Wastewater-based genomic and epidemiological monitoring of SARS-CoV-2, South Africa, 2020-2021

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NATIONAL INSTITUTE FOR COMMUNICABLE DISEASES Division of the National Health Laboratory Service

Outline

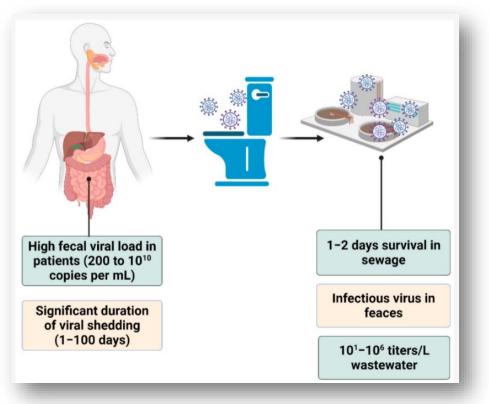
- Background
 - Principles and rationale for using wastewater-based epidemiology (WBE) for SARS-CoV-2 surveillance
 - The SACCESS network partners and sites
- Methodology
 - PCR detection and quantitation methodology
 - RNA sequencing methodology
- Results
- Conclusion







- SARS-CoV-2 replicates in cells lining respiratory and gastroenteric systems
- Whilst there is no evidence for faeco-oral transmission of SARS-CoV-2, the virus is shed in faeces
 - Viral load 10³-10⁷ copies/ml stool
 - 0-45 days after symptom onset,
 - 23% of persons with negative RT-PCR on throat swab will have positive stool



Amereh et al. Sewage Systems Surveillance for SARS-CoV-2: Identification of Knowledge Gaps, Emerging Threats, and Future Research Needs *Pathogens* 2021, *10*(8), 946; <u>https://doi.org/10.3390/pathogens10080946</u>

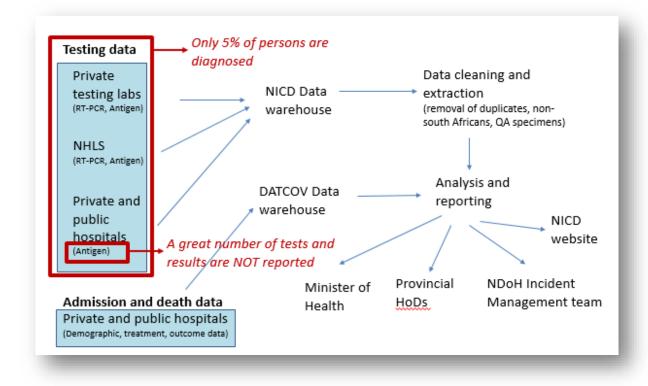








- Wastewater testing
 - overcomes the limitations of clinical surveillance
 - allows epidemiologists to infer the burden and distribution of SARS-CoV-2 in populations without the need for clinical testing,
 - Is much cheaper than clinical testing











- Wastewater testing
 - Can be used to identify SARS-CoV-2 variants that are circulating in populations under surveillance

		ELSEVIER	Available online at www.sciencedirect.com ScienceDirect	Current Opinion in Environmental Science & Health			
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	water		Ĩ				
	Article Key SARS-CoV-2 Mutations of Alpha, Gamma, and Eta Variants Detected in Urban Wastewaters in Italy by Long-Read Amplicon Sequencing Based on Nanopore Technology						
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- We aimed to show
 - <u>temporal and geographical trends in SARS-CoV-2 levels and</u> <u>sequences from wastewater</u>
 - and compare these with case loads and clinical SARS-CoV-2 sequences
 - to demonstrate the usefulness of wastewater-based epidemiology for SARS-CoV-2 as a surveillance tool.



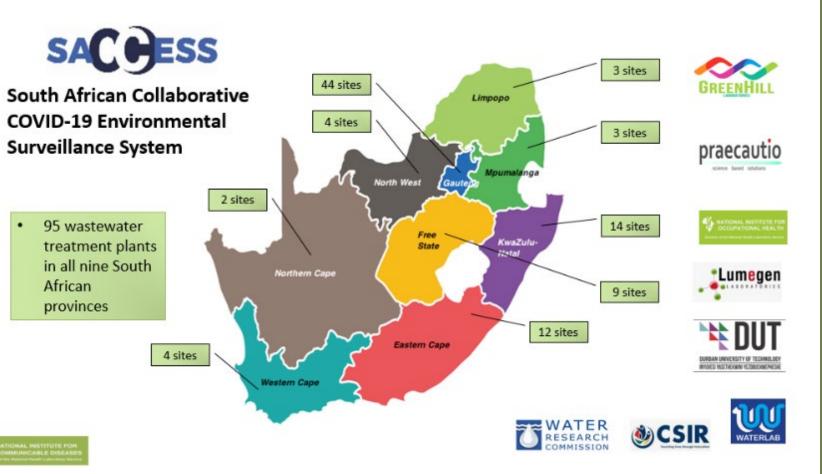






Wastewater based epidemiology in South Africa Methodology

- 7 laboratories
- 95 wastewater treatment plants
- All doing
 - Concentration
 - Qualitative PCR detection
- Genome sequencing (NICD only)
 - Proof of concept established on results from 17 specimens







Wastewater based epidemiology in South Africa

Methodology



Name of Laboratory partner	Method for virus concentration	Method for nucleic acid extraction	RT-PCR assay	Quantification
NICD	Centricon [®] Plus-70 centrifugal	QIAamp® viral RNA mini kit	Allplex™ 2019-nCoV Assay	EDX SARS-CoV-2 Standard including RNA transcripts of E, N and RdRP genes
GreenHill / Praecautio	Amicon [®] Ultra-15 Centrifugal Filter)	Omega Bio-Tek Mag-Bind® Viral DNA/RNA 96 Kit	CDC 2019-Novel Coronavirus (2019-nCoV) Real-Time RT-PCR Diagnostic Panel	Relative quantification based on CDC 2019-Novel Coronavirus (2019-nCoV) Real-Time RT-PCR Diagnostic Panel
NIOH	Skim milk flocculation	MagMAX Viral and Pathogen Nucleic Acid Isolation Kit	TaqPath COVID-19 CE-IVD RT-PCR Kit Thermo Fisher	Standard curve method using TaqPath kit positive control
WaterlLab/UP	Skim milk flocculation	QIAamp® Ultrasens® Virus kit	Allplex™ 2019-nCoV Assay	Standard curves using the 2019_nCoV_N positive control plasmid (Integrated DNA Technologies, Inc, Coralville, IA)
SAMRC-TB PLATFORM	Centrifugation	ZymoBiomics RNA Extraction Kit	AllPlex 2019-nCoV Assay	EDX SARS-CoV-2 Standard including RNA transcripts of E, N and RdRP genes
Lumegen	Passive sampling + resuspension in PBS	MN DNA/RNA pathogen extraction Kit	TaqPath COVID-19 CE-IVD RT-PCR Kit (Thermo Fisher)	5-point standard curve of the TaqPath positive control
CSIR	Polyethylene Glycol	Omega Bio-tek ENZA total RNA Kit II	2019-nCoV CDC EUA Kit	Relative quantification based on the 2019-nCoV CDC positive control



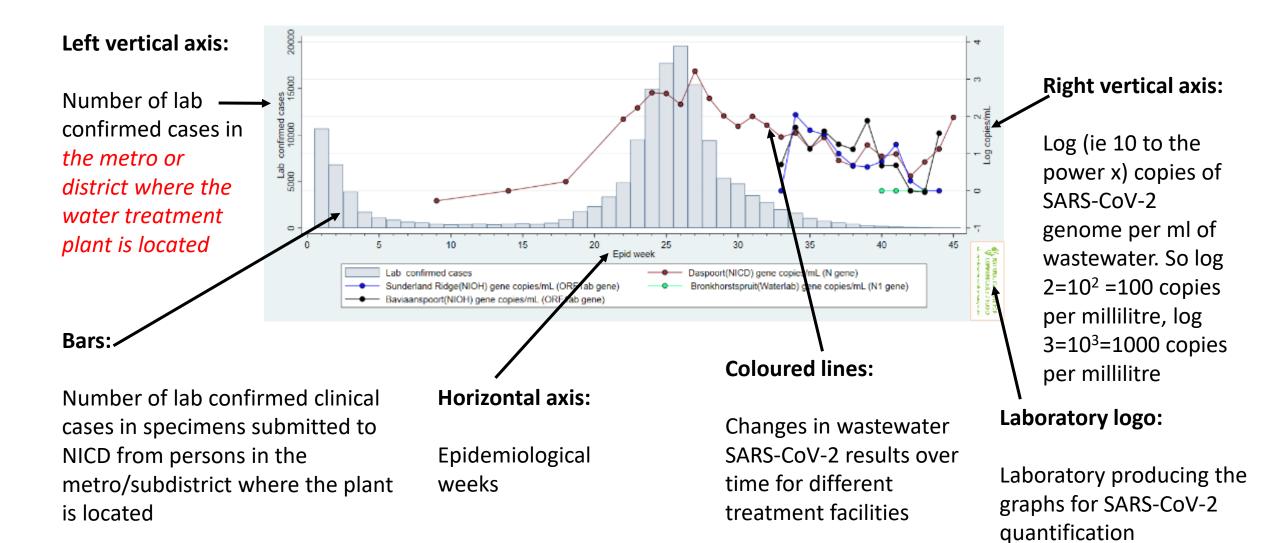




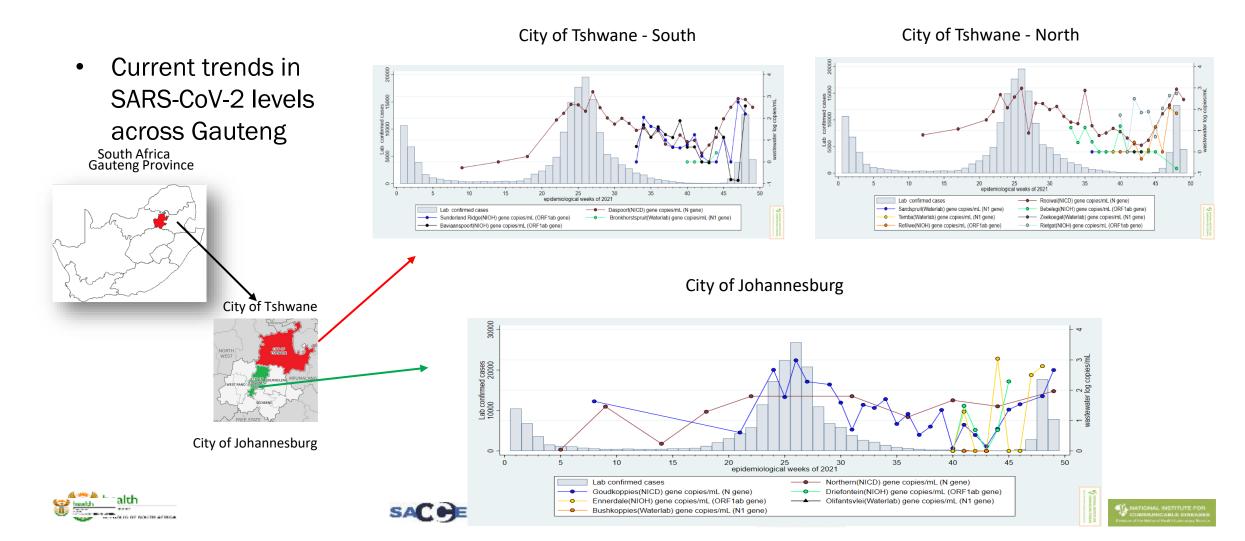


Wastewater-based Epidemiology for COVID-19

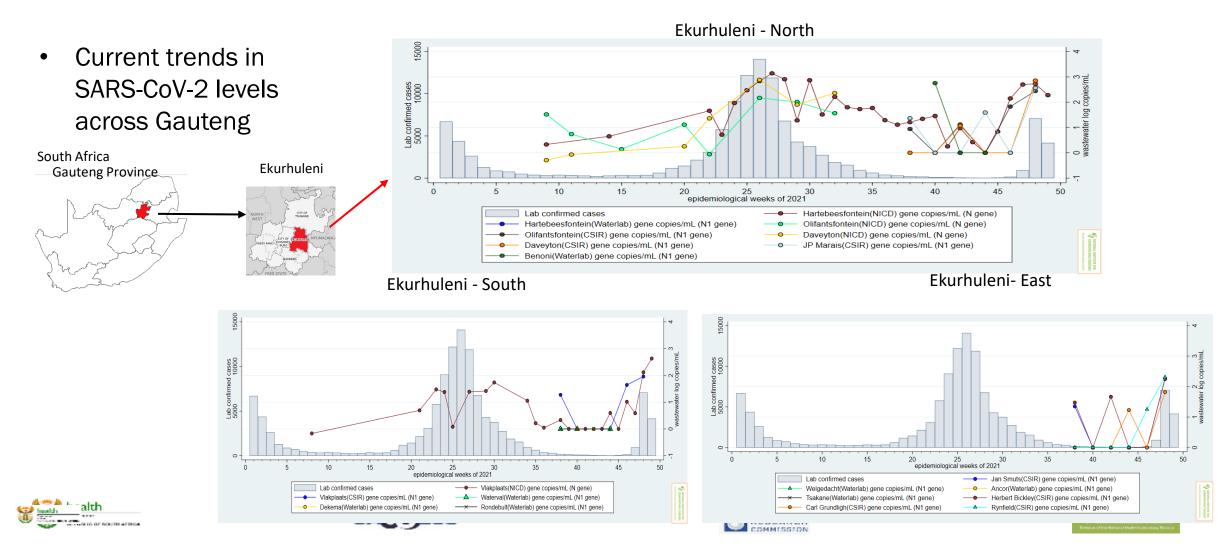
Interpretation of quantitative results

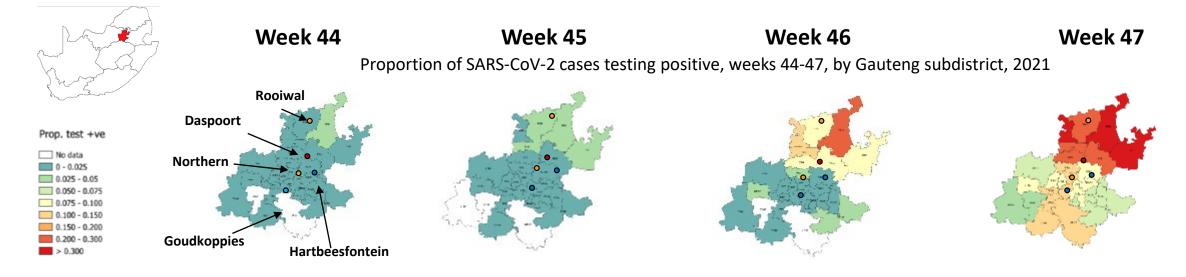


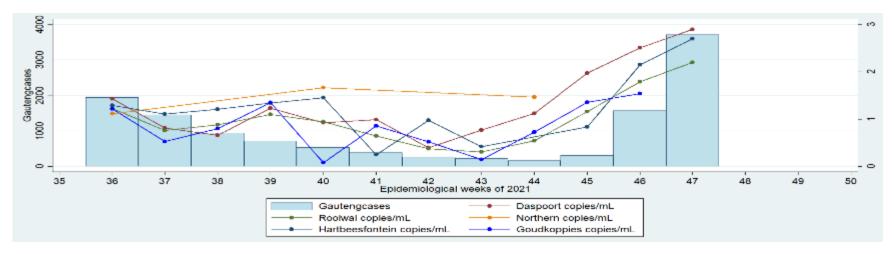
Gauteng wastewater treatment plants



Gauteng wastewater treatment plants



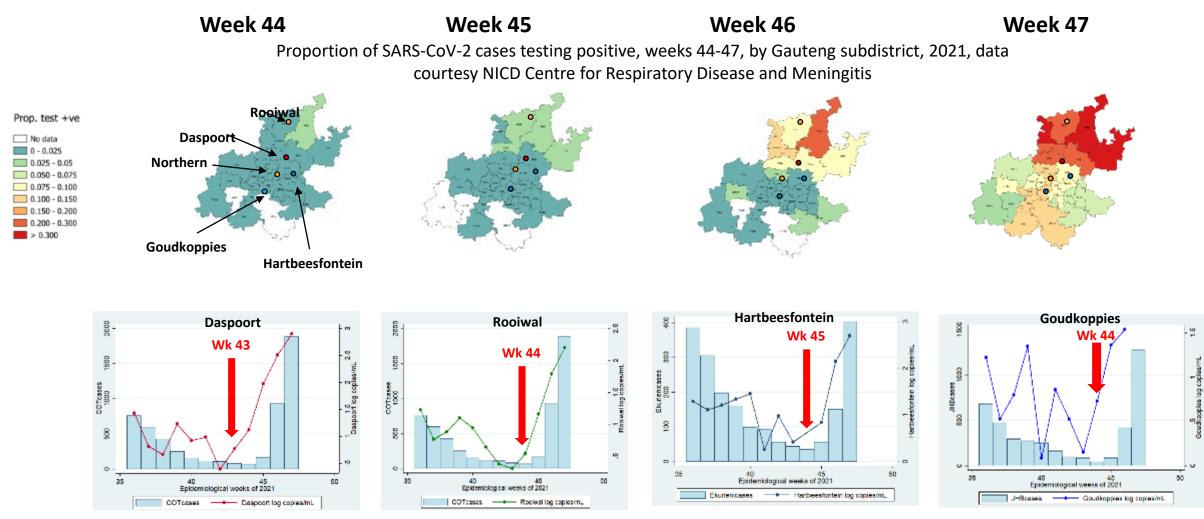






Wastewater evels (lines) at selected treatment plants, by total Gauteng cases, epi weeks 36-47, 2021



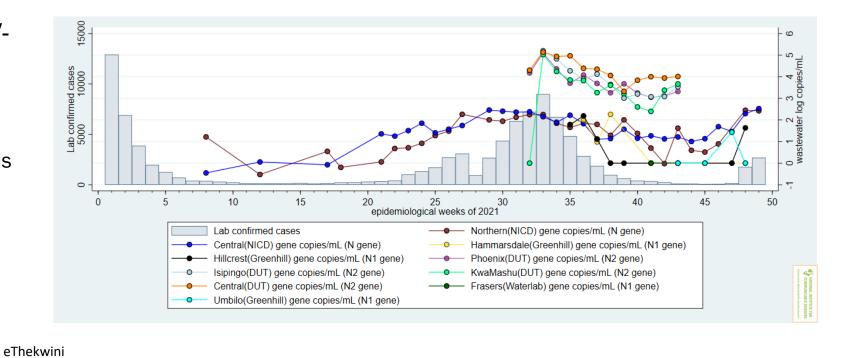


Wastewater levels (lines) at selected treatment plants, by total cases in each Gauteng district where the treatment plant is situated, epi weeks 36-47, 2021 Data courtesy NICD Centre for Vaccines and Immunology

Ethekwini wastewater treatment plants

- Current trends in SARS-CoV-2 levels across eThekwini wastewater plants
 - 2 plants
 - Ongoing upward trend which is levelling

South Africa









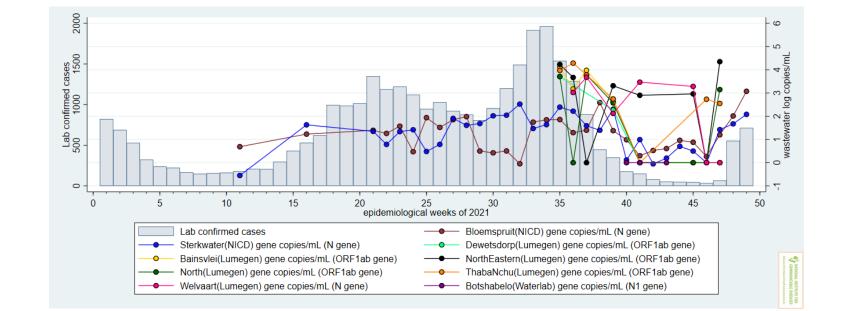
Mangaung waste water treatment plants

- Current trends in SARS-CoV-2 levels across Mangaung wastewater plants
 - 2 plants
 - One plant with increasing levels
 - One plant with decreasing levels



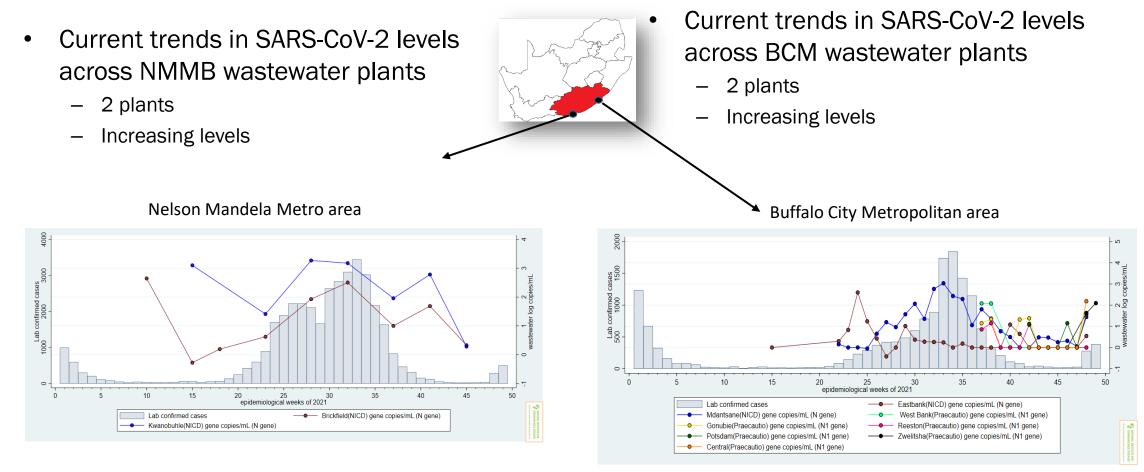








Eastern Cape wastewater treatment plants



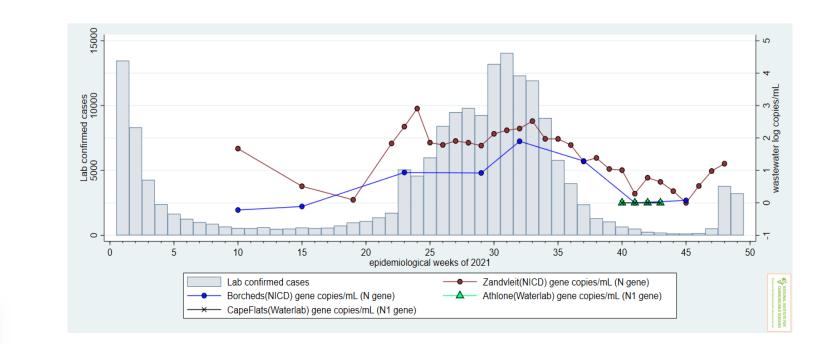






City of Cape Town wastewater treatment plants

- Current trends in SARS-CoV-2 levels across City of Cape Town wastewater plants
 - 2 plants
 - Stable/increasing levels





City of Cape Town







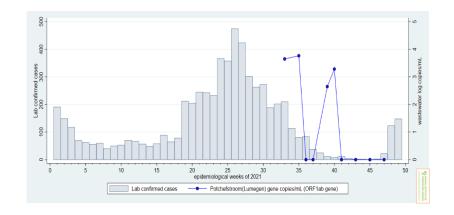


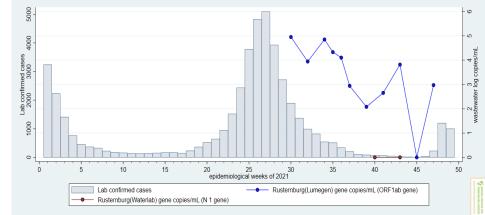
North west wastewater treatment plants

JB Marks - Potchefstroom

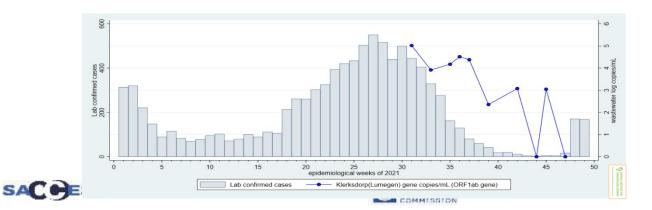
Bojanala - Rustenberg

- Current trends in SARS-CoV-2 levels across North West Province
 - 3 plants
 - Increasing levels





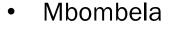
Matlosana - Klerksdorp

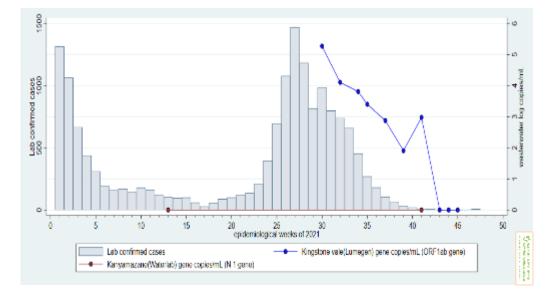


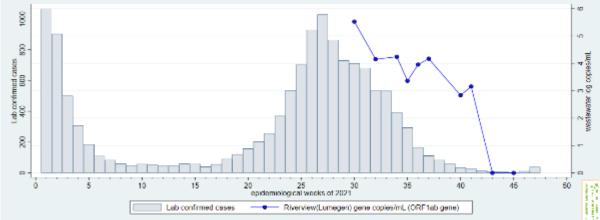


Mpumalanga wastewater treatment plants

• Emalahleni







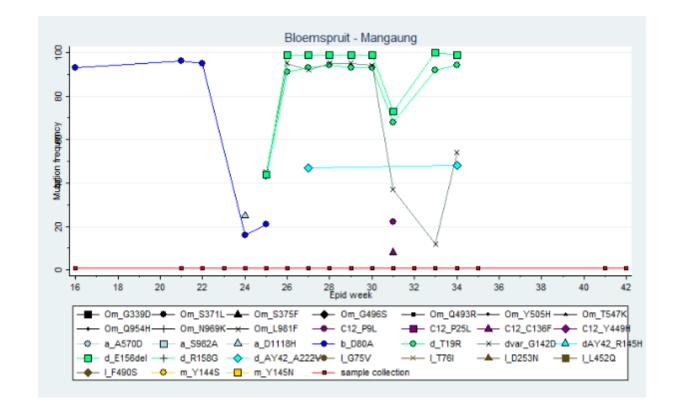






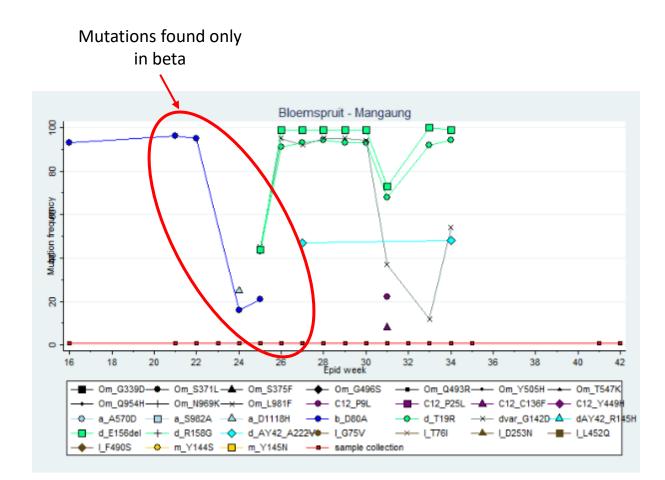


- Full genome sequencing of RNA fragments present in wastewater
 - Allows for detection of single nucleotide polymorphisms (SNPs compatible with different variants)
 - Tracking of variants and read frequency of mutations over time



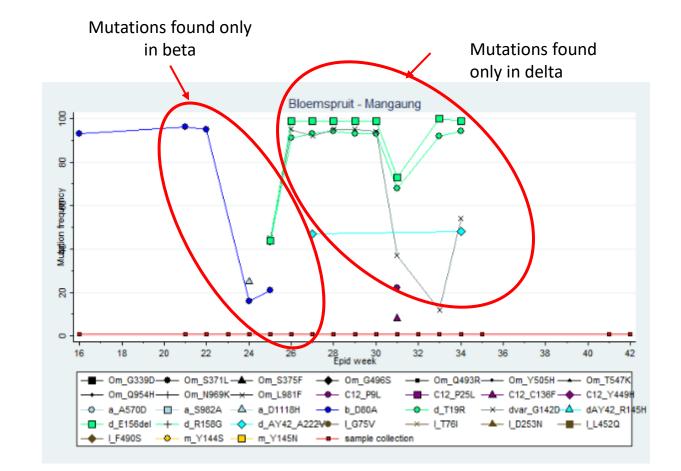


- Mangaung
 - Week 16,
 - Beta dominance
 - C.1.2
 - Week 22
 - Delta appeared and became dominant



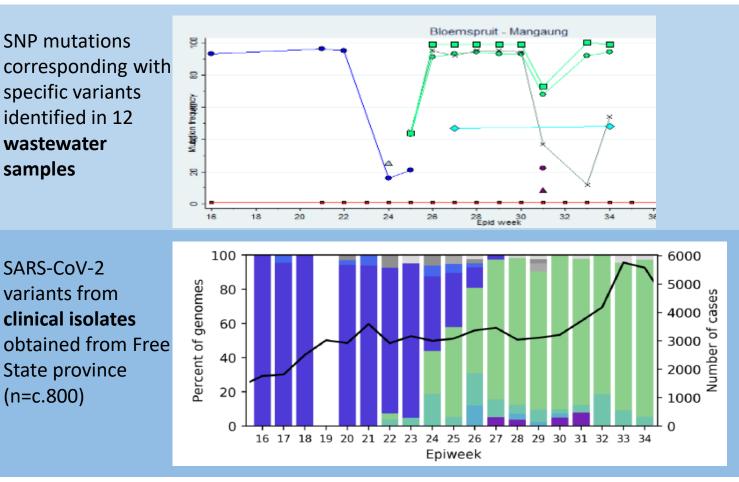


- Mangaung
 - Week 16,
 - Beta dominance
 - C.1.2
 - Week 22
 - Delta appeared and became dominant





- Mangaung Bloemspruit WWTP
 - Week 16,
 - Beta dominance
 - C.1.2
 - Week 22
 - Delta appeared and became dominant
- Good correspondence
 with clinical genomics
 and lineages



Clinical isolates figure courtesy CRDM, NICD Jinal Bihman



Conclusion

- Amongst sentinel wastewater treatment plants in South Africa
 - Quantitative SARS-CoV-2 viral load in genome copies/ml demonstrates concordance with burden of disease as determined by clinical testing
 - Sequencing of SARS-CoV-2 RNA in wastewater demonstrates the presence of single nucleotide polymorphisms compatible with variants detected in clinical specimens over time





Conclusion

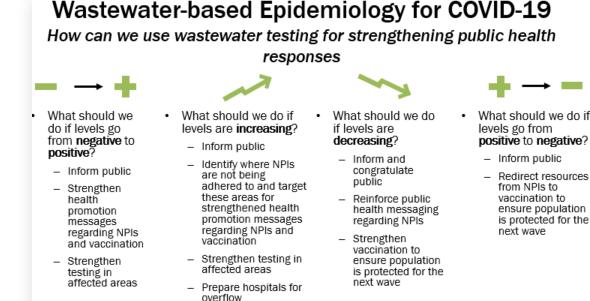
- Amongst sentinel wastewater treatment plants in South Africa
 - Quantitative SARS-CoV-2 viral load in genome copies/ml demonstrates concordance with burden of disease as determined by clinical testing
 - Sequencing of SARS-CoV-2 RNA in wastewater demonstrates the presence of single nucleotide polymorphisms compatible with variants detected in clinical specimens over time

- Wastewater-based epidemiology methods successfully detected the 4th wave in Gauteng
- Determination of the presence of omicron variant of concern in wastewater in SACCESS network laboratories is awaited, but has been identified in Cape Town wastewater treatment plants



Way forward

- Integrate wastewater-based epidemiological surveillance findings into clinical surveillance networks
- Make results accessible through web-based dashboard
- Determine thresholds and triggers for action based on wastewater levels
- Improve turn-around times for genomic sequencing







Acknowledgements

- Water Research Commission
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- Partner labs





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