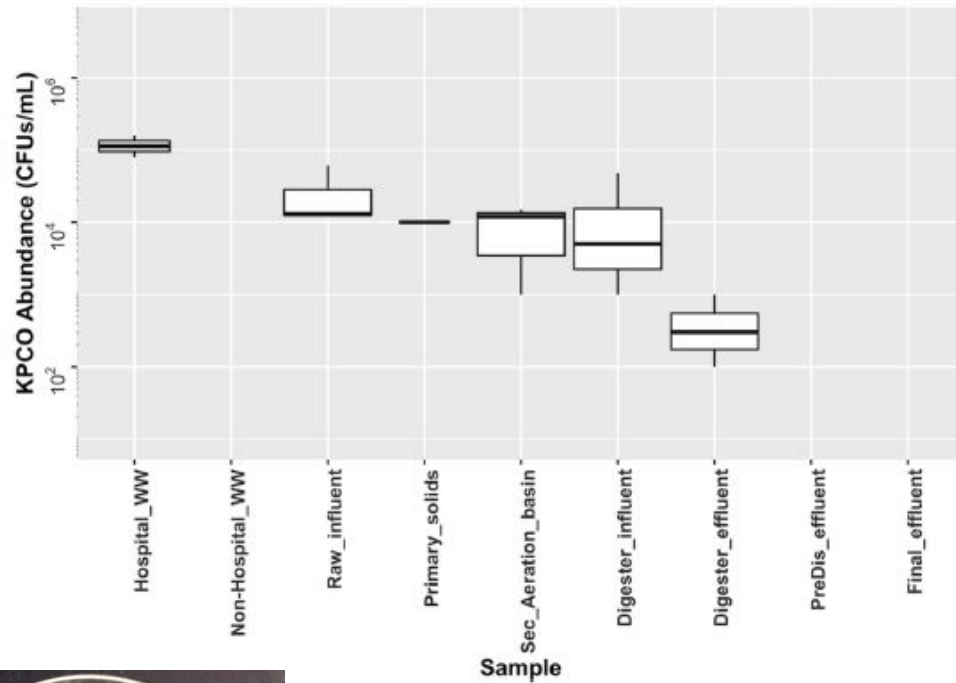
**Erica  
Loudermilk**





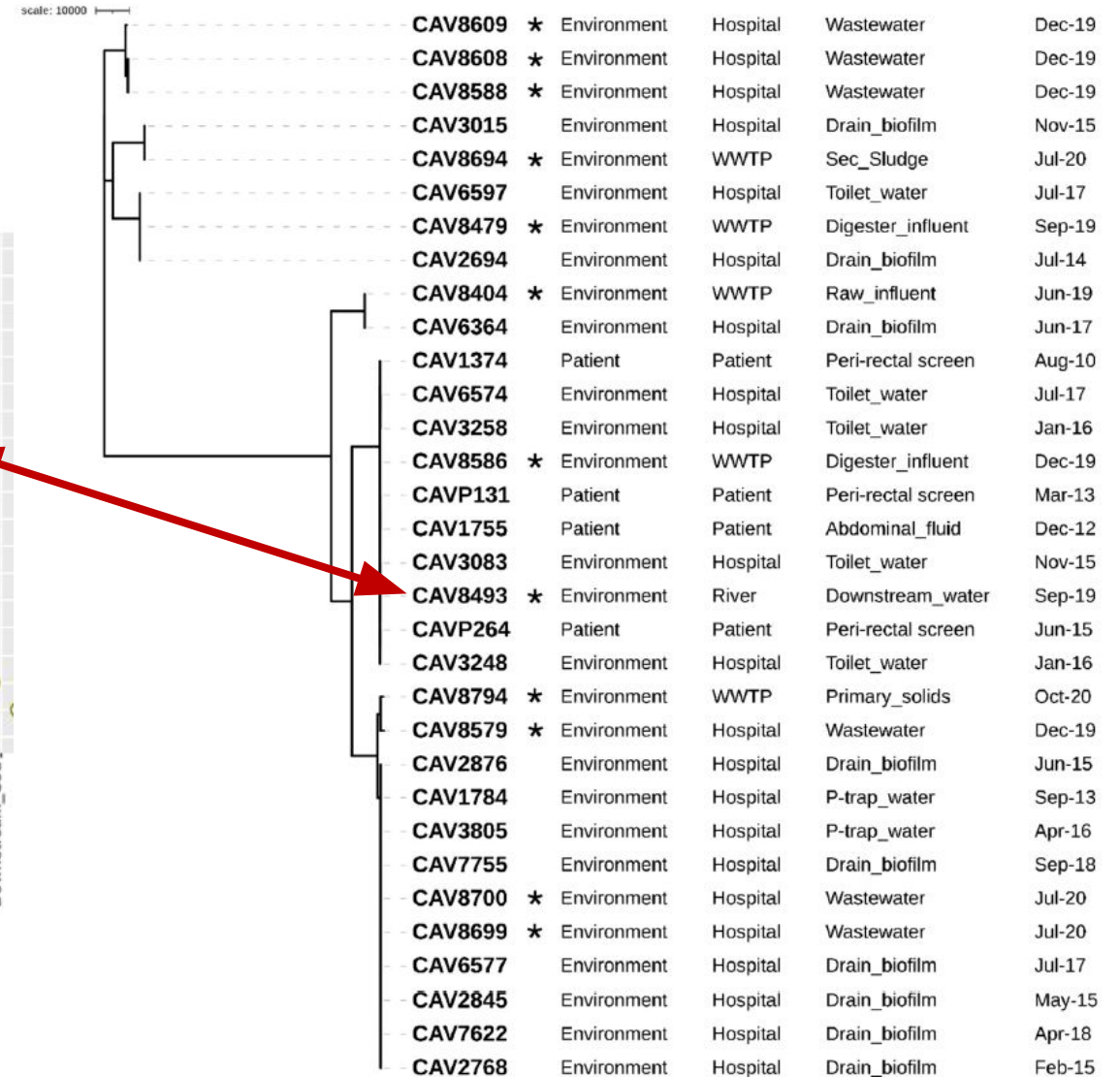
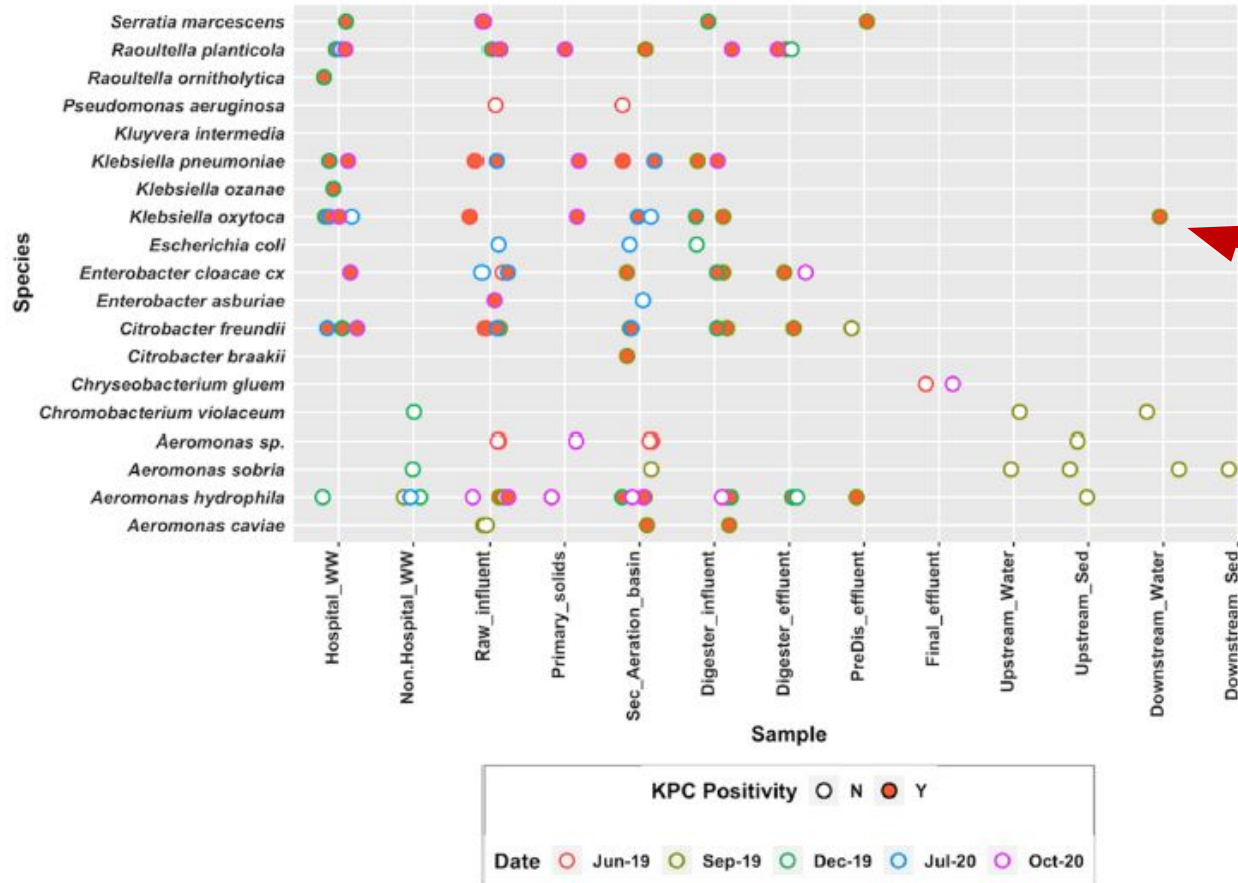
# Culture and qPCR align for KPCO

## WW works well to eliminate $bla_{KPC}$ and KPCO



# Culture data from across the continuum diverse KPCO

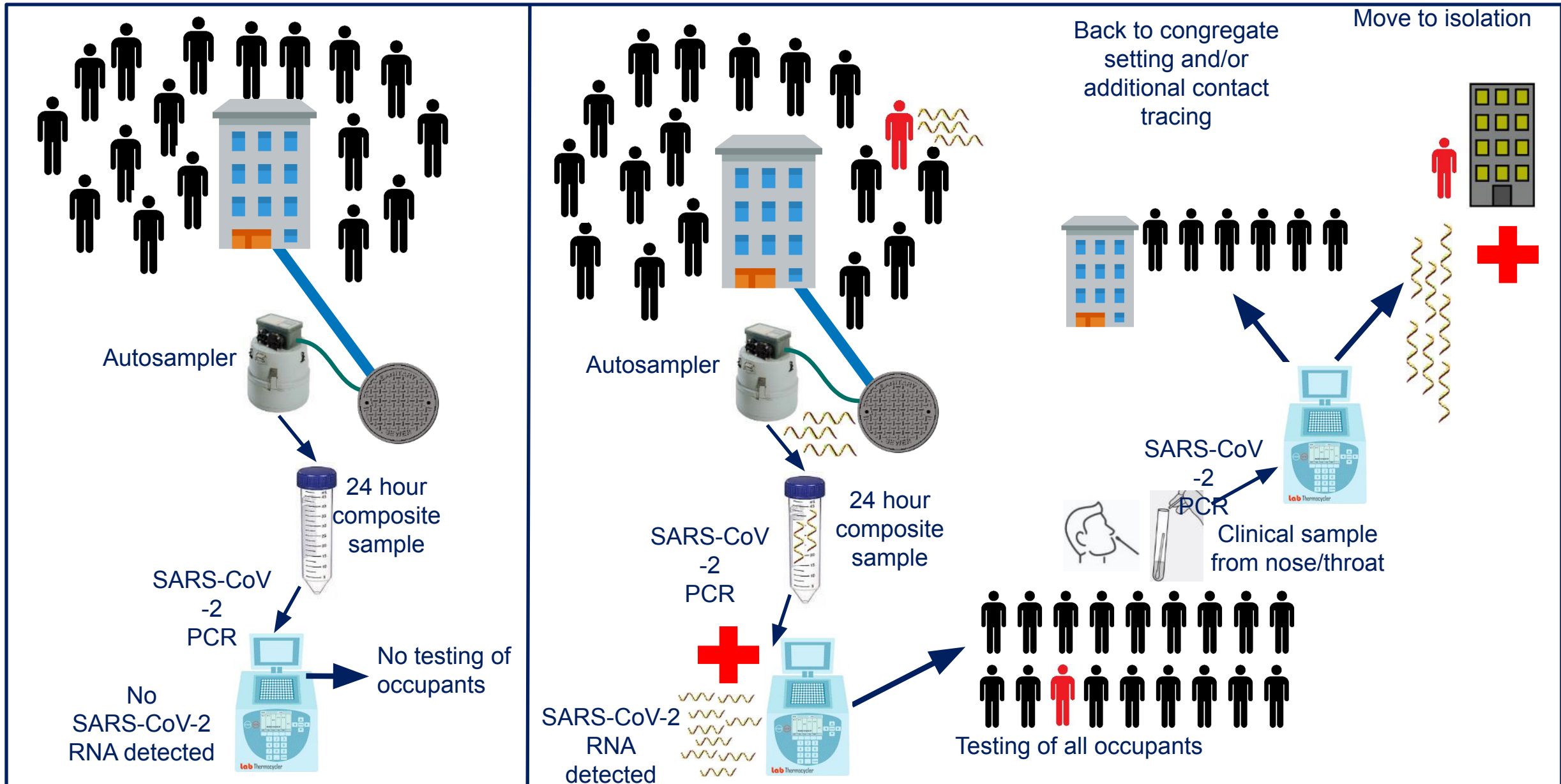
One KPC-*K. oxytoca* in downstream water identical to prior patient/hospital isolates



## Need for massive surveillance on congregate living but not enough resources available to test everyone

- Early data was emerging that SARS CoV-2 RNA was detectable in stool and municipalities were beginning to monitor community level via waste water treatment plants
- Could this technology be used to detect new positives in a building without the challenges around specimen collection
- Different group of people doing the work and could be done passively
- May require different resources and personnel who are less taxed by the pandemic

# Concept behind pooled wastewater testing

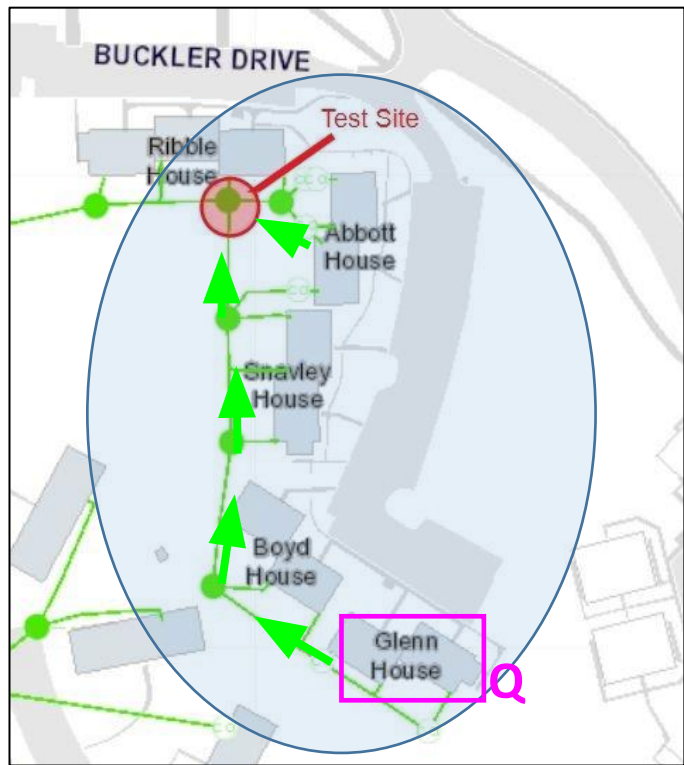


# We did not know if it would work or how it should work (and/or if we could make it work)

- Was working on a collaboration with Lisa Colosi-Peterson for monitoring antimicrobial resistance in WW effluent
- Had established relationships with facilities within the health system and across campus
- Began discussions with a few groups had sprung up across the US who were looking to do a similar project (e.g. Syracuse U.)
- There were other groups on campus with interest in WW testing for predicting trends at a larger level
- Administration was very supportive of the effort



We know we can detect very few positives from a large group



○ Buildings flowing into collected sample.

□ Q Quarantine dorm

● Sampled manhole

➔ Wastewater flow

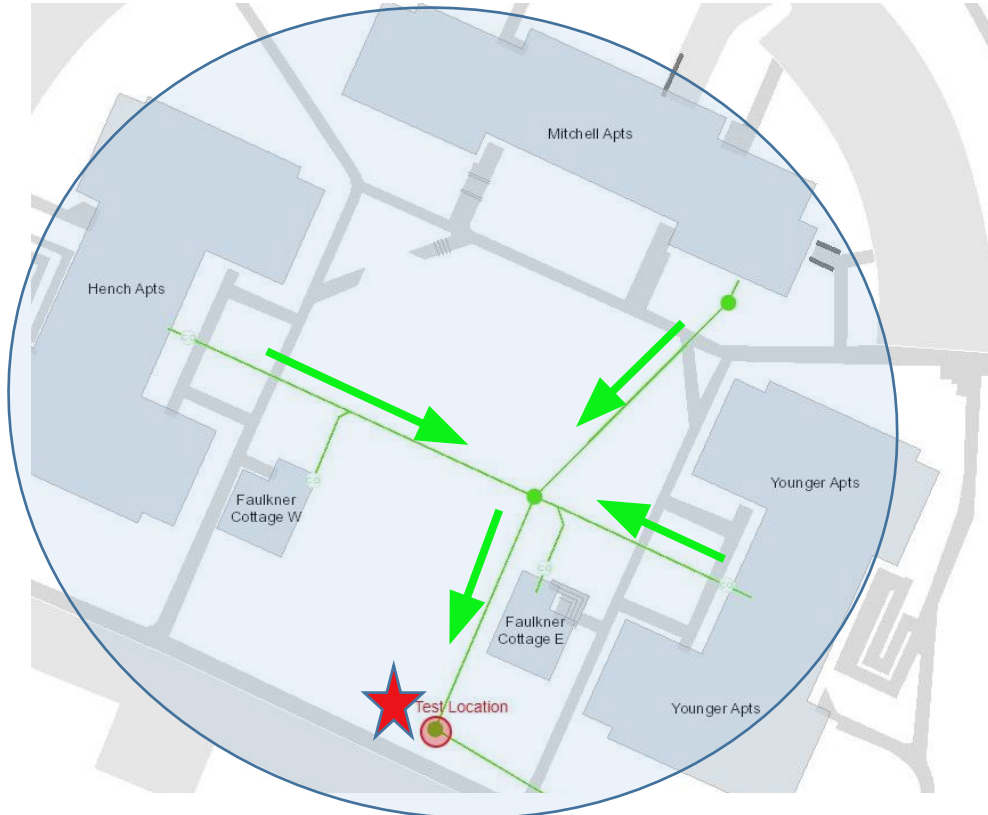
	Week								
	1	2	3	4	5	6	7	8	9**
<b>Occupants tested positive*</b>	2	1	0	0	0	0	0	0	0
<b>Occupants testing negative</b>	103	102	102	102	102	102	102	102	<102
<b>Occupants in onsite quarantine</b>	2	3	1	0	0	0	0	0	0
<b>WBT result at</b>	N/A	+	+	+	+	+	+	+	-

Key: WBT yields positive result (+) even though no contagious occupants were in residence.

\*positive occupants never retested



\*\*occupants began moving out

# When all occupants are testing negative we get a negative result initially



 Buildings flowing into collected sample.

 Sampled manhole

	Week		
	1	2	3
<b>Total occupants</b>	<b>66</b>	<b>66</b>	<b>66</b>
<b>Occupants testing negative</b>	<b>0</b>	<b>0</b>	<b>0</b>
<b>WBT result at site (Pos/Neg)</b>	<b>NA</b>		

Sensitivity of 96.2% and a specificity of 100%.

However, the method could not distinguish new infectious cases from persistent convalescent shedding of SARS-CoV-2 RNA.

If the detection of convalescent shedding is considered a false positive, then the sensitivity is 100% and specificity drops to 45%.

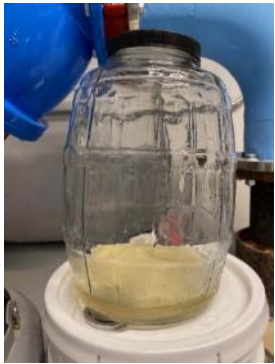
# Settled/tested workflow



24 hour collection



2 hour ultracentrifuge



Extraction and amplification 6 hours  
Using CDC EUA on the Applied Biosystems 7500



	Rp	N1	N2
Dorm A	29	27	31
Dorm B	26	32	31
Dorm C	ND	ND	30



By the time we decided to do it so had everyone else

HACH AS950



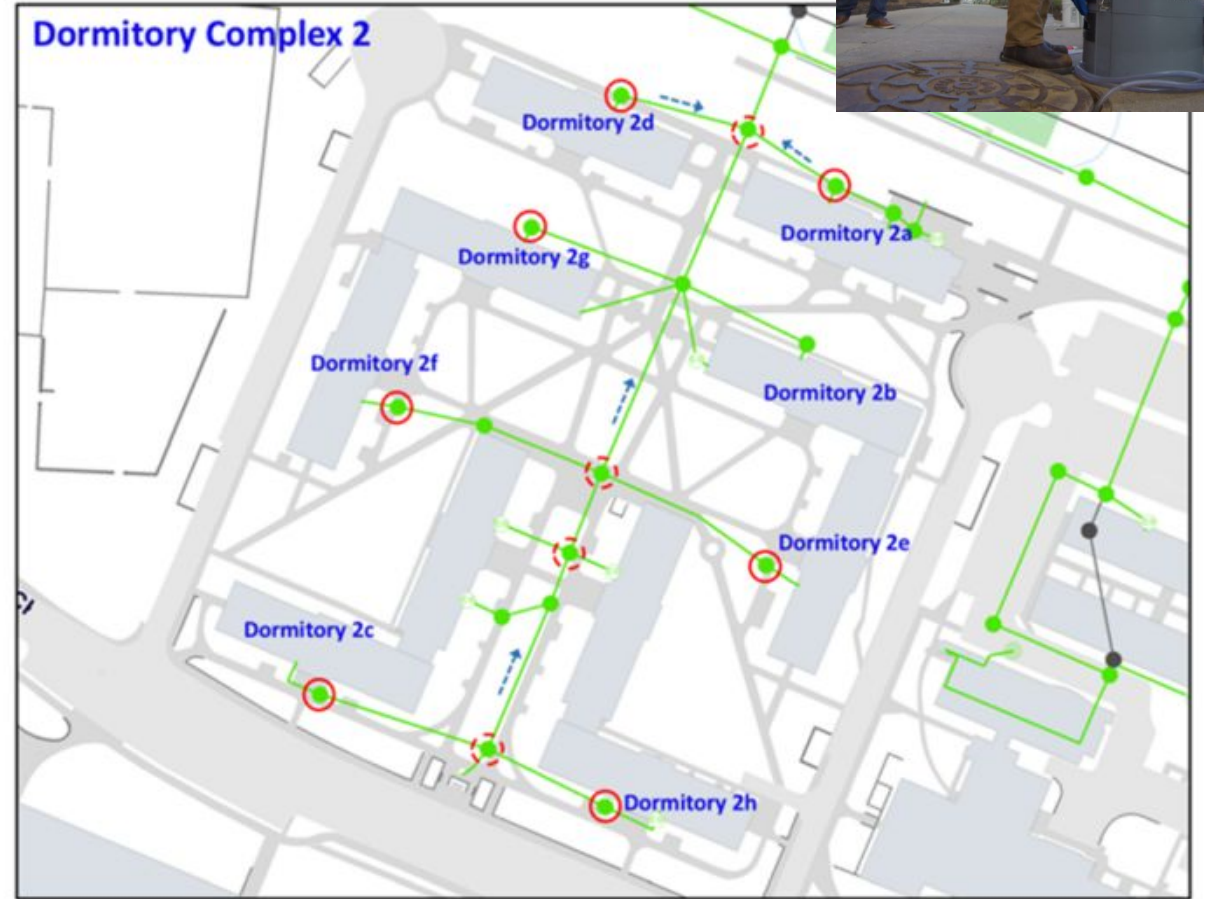
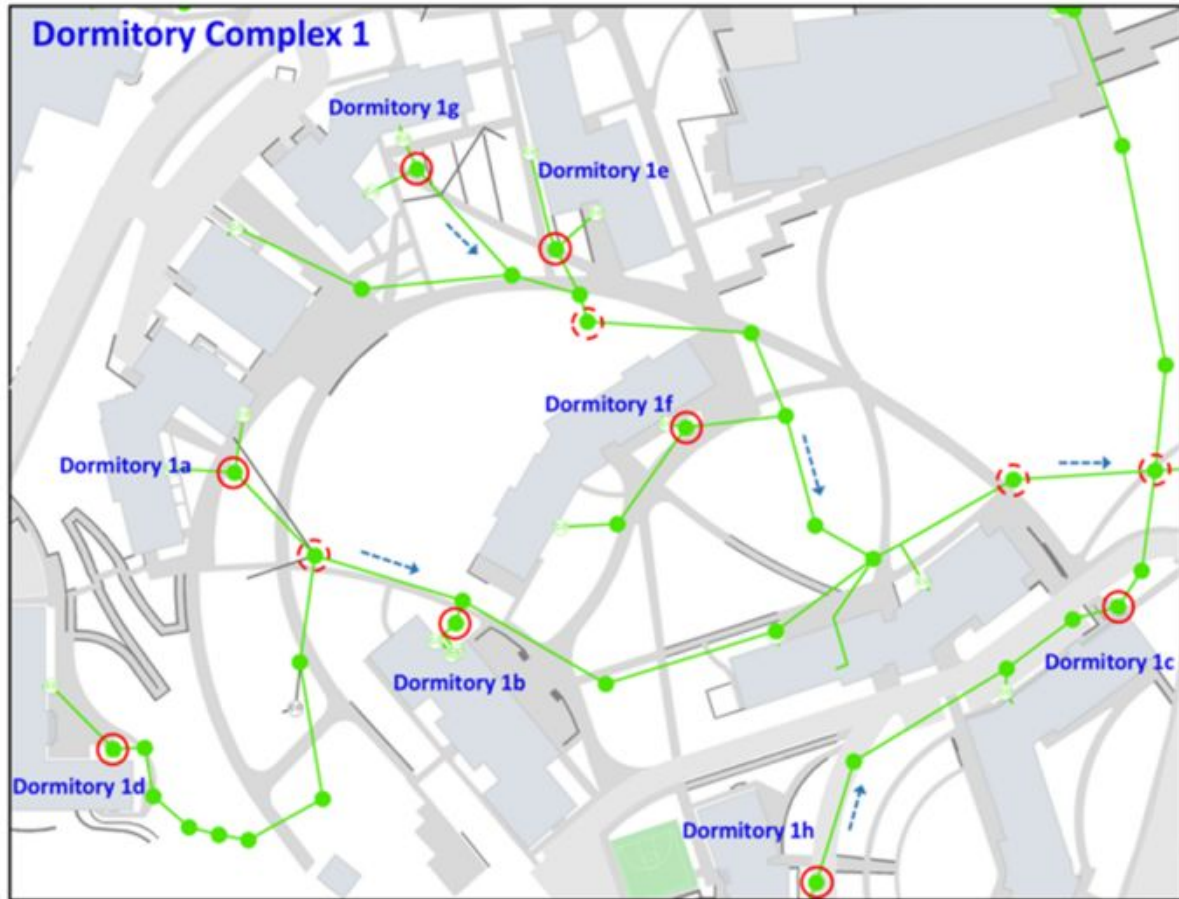
ISCO GLS Compact



UVA-AS V1



# We deployed 18 autosamplers daily to the dorms



First two weeks were very stressful but the WW testing helped limit outbreaks

## Mid-September

- PP based on WW positivity @ Balz-Dobie with 10 cases detected
- Next day PP based on WW positivity @ Lefevre 3 cases detected.
- Next Day PP @Echols- 4 positive, @Kellogg – 7 positive
- Next week PP @Hancock – 9 positive
- Next Day PP @Page – 8 positive



# Attempting to assess what the data means

Very difficult to interpret the data because of the convalescent shedding

Made an initial attempt at quantitation with the current method but difficult to normalize with molecular methods

It does seem that the data can only be interpreted in a series

So many factors the data did not always aligning with the results from the dorm occupant testing

Initial attempt at trying to understand signal strength and data interpretation

KEY:		
Negative	CT > 40	
Indeterminate		
+	CT 35-40	
++	CT 30-35	
+++	CT < 30	
Not sampled		
Failed	Did not pass QC	

# McCormick results summary example date

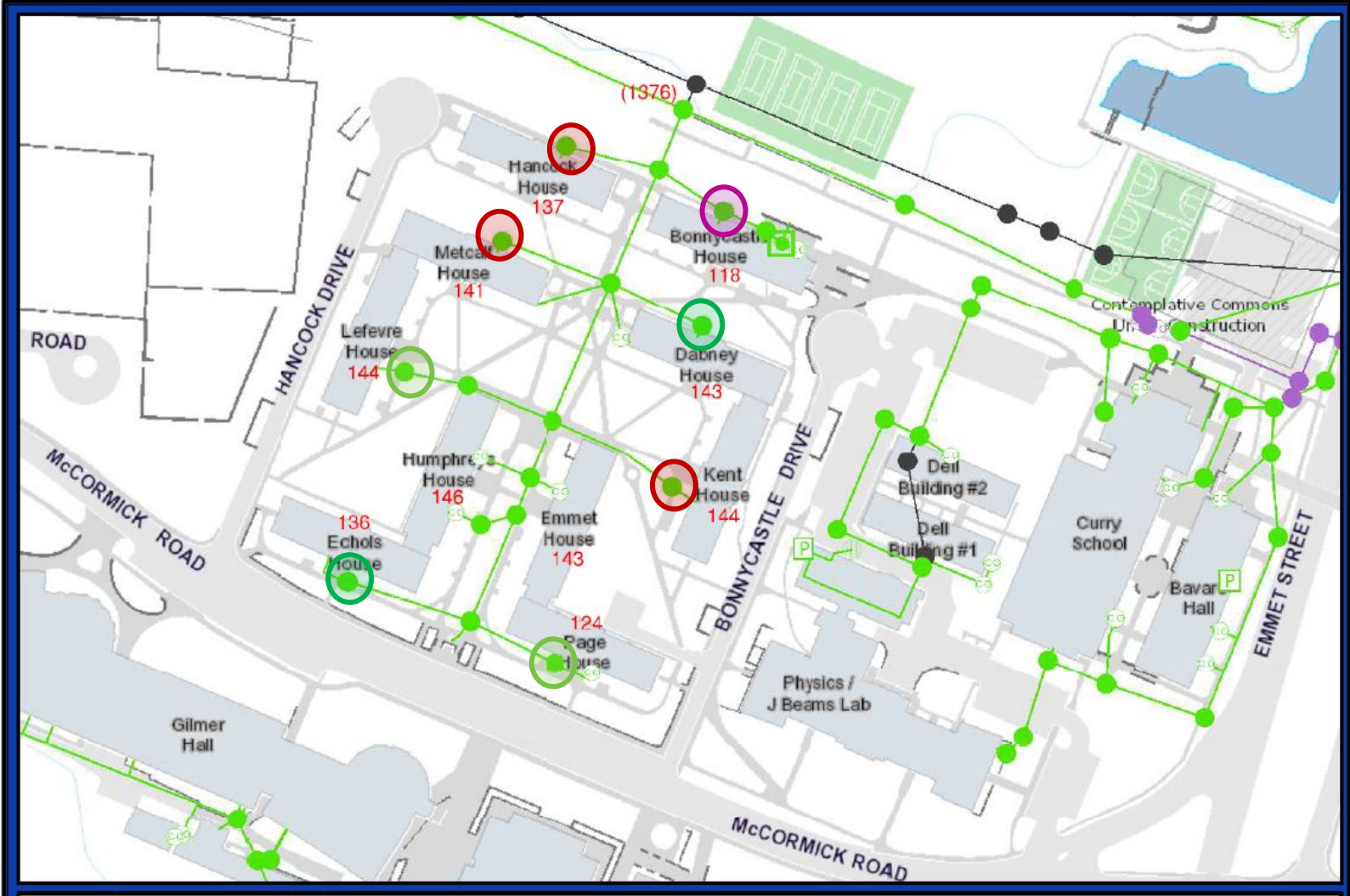
Strong positive signal X 2 days  
+Bonnycastle

Positive signal monitoring\*  
+Hancock  
+Metcalf  
+Kent

Negative  
+Echols  
+Dabney

Not able to interpret results  
+Page  
+Lefevre

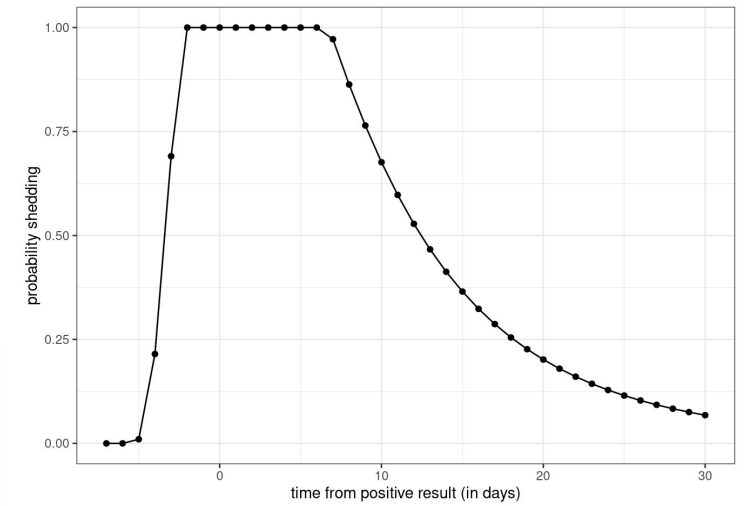
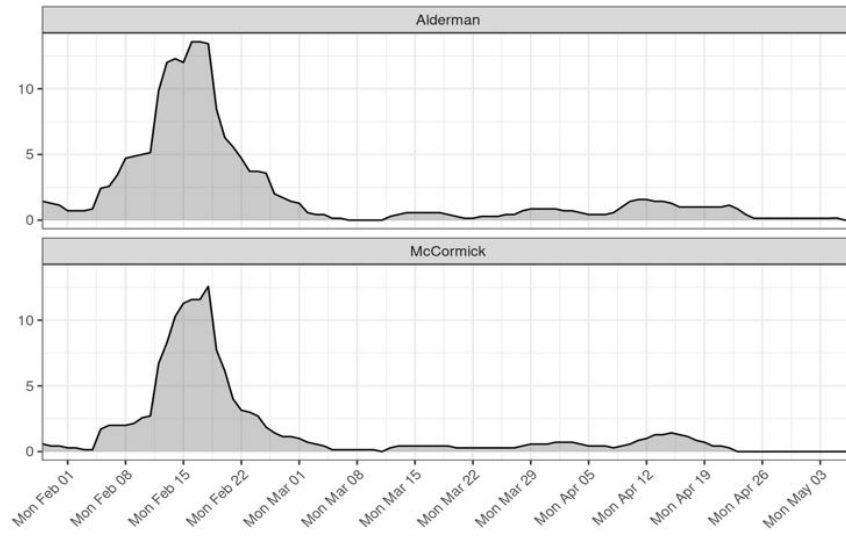
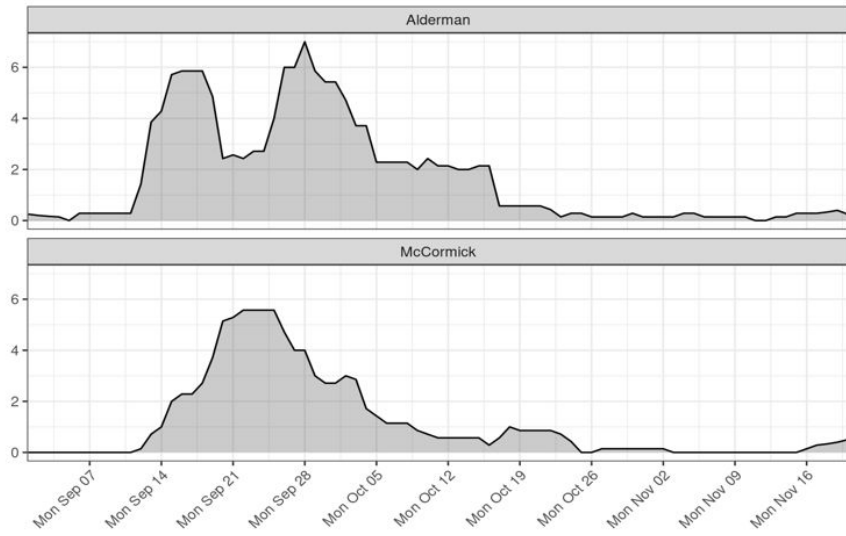
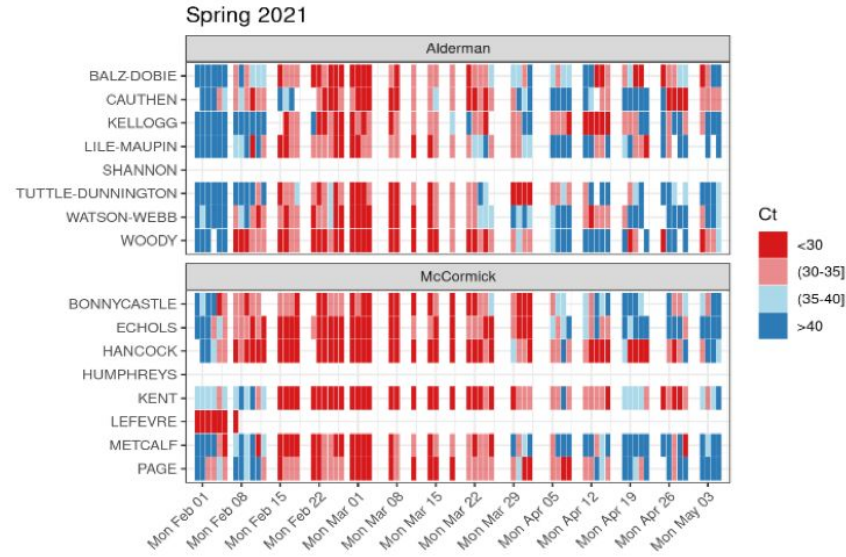
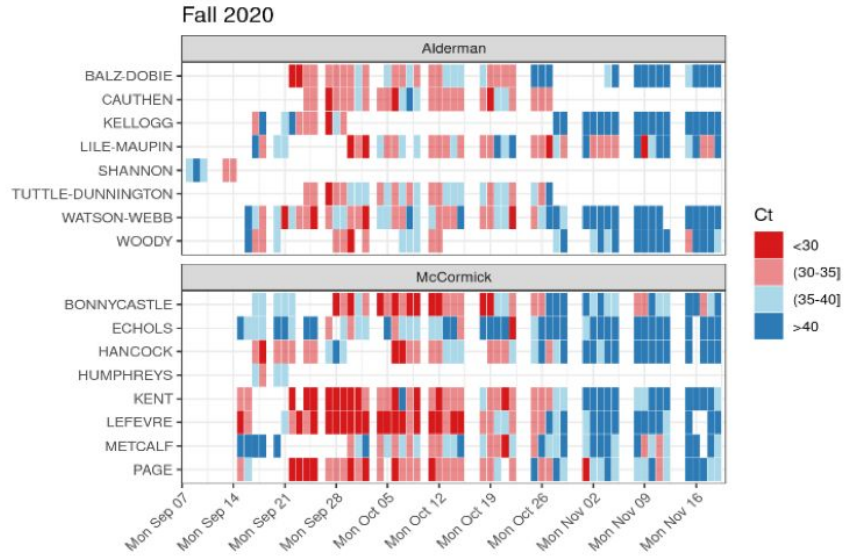
Not monitored  
+Humphreys  
+Emmett



\*Could be positive from prior shedding or new cases and trending results

Hall style dorms ~100-110 occupants

# The use of the data was different in Fall and Spring



Mike Porter, Kawai Tanabe,  
Chris Holstege and Shireen  
Kotay

## Summary

**WW surveillance may have many use cases but looking forward to further method development and refinement through collaborations**

**Passive pooled testing approach at the building level with the advantage of not having all the specimen management issues and assisting in using targeted resources**

**WW surveillance may also be adapted for understanding AMR genes of consequence into the environment**

# Too many thanks to fit on a slide

## Research Lab

Katie Barry  
Hardik Parikh  
Shireen Kotay  
Limor Steinberg

## School of Engineering

Lisa Colosi Peterson  
Erica Loudermilk  
Will Guilford

## VDH/DCLS

Denise Bonds  
Denise Toney  
Lauren Turner  
Logan Fink

## IT

Rena Morse  
David Taylor

## Facilities

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Paul Zmick  
Derrick Wilson  
Tom Harkins  
Rollie Zumbrunn

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Pace Lochte  
JJ Davis  
Susan Davis  
Chris Holstege  
Costi Sifri  
Mitch Rosner  
Michael Marquardt  
Kawai Tanabe  
Mike Porter

## LVG

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Mike Straightiff  
Bob Creeden

## UVA Clinical Laboratory

Melinda Poulter  
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Jen DeArment  
Lynne Foster  
April Attai  
Emily Snavely  
Joanne Carroll  
Dawn Dirks  
Jim Bowden  
Stacie Edmonds  
Gwen Ferguson  
Gayle Usher  
Frankie Brewster  
Lynn Hamilton  
Adam Rhodenizer  
Phoebe Gaither  
Randy Vandevander  
Dawn Burris

Many more microbiology, molecular and specimen management laboratory technologists and staff





# Questions/ Open Discussion

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thank  
you!

See you Soon!

Send inquiries / topics to:  
[rekha.singh@vdh.virginia.gov](mailto:rekha.singh@vdh.virginia.gov)