

Wastewater-based genomic epidemiology for SARS-CoV-2 surveillance in South Africa

17 May 2024

Sample collection dates up to 10 May 2024 (Epidemiological week 19)

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Summary: SARS-CoV-2 transmission and genomics based on evaluation of wastewater at sentinel sites across RSA

Wastewater <u>levels</u> Epidemiological weeks 48 (2023) –19 (2024)

- From weeks 48-52 the cumulative SARS-CoV-2 levels measured at wastewater treatment works (WWTW) has remained around one log genome copies/ml of wastewater. This has followed on from the increases observed in weeks 31-41, when the cumulative SARS-CoV-2 levels in wastewater at sentinel sites in South Africa showed increases to levels above 2 log (100) genome copies/ml of wastewater, up from below one log copy/ml in epidemiological week 22 (first week in June 2023). From week 45, levels measured below 2 log genome copies/ml of wastewater.
- In Epi weeks 1-6 lower levels were observed nationally, with an increase to above 2 log (100) genome copies/ml seen in week 7 and a subsequent decrease to about one log genome copies/ml of wastewater week 8. Levels increased to moderate in Epi week 10. Levels increased to high (above 2 log (100) genome copies/ml) in Epi week 14 and remain high in Epi week 16. Levels drop slightly to moderate in Epi week 17 and have decrease further in Epi week 19.
- Correlation with syndromic surveillance for influenza-like illness (ILI) and severe
 acute respiratory infection findings (SARI) is required to determine the clinical and
 public health significance of ongoing transmission. https://www.nicd.ac.za/diseases-a-z-index/disease-index-covid-19/surveillance-reports/weekly-respiratory-pathogens-surveillance-report-week/

Wastewater <u>genomics</u> Epidemiological weeks 48 (2023) – 17 (2024)

- Omicron lineage BA.2.86.X, and JN.1.X sub-lineages were the dominant lineages circulating in wastewater samples between January 2024 April 2024 (Epiweek 1-10)
- In clinical samples, **BA.2.86** was also the dominant lineage circulating throughout November 2023 and January 2024, followed by **XBB.1.5***, **XBB.1.19*** and **JN.1***.
- The Omicron lineage BA.2.86, JN.1* and Recombinant lineages are circulating in KwaZulu-Natal in eThekwini (in the catchments of Northern and Central WWTWs). In Gauteng, in the City of Johannesburg (in catchments of Northern and Goudkoppies WWTWs), in Ekurhuleni (in the catchments of Olifantsfontein, Vlakplaats, and Hartebeesfontein WWTWs), and the City of Tshwane (in the catchment of Daspoort WWTP) BA.2.86.X and JN.1.X are the dominantly circulating. It is also circulating in Eastern Cape in Buffalo City (in the catchment of Mdantsane WWTW), in Western Cape, in the City of Cape Town (in the catchment of Borches Quarry WWTW), Free State, in Mangaung (catchments Bloemspruit and Sterkwater WWTWs) and North West (Boitekong). In Limpopo (Musina) and Mpumalanga (Kingstonvale) BA.2.86.X lineage was the main lineage detected.

Interpretation: Ongoing transmission of SARS-CoV-2 due to Omicron lineages including the new BA.2.86 and JN.1* lineages.



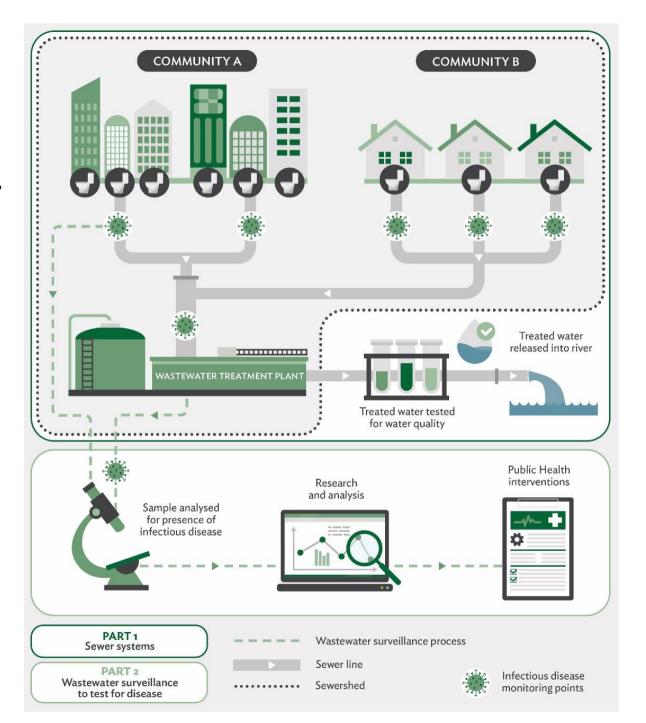
Wastewater-based Epidemiology for COVID-19 How is wastewater tested for SARS-CoV-2?

For a full description of this process, see the photo essay developed in collaboration with the Gauteng City Region Observatory https://www.gcro.ac.za/outputs/photo-essay-sewersheds-what-can-wastewater-tell-us-about-community-health/

For a technical description and analysis of wastewater levels and results see

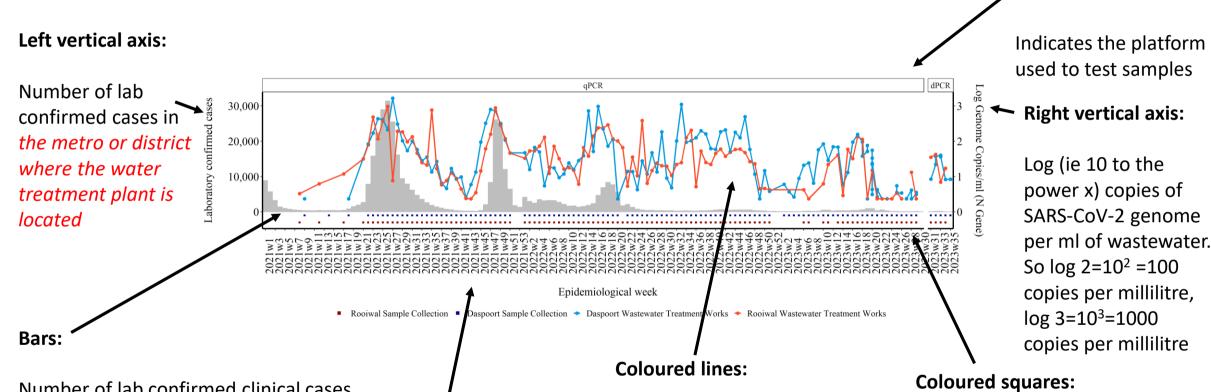
https://pubmed.ncbi.nlm.nih.gov/37506905/ https://www.medrxiv.org/content/10.1101/2022.12.15.22283506v1 (accepted by Nature Communications, publication pending)

SARS-CoV-2 is not transmitted by faeco-oral route. Wastewater with SARS-CoV-2 is not infectious



Wastewater-based Epidemiology for COVID-19

What does wastewater testing for SARS-CoV-2 mean?



Number of lab confirmed clinical cases in specimens submitted to NICD from persons in the metro/subdistrict where the plant is located

Horizontal axis:

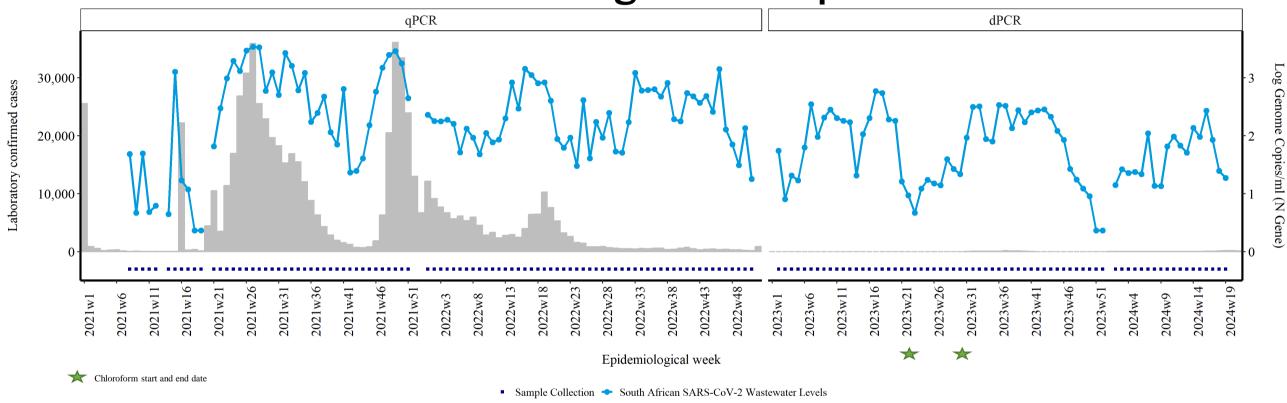
Epidemiological weeks from 2021 to 2023

Changes in wastewater SARS-CoV-2 results over time for different treatment facilities

Epi weeks during which samples were collected

Facets:

South Africa at a glance: Summed total of clinical and genome copies



Changes in levels of SARS-Cov-2 in log-transformed genome copies/ml (right vertical axis, line graph) in in-flowing untreated wastewater from plants tested by NICD, compared with <u>laboratory-confirmed cases from Tshwane, Johannesburg, Ekurhuleni, eThekwini, Mangaung, Nelson Mandela, Buffalo City, City of Cape Town, Vhembe, Ehlanzeni and Bonjanala Platinum (left vertical axis, grey bars)</u>, by epidemiological week, 2021-2023. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from qPCR to dPCR occurred for SARS-CoV-2 testing.





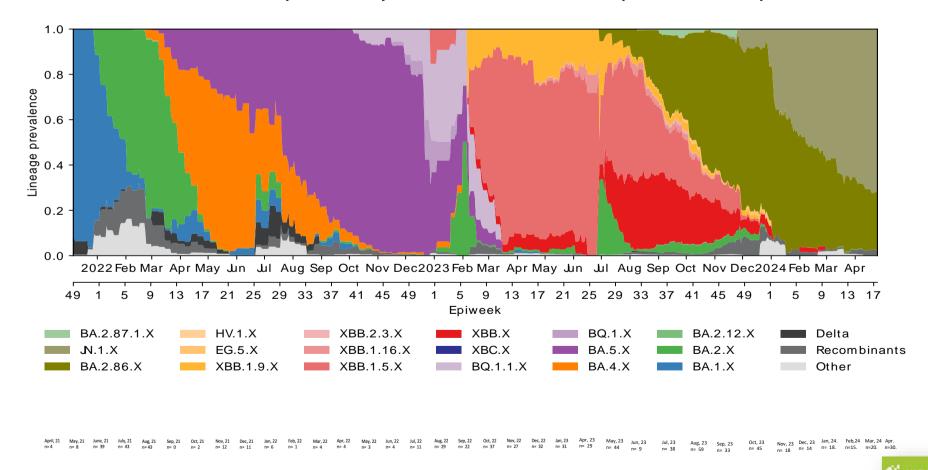
South Africa at a glance: Circulating variants as determined by Freyja deconvolution of sequence data

 SARS-CoV-2 variants in wastewater as determined by the 'Freyja' tool (Scripps

Institute)

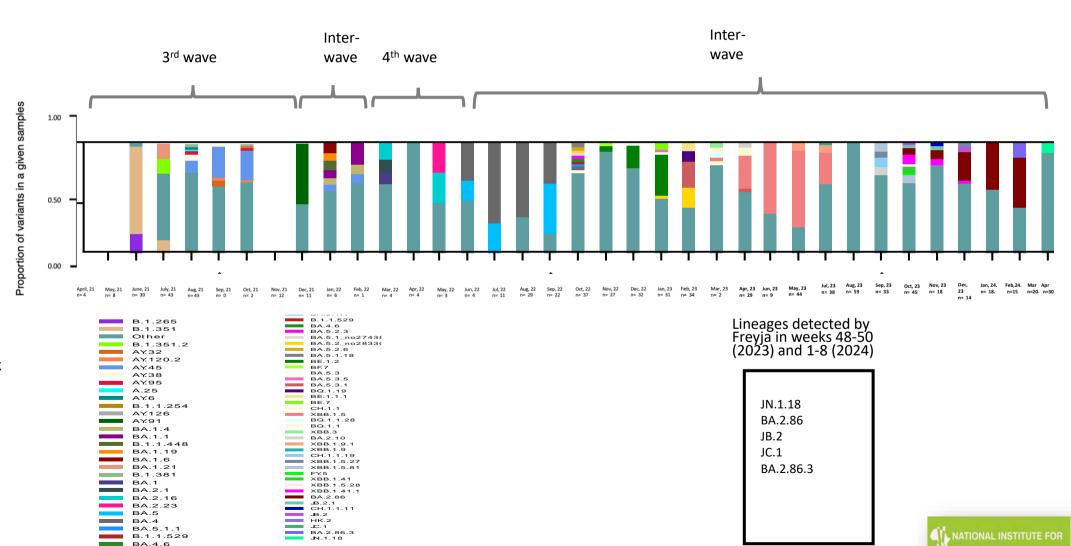
- Allows determination of variants in each wastewater sample
- Results from sequencing data ending in epi week 17 (April 2024)
 - Omicron lineages BA.2.86.X and JN.1.X were circulating from December to May, with Recombinants and XBB.X sublineages circulating at low proportion.

Inferred variants in wastewater samples from South African wastewater treatment plants by month, between April 2021- April 2024

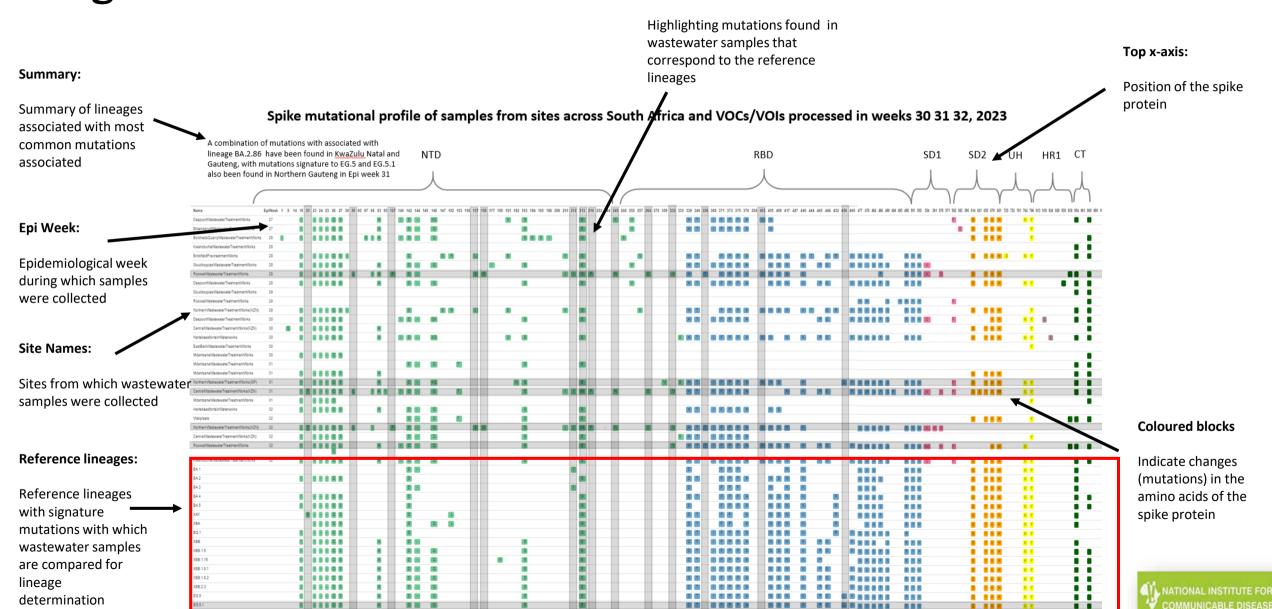


South Africa at a glance: Circulating <u>lineages</u> as determined by Freyja deconvolution of sequence data

- Results from sequencing data ending in epi week 17 (April 2024)
 - Omicron lineages BA.2.86 and JN.1 were circulating from January to March with BA.2.86 dominance.
 - Omicron XBB sublineages were in circulation throughout.
 - The predominant lineage circulating in clinical samples in the recent week is BA.2.86 and JN.1 sublineages.



What mutations in the spike protein tell us about the circulating lineages



Spike mutational profile of samples from sites across South Africa and VOCs/VOIs processed in weeks 15 - 17, 2024

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A. combination of mutations associated with lineage BA.2.86, JN.1 and XBB sublineages have been found in sites across all South African provinces during Epi weeks 15-17. Majority of mutations are signature to JN.1 and BA.2.86



Spike mutational profile of samples from sites across South Africa and VOCs/VOIs processed in weeks 15 - 17, 2024

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A. combination of mutations associated with lineage BA.2.86, JN.1 and XBB sublineages have been found in sites across all South African provinces during Epi weeks 15-17. Majority of mutations are signature to BA.2.86.X and JN.1.X



Spike mutational profile of samples from sites across South Africa and VOCs/VOIs processed in weeks 15 - 17, 2024

																				Г																		
SiteName	EpiW	eek	p679	p680	pć	581 pć	697	p704	p764	p778	p795	p796	p799	p812	p815	p848	p855	p869	p870	- 1	p939	p950	p954 p96	9 p1002	p1027	p1086	p1088 p1094	p1095	p1104	p1117 p	p1124	p1143 p11	144 pl1	.78 pll	90 p1228	8 p1234	p1240	p1249
Central Wastewater TreatmentWorks(KZN)	15	K	•		R			•	•	•	•	Y .		•							F	Н	K						ī.		L							
Goudkoppies Wastewater Treatment Works	15	K		T	R	I		K				Y									F	N H	K								L				L			
ERWAT Vlakplaat Wastewater TreatmentWorks	15	K			R			K				Y								- 1	F	Н	K								L							
Komatipoort Sewage plant	15	K			R			K				Y								- 1	F	Н	K						L		L							
Bloemspruit Wastewater TreatmentWorks	15	K			R			K				Y									F	Н	K								L							
Zandvleit Wastewater Treatment Works	15	K			R							Y								- 1			K															
Borcherds Quarry Wastewater Treatment Works	15	K			R							Y								- 1	F	Н	K			0			L		L							
Daspoort Wastewater Treatment Works	15	K			R			K				Y								- 1	F	H	K						ī		L							
Mdantsane Wastewater Treatment Works	15	-						-				-								- 1																		
Daspoort Wastewater Treatment Works	16	K			R			K				v v	7								F	н	K						T.		L							P
Northern Wastewater Treatment Works(GP)	16	K			R		1	K				v								1		н	K						-		L							
Central Wastewater Treatment Works(KZN)	16	K			P		-	K				v			9					- 1	,	Н Н	V.			,			1		L			V				
Northern Wastewater Treatment Works(KZN)	16	K			R			K				v										н н	K						ī		L							
Hartebeesfontein Waterworks	16	K K			P			K				v				v		T			F	н н	V.		9				L		L				L			
Goudkoppies Wastewater Treatment Works	16	V.			D.							v				1		1			r		V.		3						L							
ERWAT Vlakplaat Wastewater Treatment Works	16	v			D			K	A			v								- 1		u u	r.	п						T.	Ī.							
Kingstonvale	16	V.		v	D.			N.	А	1		v									(п п	V.	п						V	ī							
Komatipoort Sewageplant	16	v		1	D.							v									,	п п	V.								ī							
Bloemspruit Wastewater Treatment Works	16	K.			D.							ı v		e							1	п	Λ.			L		2			ī							
Sterkwater Wastewater Treatment Works	16	K.			R.			77				1		3							1 1	n 	7								1							
Borcherds QuarryWastewaterTreatmentWorks	10	K.			R D			K.				1	,	5							1	H	Y.								ī						F	
Zandvleit Wastewater Treatment Works	10	K			K			K													F	H	K			l.			L		L						1	
	16																						_															
Mdantsane Wastewater Treatment Works	10	K			K							1							_		F	H	K								I.							
Daspoort Wastewater Treatment Works	16	K			K							Y					L		T		F	Н	K		1	l.			L		L							
Rustenburg Wastewater Treatment Works	16	K			R			K				Y									F	H	K								L							
Boitekong	16	K			R							Y									F	H	K										ע					
Brickfield Pre-treatment Works	16	K			R							Y									F	H	K								L							
Central Wastewater Treatment Works(KZN)	17	K			R							Y									F	H	K			l			L	I	L		_					
Rooiwal Wastewater Treatment Works	16	K			R							Y									F	H	K								L		ע					
Daspoort Wastewater Treatment Works	17	K			R																F	H	K						L		L							
Northern Wastewater TreatmentWorks(KZN)	17	K		P	R							Y									F	H	K		1	l			L		L							
Komatipoort Sewageplant	17	K			R							Y											K															
Goudkoppies Wastewater TreatmentWorks	17	K			R							Y									F	H	K			l.	L		L		L				L			
Kingstonvale	17	K			R							Y									F	H	K								L	K						
Bloemspruit Wastewater Treatment Works	17	K			R							Y		S							F	H	K						L		L							
Sterkwater Wastewater Treatment Works	17	K			R							Y									F	H	K						L		L							
Mdantsane Wastewater Treatment Works	17	K			R							Y									F	H	K						L		L							
Rooiwal Wastewater Treatment Works	17	K			R							Y									F	H	K								L							
Borcherds Quarry Wastewater Treatment Works	17	K			R							Y									F	H	K						L		L							
Rustenburg Wastewater Treatment Works	17	K			R							Y									F	Н	K															
Variants																																						
BA.2		K			H			K				Y										H																
BA 2 XBB.1.5 XBB.1.16		K			H			K				Y										H	K															
XBB.1.16		K			H			K				Y										Н	K															
XBB.1.9.1		K			H			K				Y										н	K															
XBB.1.9.2		K			H			K				Y										н н	K															
XBB.2.3		K			H			K				Y										н	K															
XBB.2.3 EG.5		K			H			K				Y										11	K															
EG.3.1		K			Н			K				Y								\rightarrow		H	K															
BA.2.86		K			R			K				Y									F	H H	K.															
JN.1		K			R			K				Y									1	H H																
H												-									1	н	7															

A. combination of mutations associated with lineage BA.2.86, JN.1 and XBB sublineages have been found in sites across all South African provinces during Epi weeks 15-17



Amino acid mutations and frequency — Spike protein

XBB* is a recombinant of BA.2.10.1 and BA.2.75 that is characterised by one or more of the following mutations in the spike protein: V83A, Y144-, H146Q, Q183E, V213E,G252V, G339H, R346T, L368I, V445P, G446S, N460K, F486S, F490S

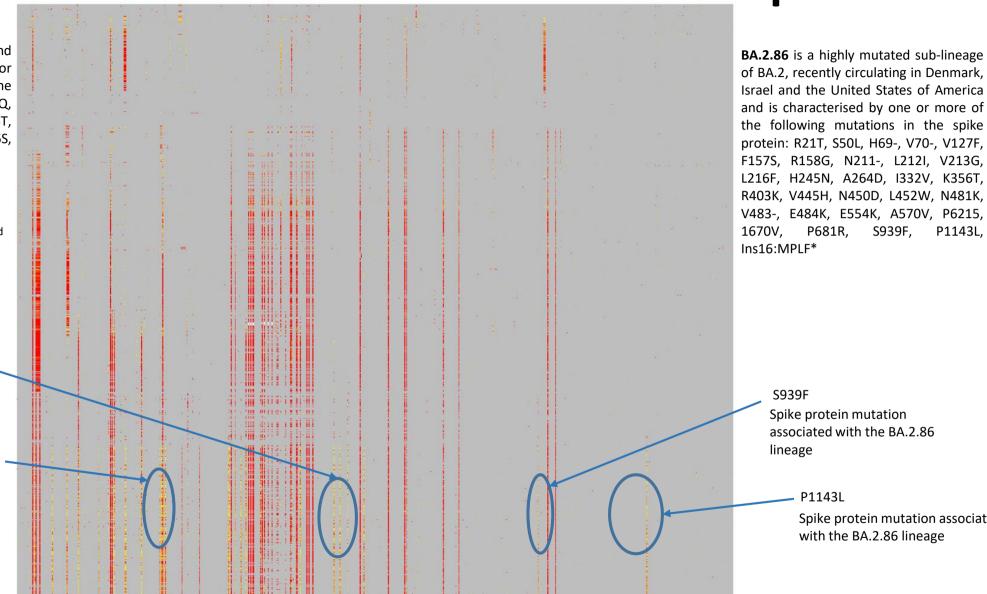
List of variants and sub-lineages of interest and

(https://www.who.int/en/activities/tracking-SARS-CoV-2-variants)

> E554K Mutations in spike protein associated with BA.2.86

> > V213E, R346T

Mutations in spike protein associated with XBB* sublineages



Heatmap showing patterns of emerging mutations in the spike region of SAKS-COV-2, collected from April, 2021 - May, 2024. Mutations appearing in yellow have a low read frequency, those appearing in orange have a medium read frequency and those appearing in red have a high read frequency. Mutations are included and updated weekly.



P681R,

S939F

P1143L

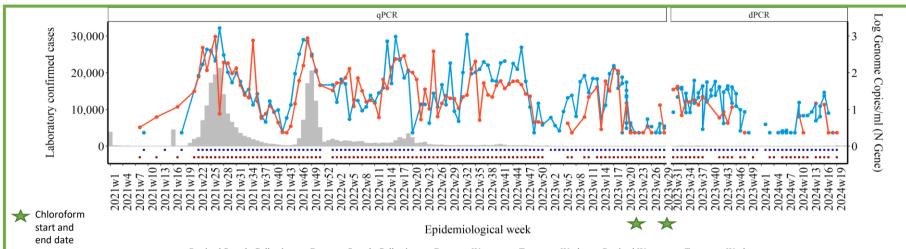
S939F.

Spike protein mutation associated

with the BA.2.86 lineage

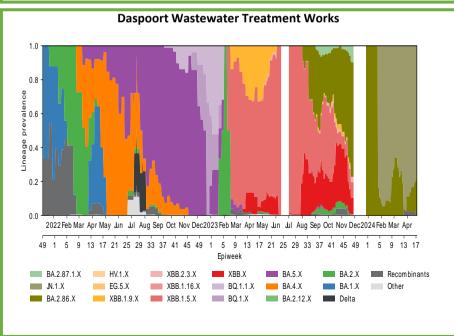
P1143L.

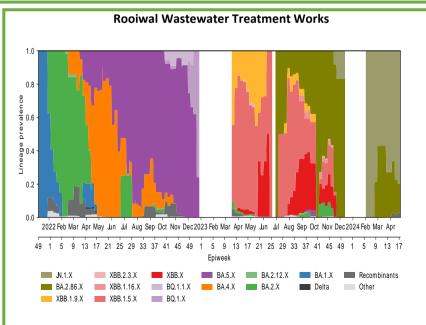
Gauteng - Tshwane



• Rooiwal Sample Collection • Daspoort Sample Collection • Daspoort Wastewater Treatment Works • Rooiwal Wastewater Treatment Works Wastewater Ievels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Tshwane (left

Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Tshwane (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from qPCR to dPCR occurred for SARS-CoV-2 testing.





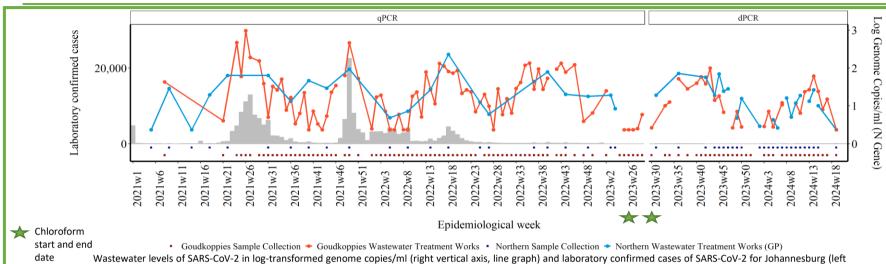
SARS-CoV-2 levels and Genomic Results in Epi week 19:

- SARS-CoV-2 levels in Daspoort WWTW increase to moderate in Epi week 14 and decrease to low again in Epi week 15. Levels increase to moderate in Epi week 16 and decrease to low in Epi week 17 and remain low in Epi week 19.
- SARS-CoV-2 levels in Rooiwal WWTW have decreased in Epi week 47. Levels remain low in Epi week 11 and increase to moderate in Epi week 14. Levels remain moderate in Epi week 16 and decrease to low in Epi week 17 and remain low in Epi week 19.
- * Sequencing data ending in Epi week 17 in Daspoort and in Rooiwal.
 - Omicron lineages JN.1.X and BA.2.86.X were circulating in Daspoort during Epi week 1 to 5. Recombinants were also detected at low proportions during Epi week 13.
 - BA.2.86.X and JN.1.X was also the dominant lineage detected in Rooiwal during Epi week 5-17, with JN.1.X sub-lineages dominating

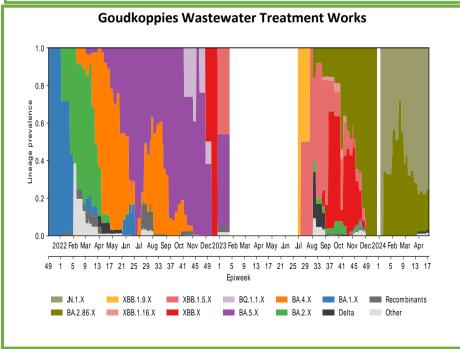
SNP Analysis:

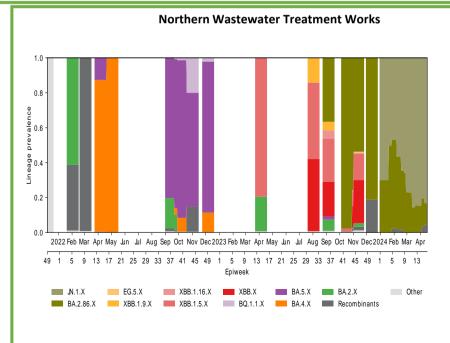
A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 and JN.1 were found in both Daspoort and Rooiwal.

Gauteng - Johannesburg



• Goudkoppies Sample Collection • Goudkoppies Wastewater Treatment Works • Northern Sample Collection • Northern Wastewater Treatment Works (GP) Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Johannesburg (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from qPCR to dPCR occurred for SARS-CoV-2 testing.





SARS-CoV-2 levels and Genomic Results in Epi week 19:

- SARS-CoV-2 levels in Goudkoppies WWTW increased from Epi week 5 to Epi week 7, but levels remain low. Levels increase to moderate in Epi week 12. Levels decreased in Epi week 15 but remain moderate. Levels decreased further in Epi week 16 to low. Levels increase slightly in Epi week 17 to moderate and decrease to low in Epi week 19.
- In Northern WWTW, SARS-CoV-2 levels increased to moderate in week 8 from low levels seen in week 6. Levels decreased to low in Epi week 9, followed by an increase to moderate in Epi week 10. Levels remain moderate in Epi week 14. Levels decrease to low in Epi week 15 and decrease further in Epi week 19.

* Sequencing data ending in Epi week 17 in Goudkoppies and 13 in Northern Wastewater Treatment Works.

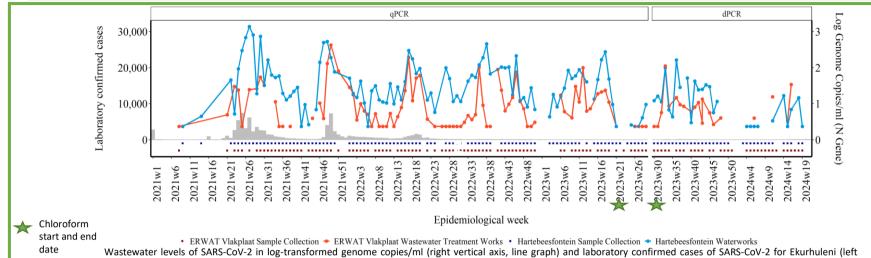
- JN.1.X and BA.2.8.X lineages were dominating during epiweek 1 - 17 in Goudkoppies. Recombinants and Other lineages were also detected in the recent weeks (at very low proportion)
- During Epi week 1 13 BA.2.86.X, JN.1.X lineages were circulating in Northern (GP). In Epi week 5. Recombinants were also observed.

SNP Analysis:

- A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 were found in Goudkoppies.
- A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 and JN.1were found in Northern Johannesburg.

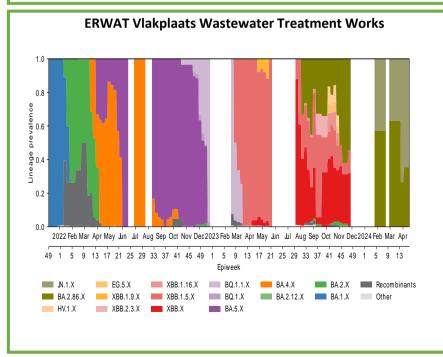
Division of the National Health Laboratory Service

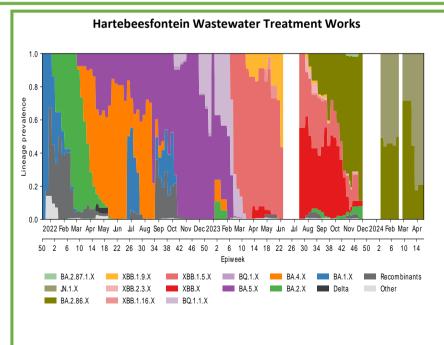
Gauteng - Ekurhuleni



• ERWAT Vlakplaat Sample Collection • ERWAT Vlakplaat Wastewater Treatment Works • Hartebeesfontein Sample Collection • Hartebeesfontein Waterworks

Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Ekurhuleni (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from qPCR to dPCR occurred for SARS-CoV-2 testing.





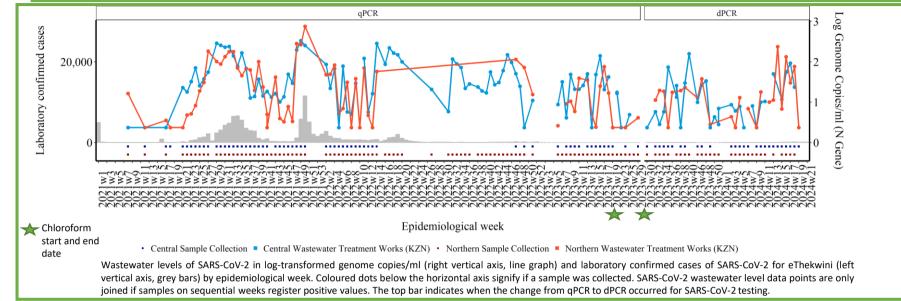
SARS-CoV-2 levels and Genomic Results in Epi week 19:

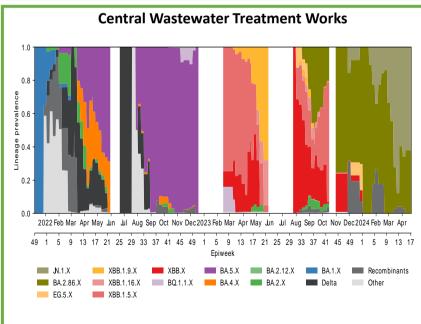
- The SARS-CoV-2 levels in Hartebeesfontein WWTW decreased in Epi weeks 4 and remain low in Epi week 7. Levels remain low in Epi week 11. Levels increase to moderate in Epi week 14. Levels decrease to low in Epi week 15 and remain low in Epi week 16. Levels increase to moderate in Epi week 18 and decrease to low in Epi week 19.
- As of Epi week 43, there was a slight increase in SARS-CoV-2 levels in Vlakplaats WWTW, after a decrease from Epi week 35. Levels remain low in Epi week 6. Levels rise to moderate in Epi week 11. Levels decrease to low in Epi week 15 and increased to moderate in Epi week 16. No new results available for Epi week 19.
- * Sequencing data ending in Epi week 5 in Vlakplaats and 6 in Hartebeesfontein.
 - Omicron lineages BA.2.86.X and JN.1.X were circulating in Vlakplaats during Epi week 13.
 - Lineages BA.2.86 and JN.1.X were predominantly circulating during Epi week 10 to 14 at the Hartebeesfontein water treatment plant.

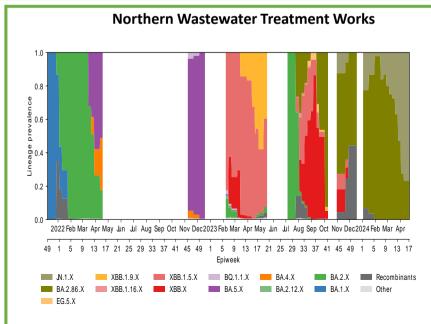
SNP Analysis:

 A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 and JN.1were found in both Ekurhuleni treatment plants.

KwaZulu-Natal - eThekwini







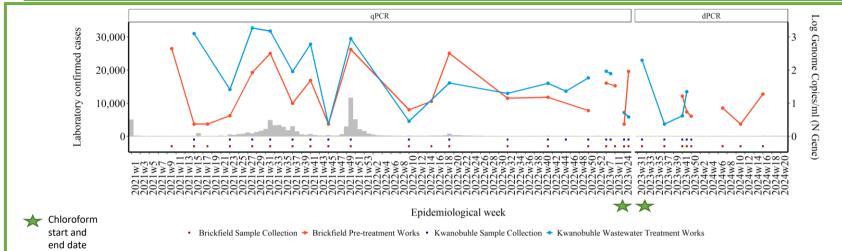
SARS-CoV-2 levels and Genomic Results in Epi week 19:

- SARS-CoV-2 levels in Central WWTW increase in Epi week 10 but remain low. Levels increase to moderate in Epi week 13. Levels decreased to low in Epi week 15 and increased to moderate in Epi week 16 and remain moderate in Epi week 18. No new results for Epi week 19 are available.
- SARS-CoV-2 levels increased in Epi week 5 in Northern WWTW. Levels increase to moderate in Epi week 10 and remain moderate in Epi week 13. Levels increase to high in Epi week 14. Levels decrease to low in Epi week 15 and increase to high in Epi week 16. Levels decrease to moderate in Epi week 17 and drop to low in Epi week 19.
- * Sequencing data ending in Epi week 9 in Central eThekwini and in eThekwini North.
 - During Epiweek 1 17 Lineage BA.2.86.X, JN.1.X were dominantly circulating in eThekwini Central. XBB.X, XBB.1.9.X and Recombinants were also circulating.
 - In eThekwini North, BA.2.86.X lineages were dominating. In circulation there were also lineages JN.1.X and Recombinant lineages during Epi week 1 - 17.

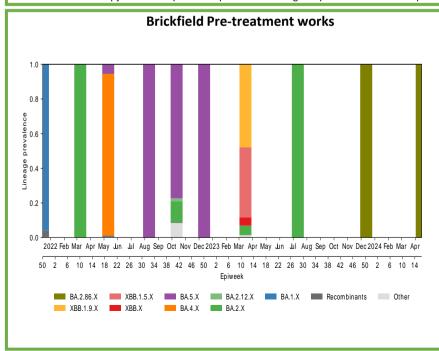
SNP Analysis:

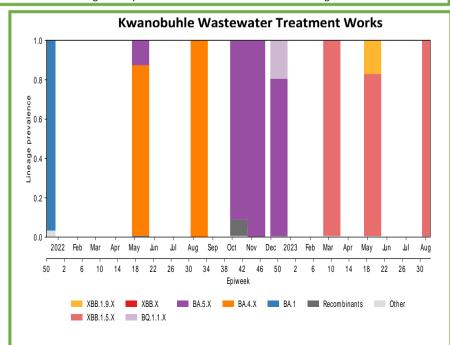
 A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 and JN.1were found in both eThekwini wastewater treatment plants.

Eastern Cape – Nelson Mandela



Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Nelson Mandela Bay (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from gPCR to dPCR occurred for SARS-CoV-2 testing.





SARS-CoV-2 levels and Genomic Results in Epi week 19:

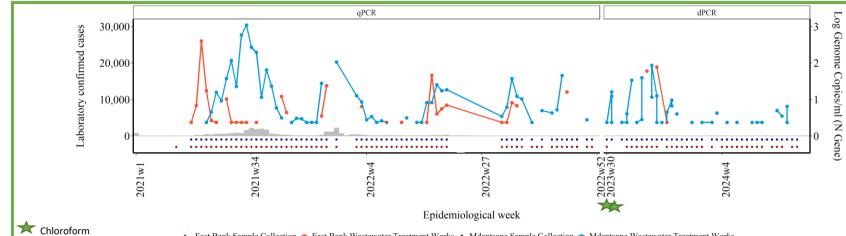
- As of Epi week 46, SARS-CoV-2 levels are moderate after an increase was observed after Epi week 36 in the Kwanobuhle WWTW. No new results are available for Epi week 19.
- SARS-CoV-2 levels decreased from moderate to low from Epi week 43 to Epi week 46 in Brickfield Pre-treatment works. There was a slight increase in level in Epi week 7, but levels remain low. Levels decrease and remain low in Epi week 11. Levels increase to moderate in Epi week 16. No new results for Epi week 19 available.
- * Sequencing data ending in Epi week 14 in Brickfield and 30 in Kwanobuhle.
 - During Epi week 2 to 10, no lineages were detected. BA.2.86.X lineage was dominating during Epi week 14 in Brickfield Pre-treatment Works.
 - Omicron lineages XBB.1.5.X sub-lineages were circulating in Kwanobuhle during Epi week 30.No new sequence data available.

SNP Analysis:

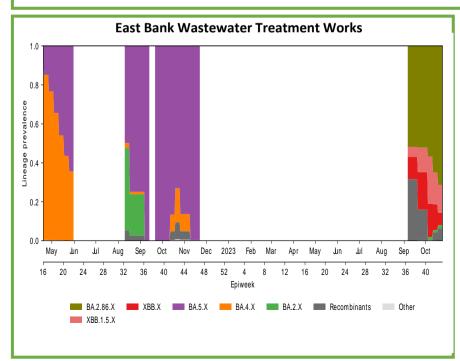
- SNP analysis could not be performed as the SARS-CoV-2 sequencing coverage in the Brickfield samples collected during Epi weeks 30-39 were too low for meaningful interpretation.
- A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 were found in the Kwanobuhle wastewater treatment plants.



Eastern Cape – Buffalo City

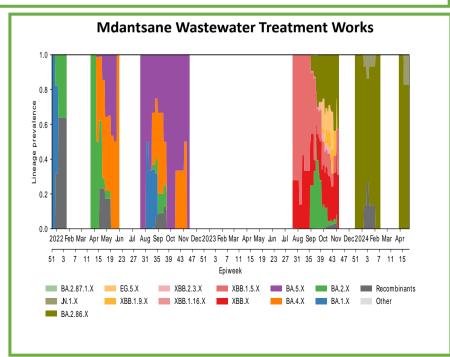


• East Bank Sample Collection • East Bank Wastewater Treatment Works • Mdantsane Sample Collection • Mdantsane Wastewater Treatment Works Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Buffalo City (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from qPCR to dPCR occurred for SARS-CoV-2 testing.



start and end

date



SARS-CoV-2 levels and Genomic Results in Epi week 19:

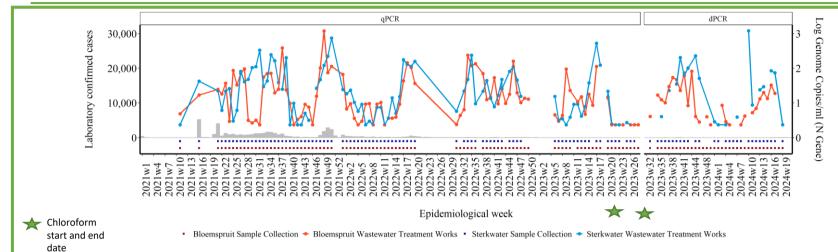
- In Epi week 46, SARS-CoV-2 levels in Mdantsane WWTW decreased and levels are low. Levels in Epi week 12 remain low. Levels increase slightly in Epi week 15 but remain low. Levels remain low in Epi week 17. No new results for Epi week 19 are available.
- SARS-CoV-2 levels in East Bank WWTW in Epi week 44 remain low after decrease was observed from Epi week 38. No new results for Epi week 19 are available.
- * Sequencing data ending in Epi week 40 in Eastbank and Epiweek 15 in Mdantsane.
 - Omicron lineages BA.2.86.X, XBB.1.5.X, XBB.X and Recombinantswere circulating in Eastbank during Epi week 40.
 - Lineage BA.2.86.X was dominating in Mdantsane during Epi week 15. JN.1.X sub-lineages were also detected in Epi week 15.

SNP Analysis:

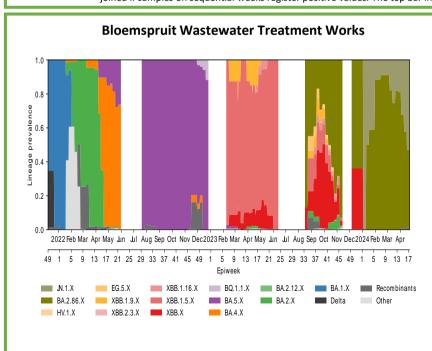
A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 were found in both Eastbank and Mdantsane.

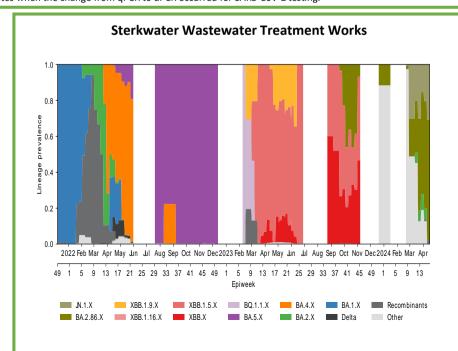


Free State – Mangaung



Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Manguang (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from gPCR to dPCR occurred for SARS-CoV-2 testing.





SARS-CoV-2 levels and Genomic Results in Epi week 19:

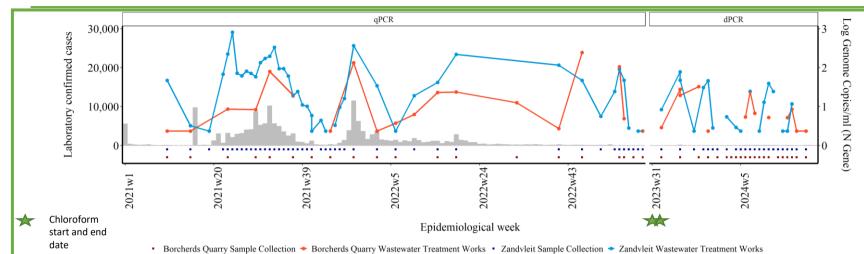
- In Bloemspruit WWTW, a 2-fold increase in SARS-CoV-2 levels were seen in Epi week 44. In Epi week 4 levels decreased and remain low in Epi week 9. Levels increase to moderate in Epi week 13 and remain moderate in Epi week 17. No new results are available for Epi week 19.
- Sterkwater SARS-CoV-2 levels in Epi week 10 increase to high (above 3 genome copies/ml). Levels drop to low in Epi week 11 and increase to moderate in Epi week 13 and remain moderate in Epi week 14. Levels increase further in Epi week 16 and remain moderate in Epi week 17 and decrease to low in Epi week 19.
- * Sequencing data ending in Epi week 17 in Bloemspruit and Epi week 13 in Sterkwater Treatment Works.
 - BA.2.86.X, JN..1.X and XBB.X sub-lineages were the dominant lineage circulating in Bloemspruit during Epi week 1 to 17.
 - BA.2.86.X, JN.1.X and Other Omicron lineages were circulating in Sterkwater in Epi week 9 to 13.

SNP Analysis:

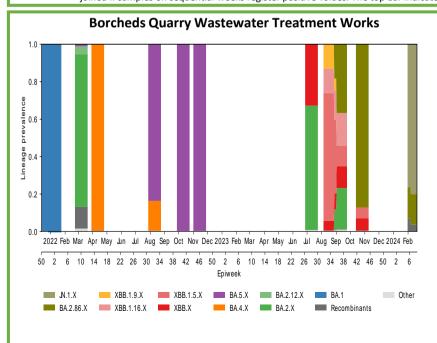
A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 were found in both Bloemspruit and Sterkwater.

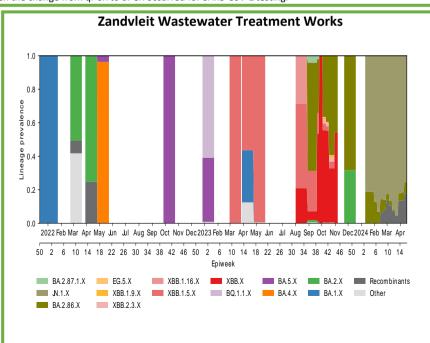


Western Cape – City of Cape Town



Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Cape Town (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from gPCR to dPCR occurred for SARS-CoV-2 testing.





SARS-CoV-2 levels and Genomic Results in Epi week 19:

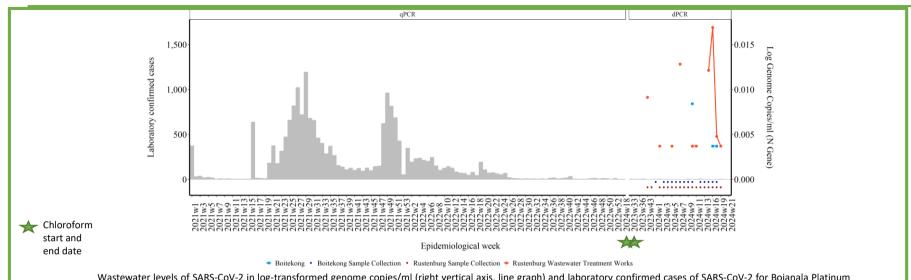
- After a sharp increase in SARS-CoV-2 levels was seen in Epi week 37, a subsequent decrease in SARS-CoV-2 levels in Borcherds Quarry WWTW was seen. Levels are low in Epi week 6 and increase to moderate in Epi week 7. Levels decrease in Epi week 8 to low and remain low in Epi week 19.
- In Epi week 4, SARS-CoV-2 levels were low at Zandvleit WWTW. In Epi week 7 levels increase to moderate. Levels drop to low in Epi week 9 and increase to moderate in Epi week 11. Levels remain moderate in Epi week 12 and decrease to low in Epi week 14 and remain low in Epi week 15. Levels increase to moderate in Epi week 16. No new results for Epi week 19 are available.
- * Sequencing data ending in Epi week 6 in Borcheds Quarry and Epi week 10 in Zandvleit.
 - During Epi week 6, JN.1.X sub-lineages were dominantly circulating in Borcheds. BA.2.86.X and Recombinants were also detected.
 - During Epi week 2 6, lineages BA.2.86.X and JN.1.X were circulating in Zandvlet, with JN.1.X sub-lineages being dominant.
 - In epi week 10 and 14 BA.2.86,X, JN.1.X and Recombinants lineages were circulating in Zandvlet, with JN.1.X lineages at highest proportion

SNP Analysis:

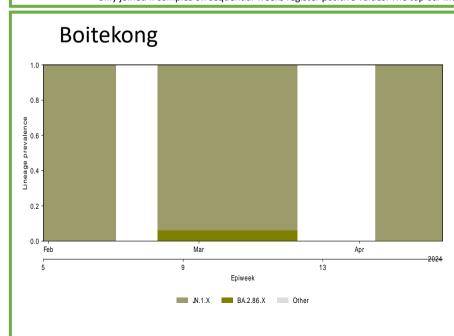
 A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 and JN.1 were found in both Borcheds Quarry and Zandevleit.

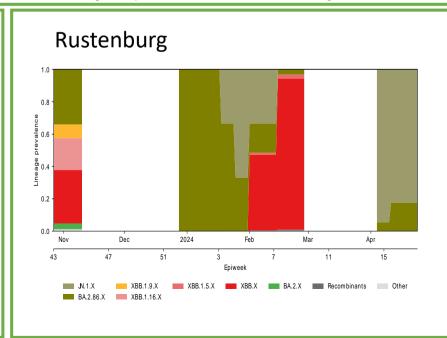


North West – Bojanala Platinum



Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Bojanala Platinum (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from qPCR to dPCR occurred for SARS-CoV-2 testing.





- * SARS-CoV-2 wastewater sample collection and testing at Rustenburg WWTW began in Epi week 43, 2023.
- * SARS-CoV-2 wastewater sample collection and testing at Boitekong began in Epi week 1, 2024.

SARS-CoV-2 levels and Genomic Results in Epi week 19:

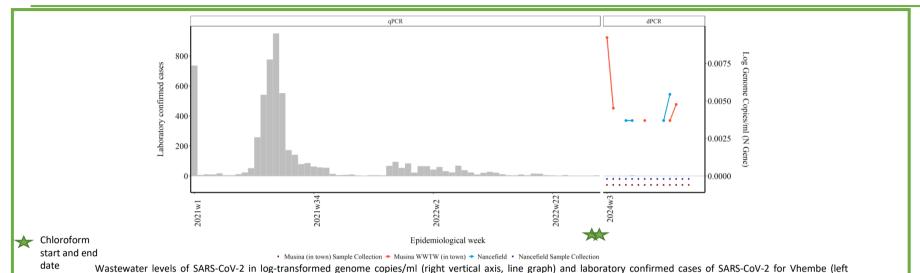
- At Rustenburg WWTW levels from Epi week 43 remain below 0.015 log genome copies/ml and remain low up until Epi week 19.
- At Boitekong wastewater results for SARS-CoV-2 were low in Epi week 10, 2024. Levels remain low in Epi week 16. No new results for Epi week 19 are available.
- * Sequencing data ending in Epi week 9 in Boitekong and Epi week 15 in Rustenburg.

SARS-CoV-2 levels and Genomic Results in Epi week 13:

- At Boitekong, JN.1.X sub-lineages were dominantly circulating in Epi week 15. In Epi week 13, no lineages were detected.
- In Rustenburg, JN.1.X and BA.2.86.X were circulating during Epi week 15. No lineages were detected during Epi week 11.

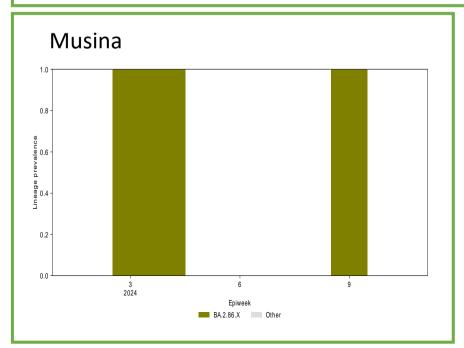


Limpopo – Vhembe



vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only

joined if samples on sequential weeks register positive values. The top bar indicates when the change from qPCR to dPCR occurred for SARS-CoV-2 testing.





- * SARS-CoV-2 wastewater sample collection and testing at Musina WWTW began in Epi week 3, 2024.
- * SARS-CoV-2 wastewater sample collection and testing at Nancefield began in Epi week 3, 2024.

SARS-CoV-2 levels and Genomic Results in Epi week 19:

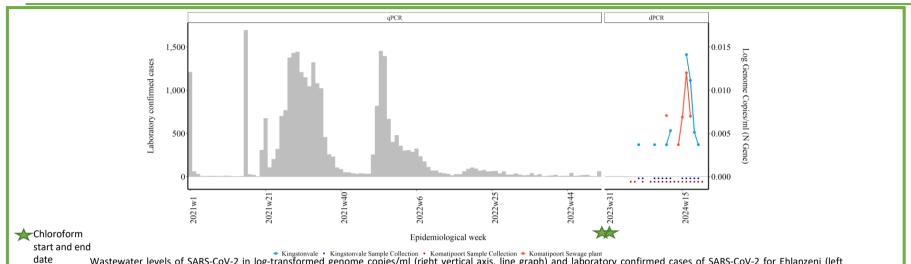
- At Musina WWTW levels from Epi week 3 remain below 0.01 log genome copies/ml and remain low up until Epi week 4, 2024. Levels remain low in Epi week 9. No new results for Epi week 19 are available.
- At Nancefield levels were low in Epi weeks 6 and 7, below 0.005 log genome copies/ml. They remain low in Epi week 12. No new results for Epi week 19 are available.
- * Sequencing data ending in Epiweek 9 in Musina and Epi week 15 in Musina

SARS-CoV-2 levels and Genomic Results in Epi week 13:

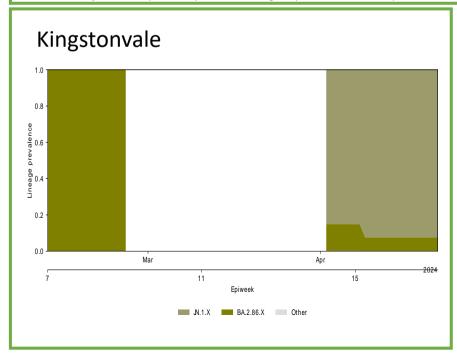
- Sequencing in Musina began in Epiweek 3, and BA.2.86.X sub-lineages were found to be the main circulating lineages. In Epi week 9 BA.2.86.X sub-lineages were still the dominant lineages detected in Musina WWTW
- BA.2.86.X lineages were detected in Nancefield during Epi weeks 7 and 15, in Epi week 11 no lineages were detected.



Mpumalanga – Ehlanzeni



* Kingstonvale * Kingstonvale Sample Collection * Komatipoort Sample Collection * Komatipoort Sewage plant
Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Ehlanzeni (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 wastewater level data points are only joined if samples on sequential weeks register positive values. The top bar indicates when the change from qPCR to dPCR occurred for SARS-CoV-2 testing.



Komatipoort

*No sequencing data currently available

- * SARS-CoV-2 wastewater sample collection and testing at Komatipoort WWTW began in Epi week 2, 2024.
- * SARS-CoV-2 wastewater sample collection and testing at Kingstonvale began in Epi week 4, 2024.

SARS-CoV-2 levels and Genomic Results in Epi week 19:

- At Komatipoort samples collected between Epi week 2 and Epi week 10 results were negative for SARS-CoV-2. Levels were low in Epi week 11 and remain low in Epi week 17. No new results for Epi week 19 are available.
- At Kingstonvale SARS-CoV-2 levels remain below 0.01 log genome copies/ml from Epi week 4 to Epi week 8. Levels remain low in Epi week 19.
- * Sequencing data ending in Epi week 15 in Kingstonvale
- No lineages were detected in Epi week 11. In Epi week 15 BA.2.86.X and JN.1.X lineages were detected, with JN.1.X dominating.



COLLABORATORS TEAM

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