



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

**17 June 2024**

Sample collection dates up to 7 June 2024  
(Epidemiological week 23)

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

## Wastewater levels

### Epidemiological weeks 48 (2023) –23 (2024)

- From weeks 48-13 the cumulative SARS-CoV-2 levels measured at wastewater treatment works (WWTW) **have 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 weeks 1-6 lower levels were observed nationally, with an increase seen in week 7 and a subsequent decrease in week 8. Levels increase 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 decreased further in Epi week 19. Levels increase to high (above 2 log (100) genome copies/ml) in Epi week 21 and decrease to moderate in Epi week 23.**
- 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 genomics

### Epidemiological weeks 48 (2023) – 23 (2024)

- **Omicron lineage BA.2.86.X (this includes KP.1 and BA.2.86.1), and JN.1.X sub-lineages** were the dominant lineages circulating in wastewater samples between January 2024 – May 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 eThekweni (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

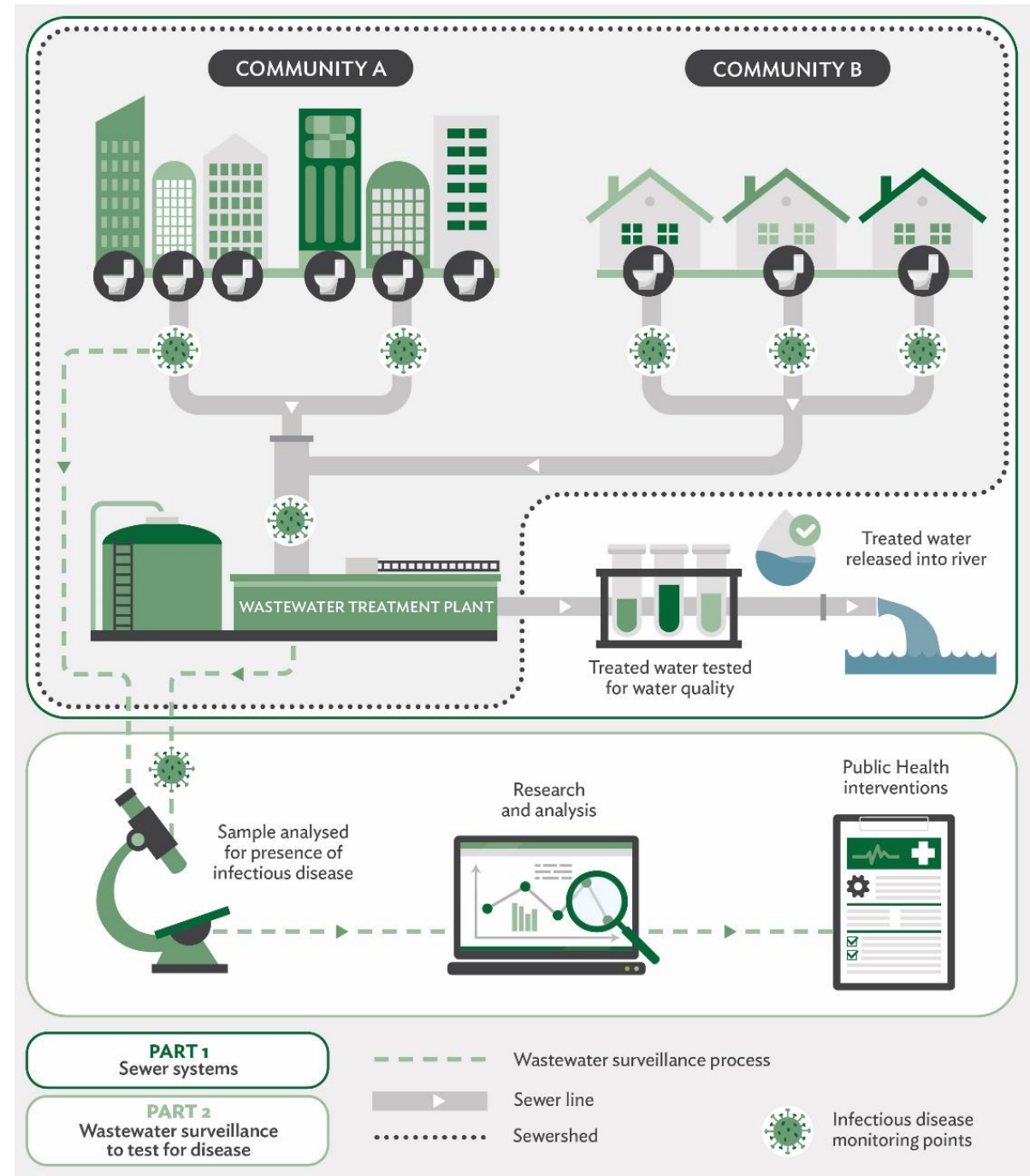
## How is wastewater tested for SARS-CoV-2?

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-essays/detail/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?

Left vertical axis:

Number of lab confirmed cases in *the metro or district where the water treatment plant is located*

Bars:

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

Coloured lines:

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

Coloured squares:

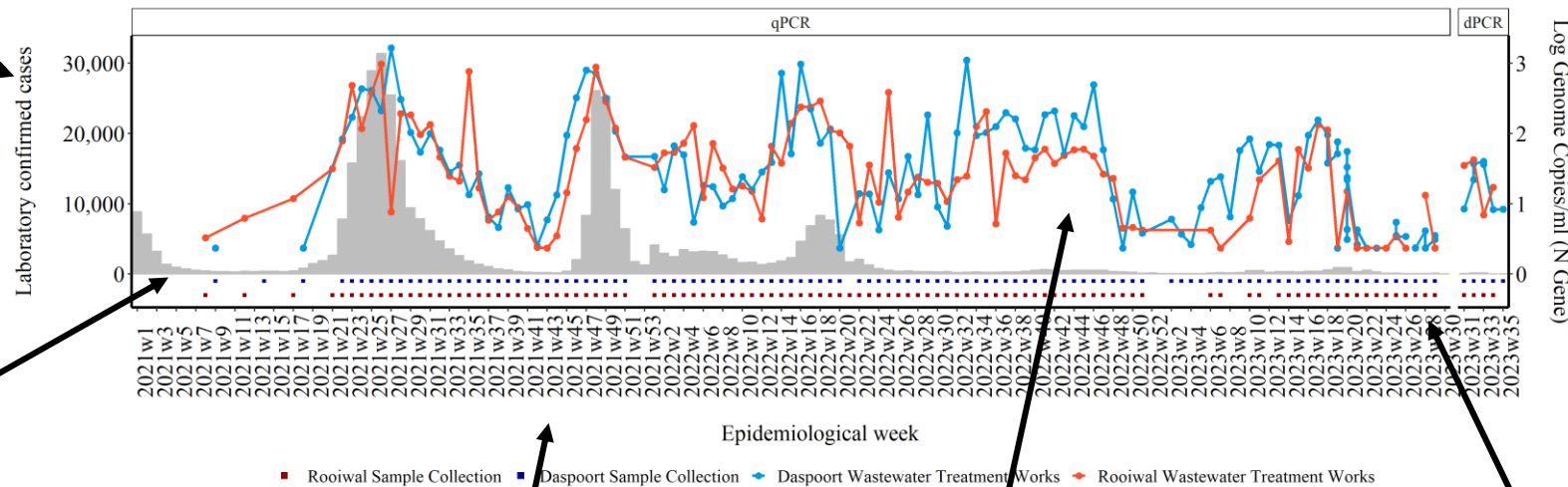
Epi weeks during which samples were collected

Facets:

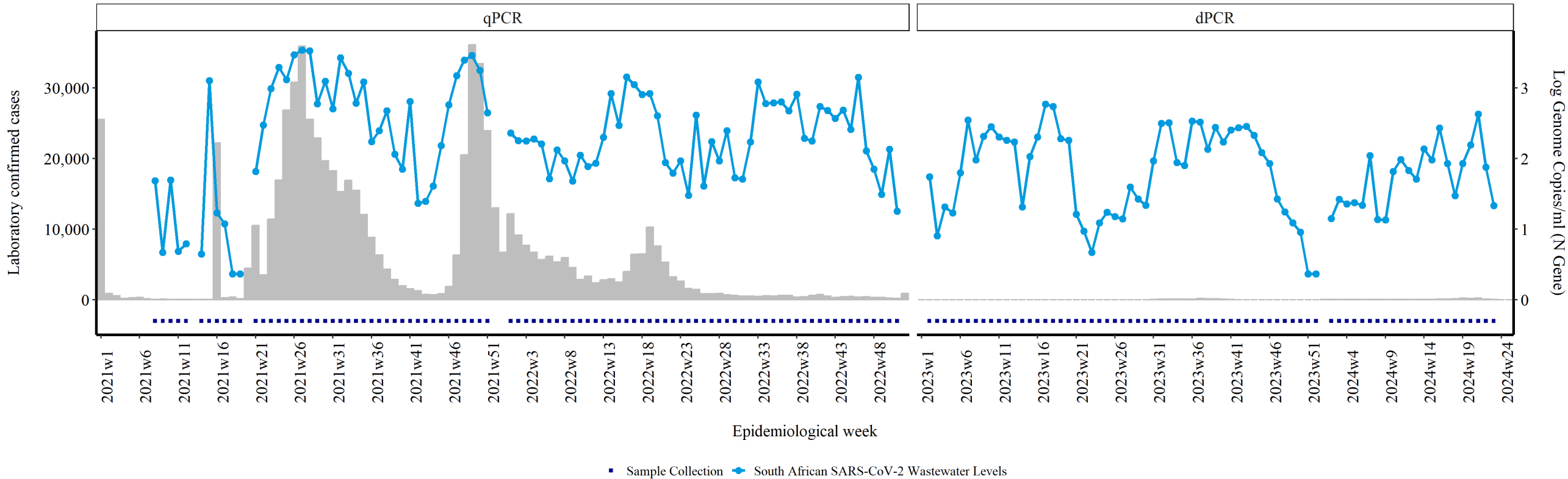
Indicates the platform used to test samples

Right vertical axis:

Log (ie 10 to the power x) copies of SARS-CoV-2 genome per ml of wastewater. So  $\log 2=10^2=100$  copies per millilitre,  $\log 3=10^3=1000$  copies per millilitre



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

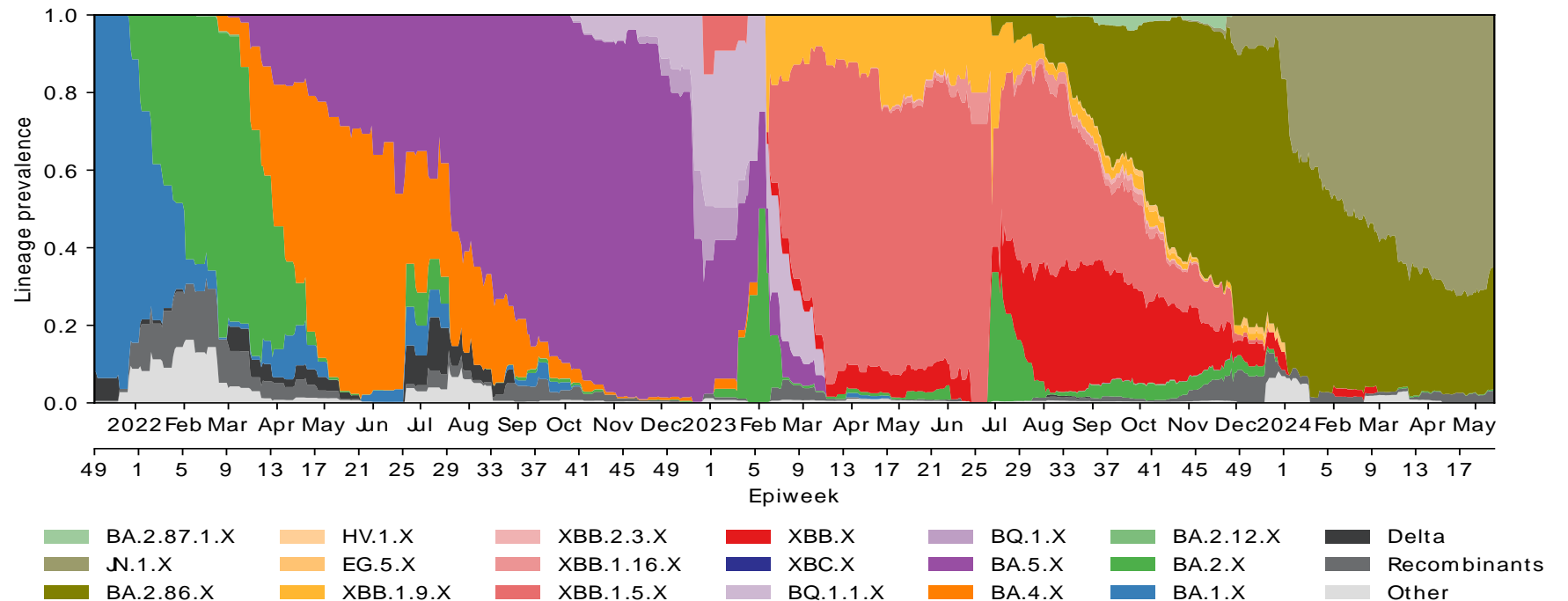


plants tested by NICD, compared with laboratory-confirmed cases from [Tshwane, Johannesburg, Ekurhuleni, eThekweni, Mangaung, Nelson Mandela, Buffalo City, City of Cape Town, Vhembe, Ehlanzeni and Bonjanala Platinum](#) (left vertical axis, grey bars), by epidemiological week, 2021-2023. Coloured dots 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

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

- 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 19 (May 2024)
  - Omicron lineages BA.2.86.X and JN.1.X were circulating from December to May, with Recombinants and XBB.X sub-lineages circulating at low proportion.

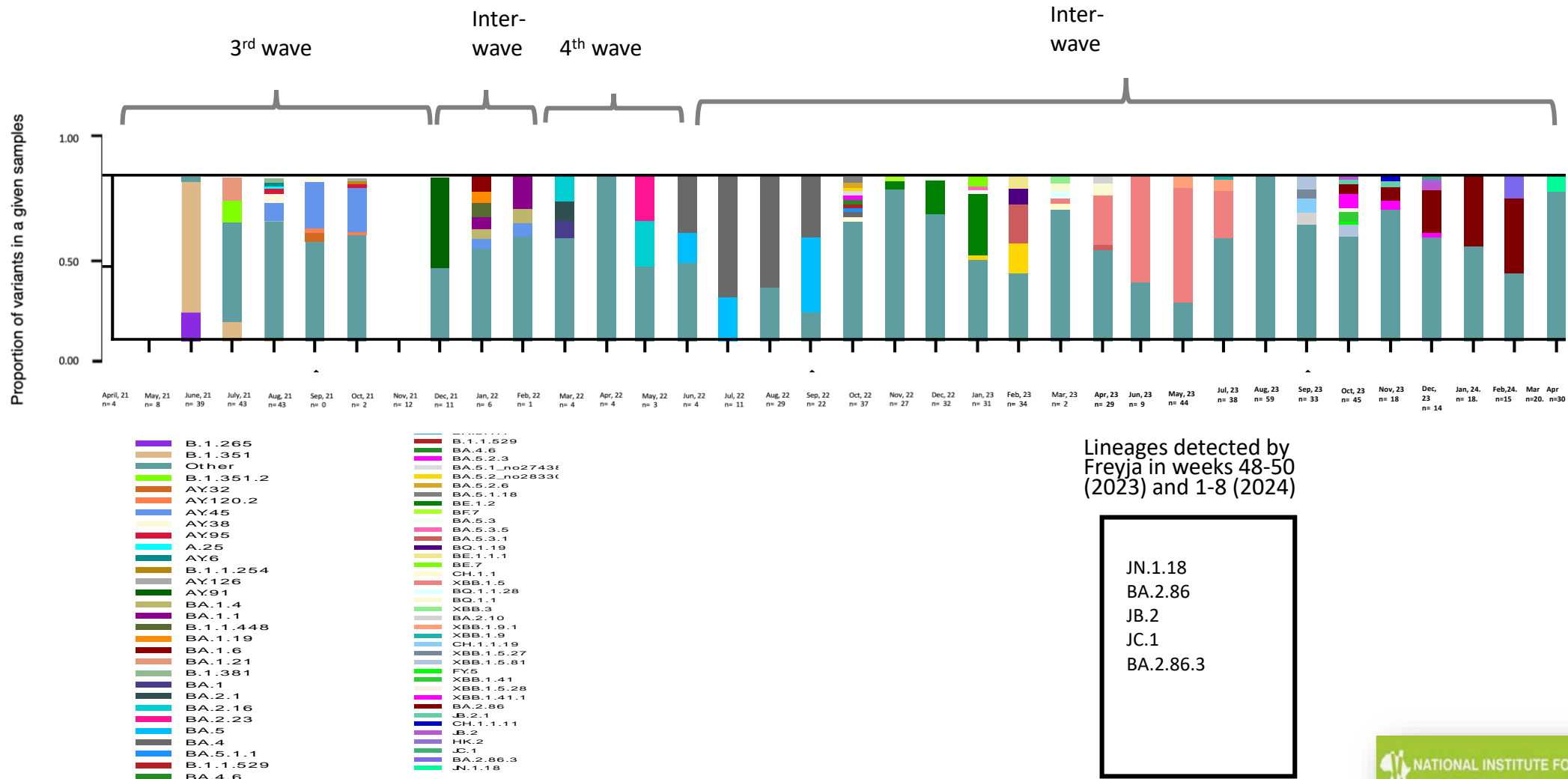


April, 21 n=4   May, 21 n=8   June, 21 n=39   July, 21 n=43   Aug, 21 n=43   Sep, 21 n=0   Oct, 21 n=2   Nov, 21 n=12   Dec, 21 n=11   Jan, 22 n=6   Feb, 22 n=1   Mar, 22 n=4   Apr, 22 n=4   May, 22 n=3   Jun, 22 n=4   Jul, 22 n=11   Aug, 22 n=29   Sep, 22 n=22   Oct, 22 n=37   Nov, 22 n=27   Dec, 22 n=32   Jan, 23 n=31   Apr, 23 n=29   May, 23 n=44   Jun, 23 n=9   Jul, 23 n=38   Aug, 23 n=59   Sep, 23 n=33   Oct, 23 n=45   Nov, 23 n=18   Dec, 23 n=14   Jan, 24 n=18   Feb, 24 n=15   Mar, 24 n=20   Apr, 24 n=30   May n=10

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

- Results from sequencing data ending in epi week 19 (May 2024)

- Omicron lineages BA.2.86 and JN.1 were circulating from January to March with BA.2.86 dominance.
- Omicron XBB sub-lineages 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

## Summary:

Summary of lineages associated with most common mutations associated

## Epi Week:

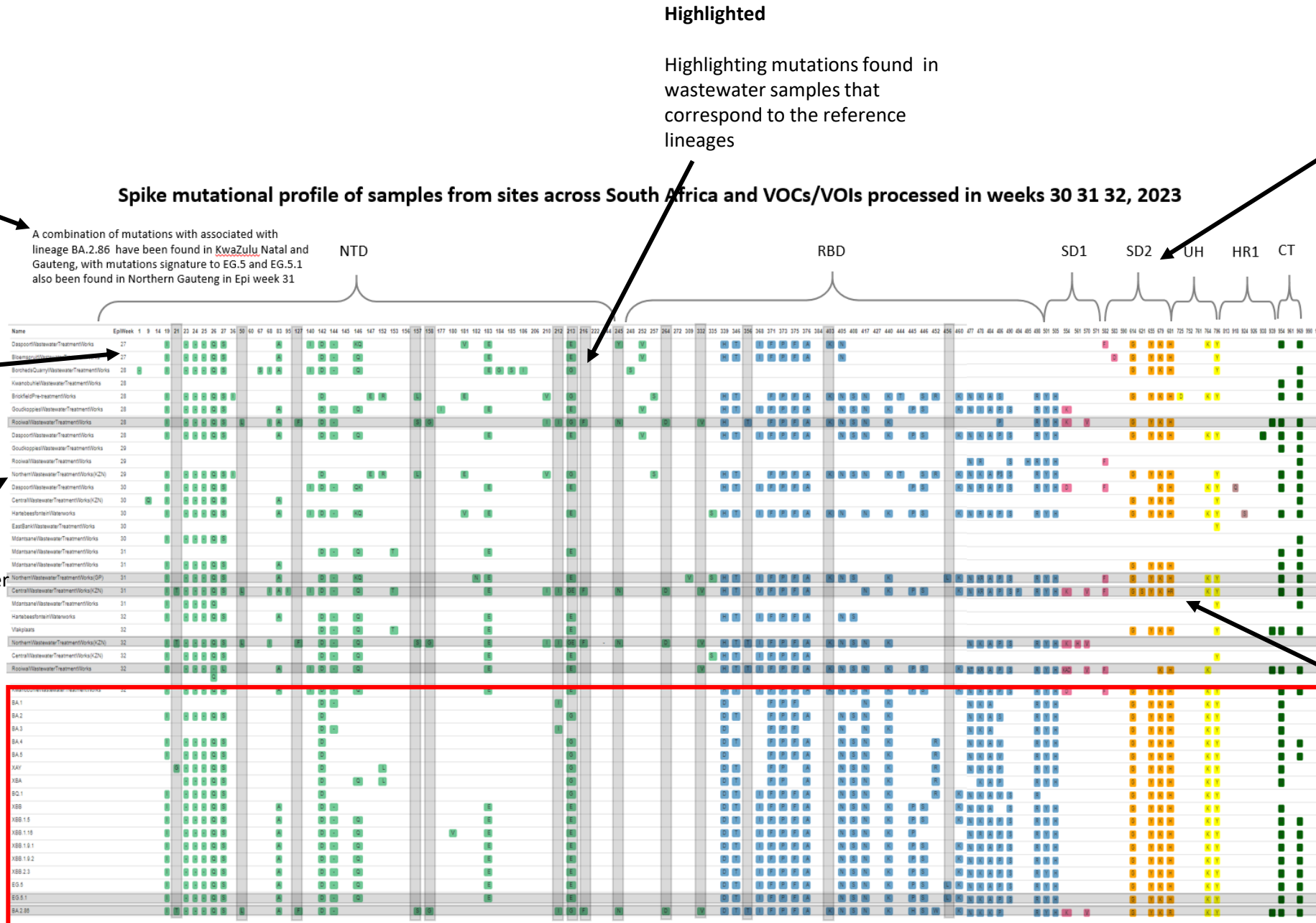
Epidemiological week during which samples were collected

## Site Names:

Sites from which wastewater samples were collected

## Reference lineages:

Reference lineages with signature mutations with which wastewater samples are compared for lineage determination



Top x-axis:  
Position of the spike protein

Coloured blocks  
Indicate changes (mutations) in the amino acids of the spike protein









# 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 concern  
<https://www.who.int/en/activities/tracking-SARS-CoV-2-variants>

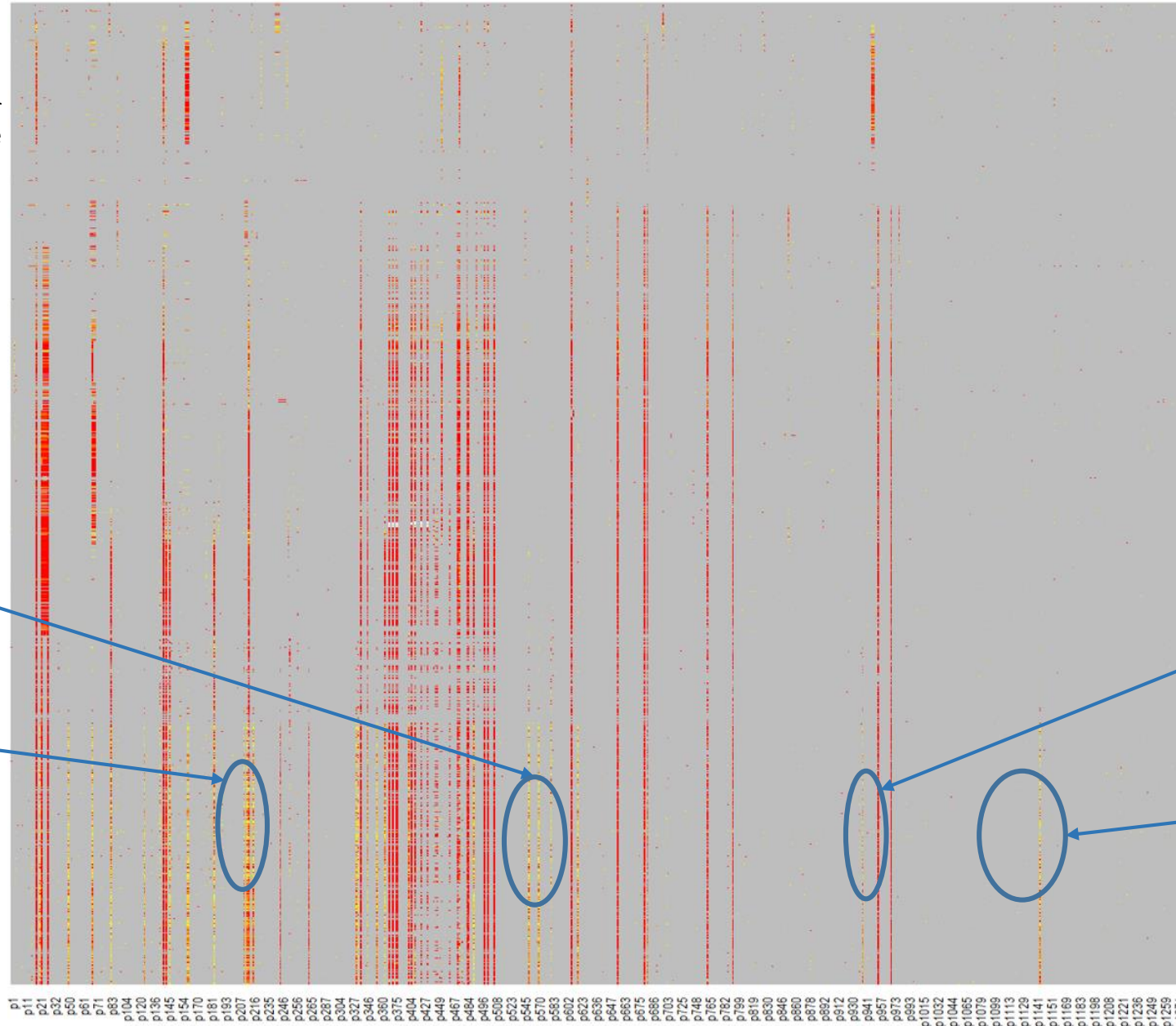
**BA.2.86** is a highly mutated sub-lineage of BA.2, recently circulating in Denmark, Israel and the United States of America and is characterised by one or more of the following mutations in the spike protein: R21T, S50L, H69-, V70-, V127F, F157S, R158G, N211-, L212I, V213G, L216F, H245N, A264D, I332V, K356T, R403K, V445H, N450D, L452W, N481K, V483-, E484K, E554K, A570V, P621S, 1670V, P681R, S939F, P1143L, Ins16:MPLF\*

E554K  
Mutations in spike protein associated with BA.2.86

V213E,  
R346T  
Mutations in spike protein associated with XBB\* sub-lineages

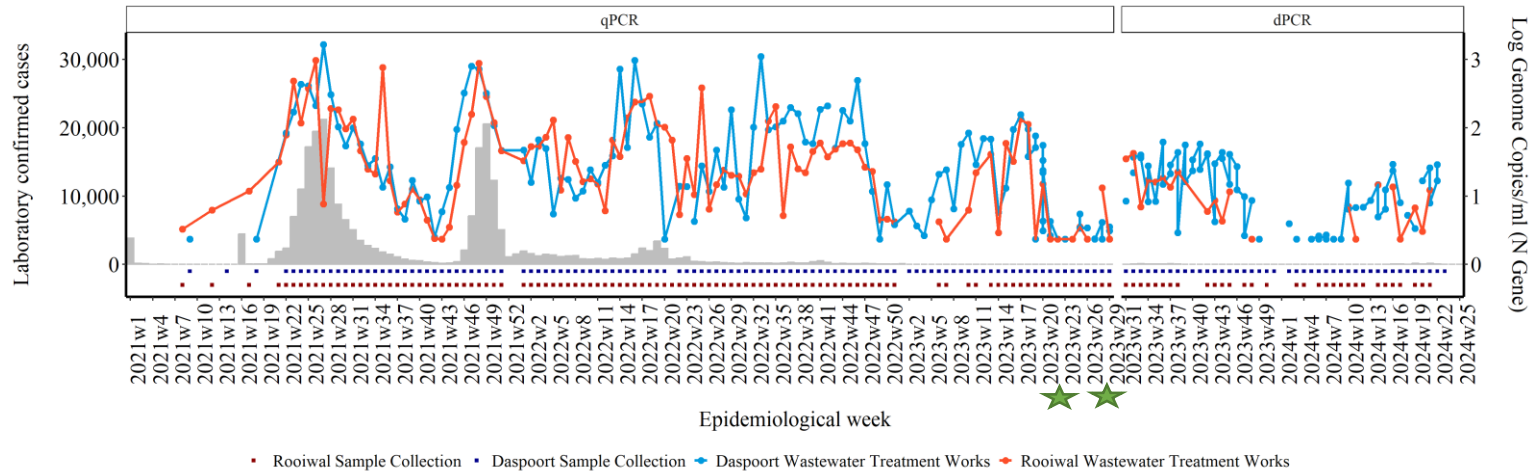
S939F  
Spike protein mutation associated with the BA.2.86 lineage

P1143L  
Spike protein mutation associated with the BA.2.86 lineage



Heatmap showing patterns of emerging mutations in the spike region of SARS-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.

# Gauteng - Tshwane



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 23:

- 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. Levels rise to moderate in Epi week 22 and remain moderate in Epi week 23.
- 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 rise to moderate in Epi week 21. No new results are available for Epi week 23.

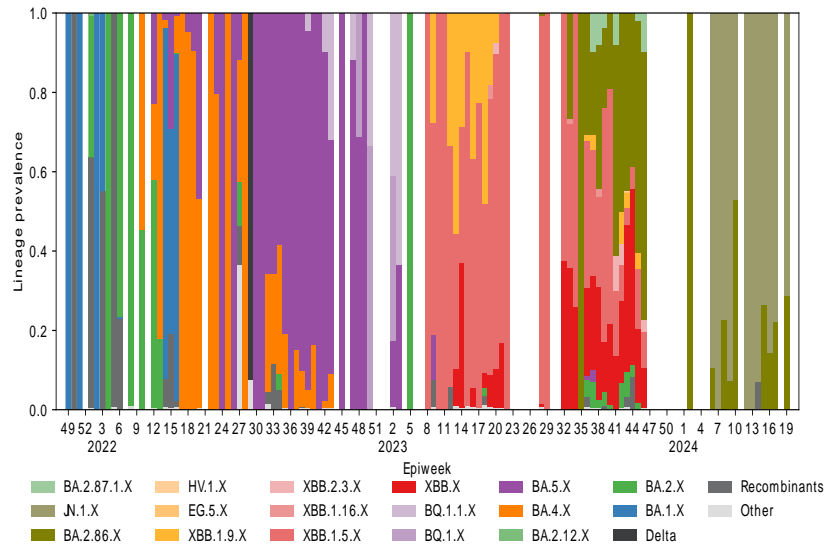
**\* Sequencing data ending in Epi week 19 in Daspoort and in Rooiwal.**

- 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

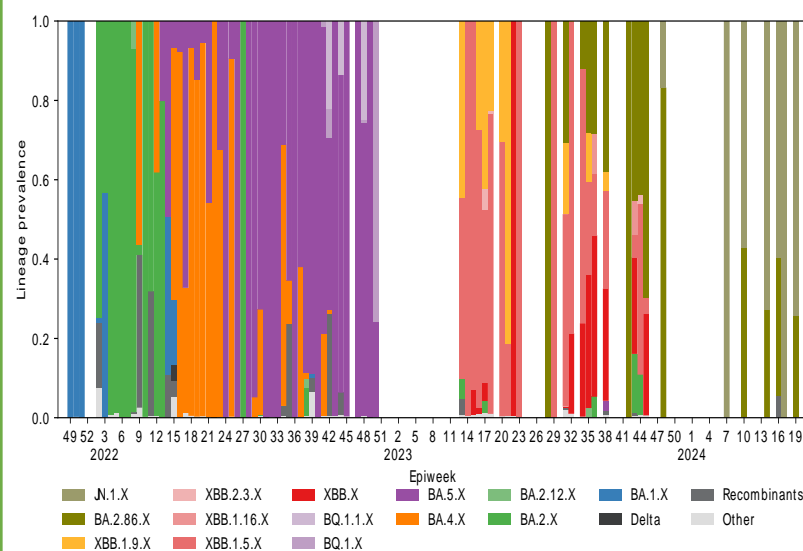
## SNP Analysis:

- 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.

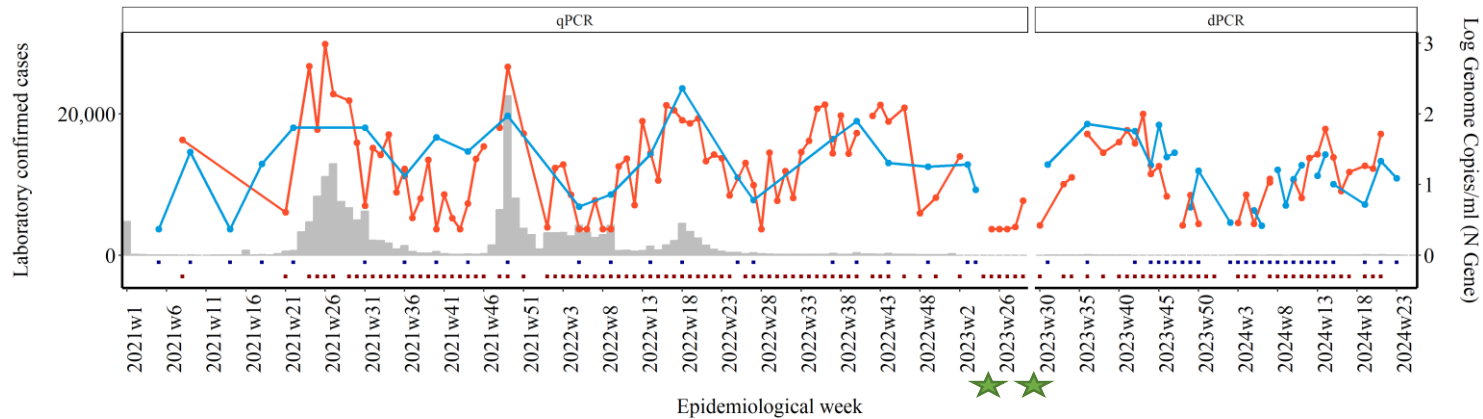
### Daspoort Wastewater Treatment Works



### Rooiwal Wastewater Treatment Works



# Gauteng - Johannesburg



Chloroform start and end date

• 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 23:

- 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 remain low in Epi week 19 and increase to moderate in Epi week 21. No new results for Epi week 23 are available.
- 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 decrease to low in Epi week 19 and increase to moderate in Epi week 21. Level remain moderate in Epi week 23.

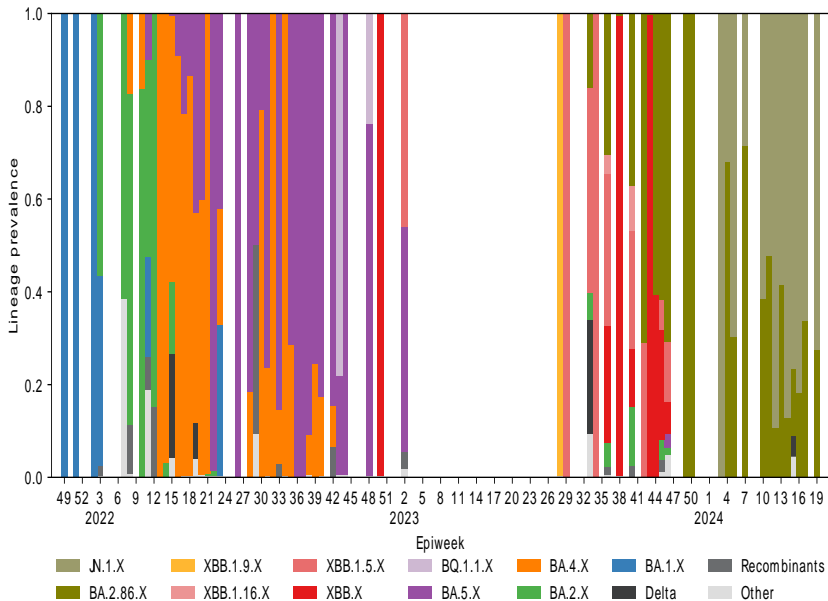
**\* Sequencing data ending in Epi week 19 in Goudkoppies and in Northern Wastewater Treatment Works.**

- 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.

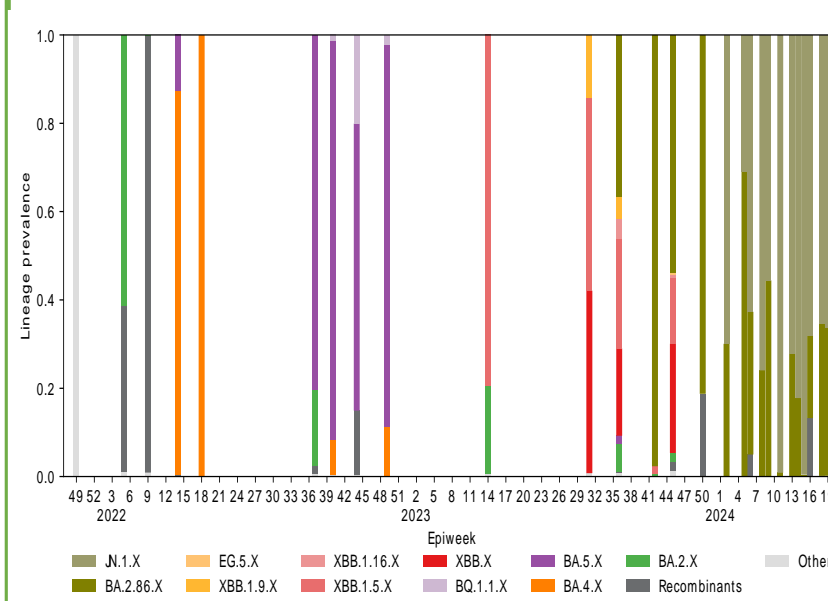
## SNP Analysis:

- 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.1 were found in Northern Johannesburg.

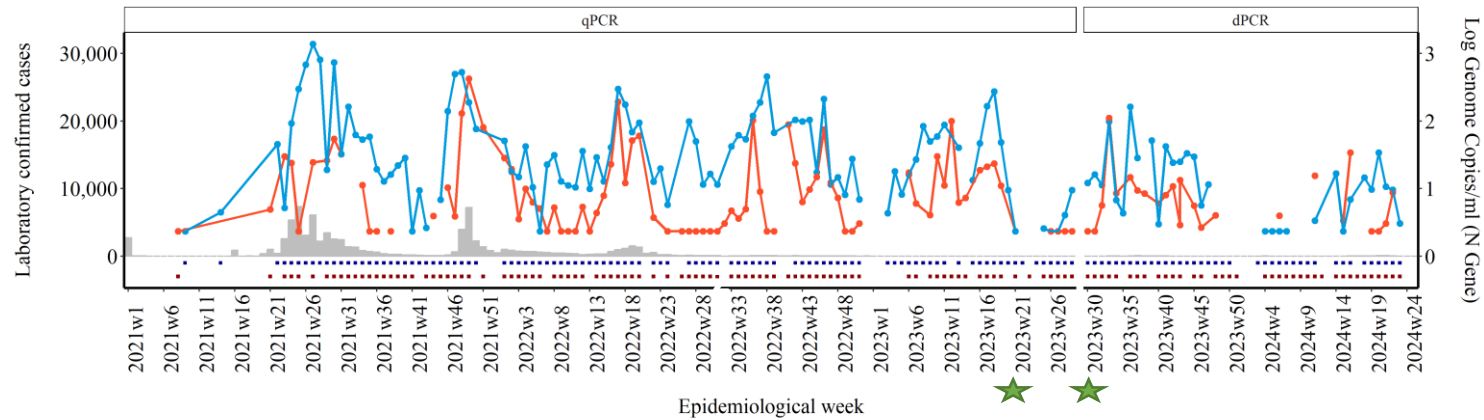
## Goudkoppies Wastewater Treatment Works



## Northern Wastewater Treatment Works



# Gauteng - Ekurhuleni



★ Chloroform start and end date

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

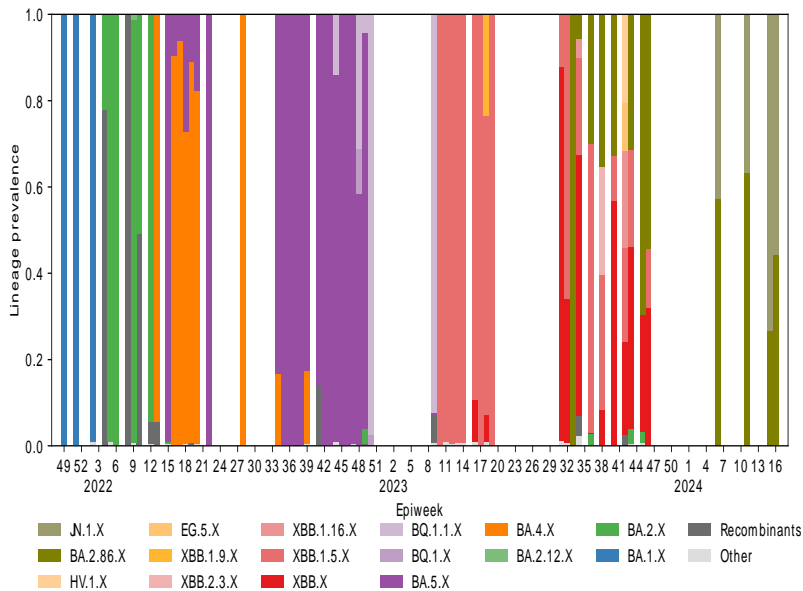
Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Ekurhuleni (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2 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 23:

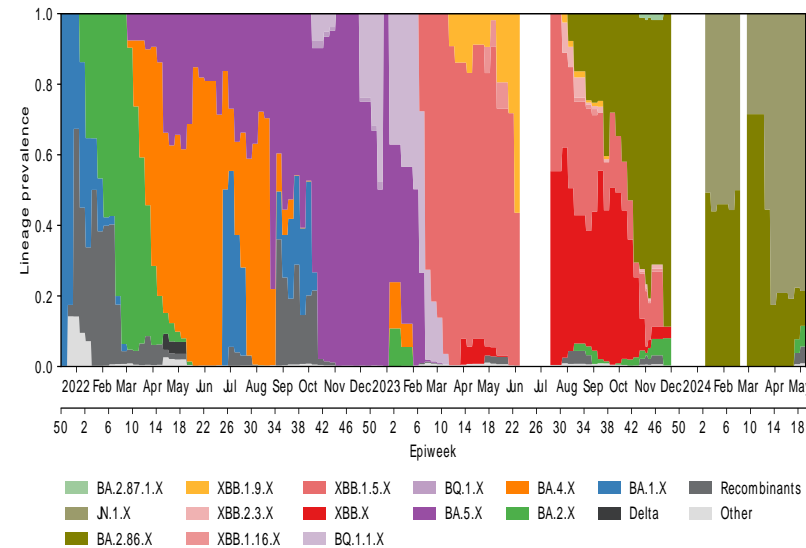
- The SARS-CoV-2 levels in Hartebeesfontein WWTW decreased in Epi weeks 4 and remain low in Epi week 7. Levels remain low in Epi week 11. Levels increase to moderate in Epi week 14. Levels decrease to low in Epi week 15. Levels increase to moderate in Epi week 20 and decrease to low in Epi week 23.
- 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 20 and decrease to low in Epi week 22. No new results for Epi week 23 are available.

**\* Sequencing data ending in Epi week 16 in Vlakplaats and 18 in Hartebeesfontein.**

## ERWAT Vlakplaats Wastewater Treatment Works



## Hartebeesfontein Wastewater Treatment Works

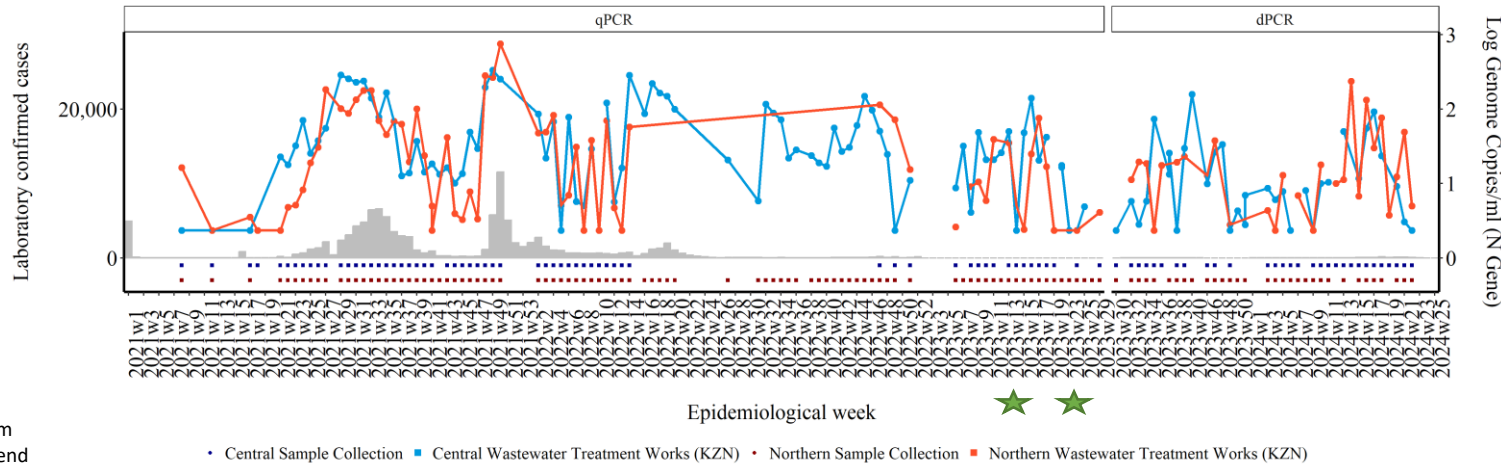


- Omicron lineages BA.2.86.X and JN.1.X were circulating in Vlakplaats during Epi week 13.
- Lineages BA.2.86 and JN.1.X were predominantly circulating during Epi week 10 to 14 at the Hartebeesfontein water treatment plant.

## SNP Analysis:

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

# KwaZulu-Natal - eThekweni



★ Chloroform start and end date

• Central Sample Collection • Central Wastewater Treatment Works (KZN) • Northern Sample Collection • Northern Wastewater Treatment Works (KZN)

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 eThekweni (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 23:

- 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. Levels decreased to low in Epi week 22. No new results for Epi week 23 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 moderate in Epi week 17 and drop to low in Epi week 19. Levels increase again to moderate in Epi week 21 and decrease to low in Epi week 22. No new results for Epi week 23 are available.

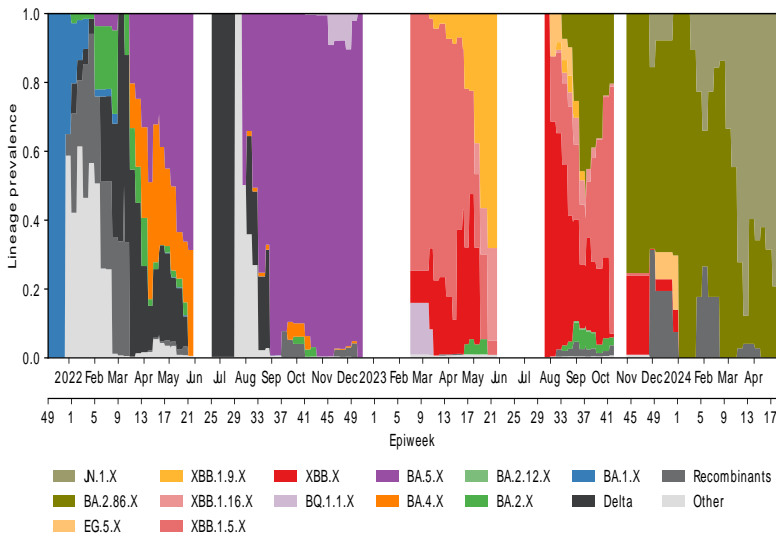
**\* Sequencing data ending in Epi week 17 in Central eThekweni and in eThekweni North.**

- During Epiweek 1 - 17 Lineage BA.2.86.X, JN.1.X were dominantly circulating in eThekweni Central. XBB.X, XBB.1.9.X and Recombinants were also circulating.
- In eThekweni 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.

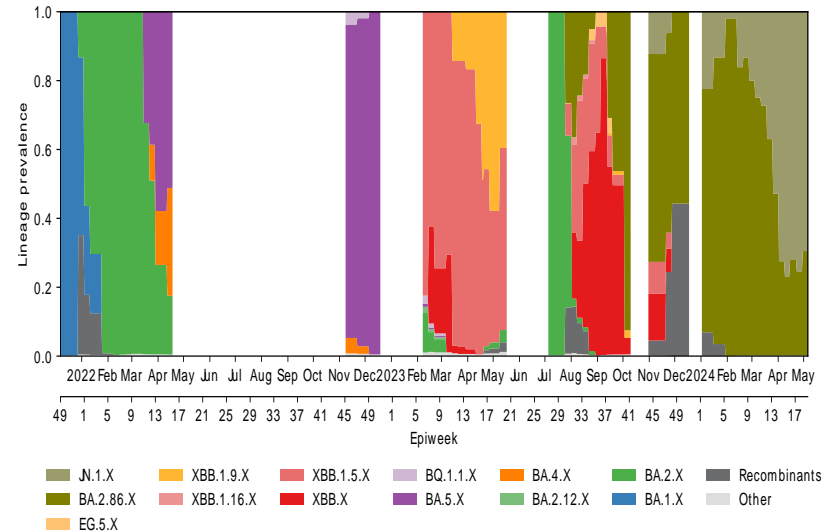
## SNP Analysis:

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

## Central Wastewater Treatment Works

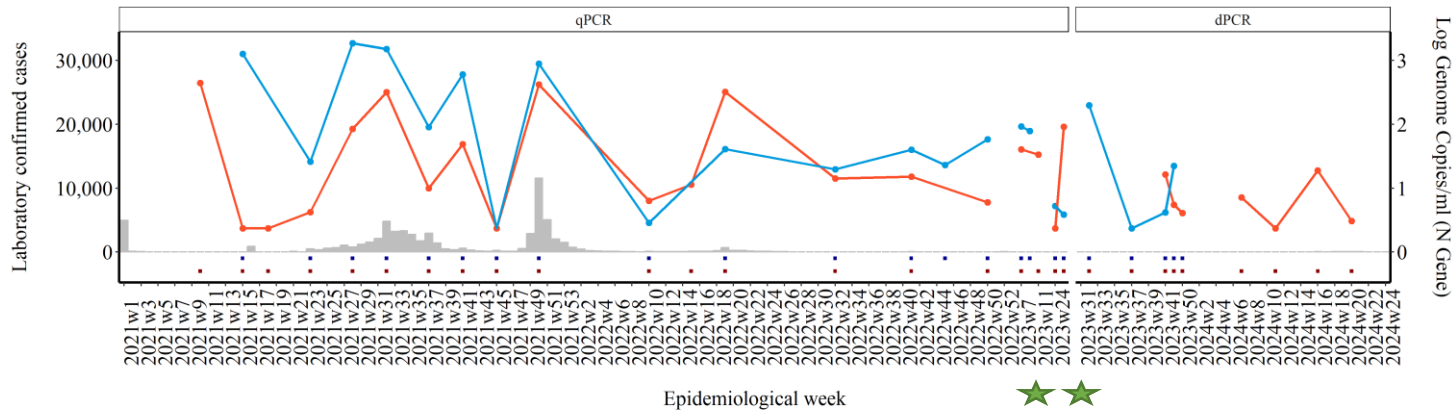


## Northern Wastewater Treatment Works





# Eastern Cape – Nelson Mandela



★ Chloroform start and end date

• Brickfield Sample Collection • Brickfield Pre-treatment Works • Kwanobuhle Sample Collection • Kwanobuhle 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 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 qPCR to dPCR occurred for SARS-CoV-2 testing.

## SARS-CoV-2 levels and Genomic Results in Epi week 23:

- 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 23.
- 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 and decrease to low in Epi week 19. No new results are available for Epi week 23.

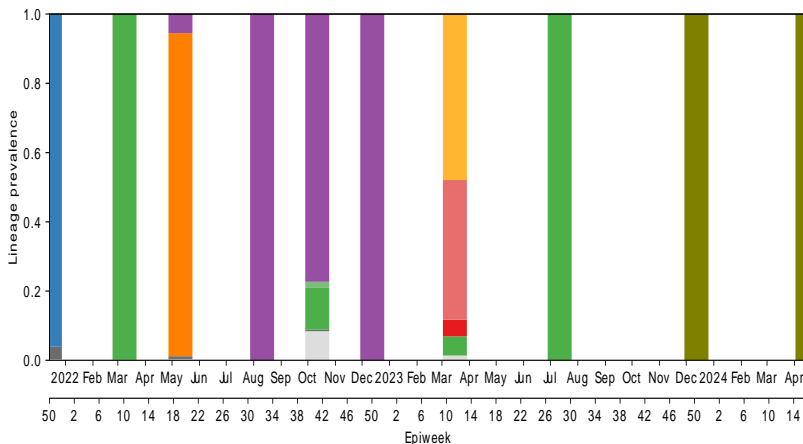
**\* Sequencing data ending in Epi week 14 in Brickfield and 30 in Kwanobuhle.**

- 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:

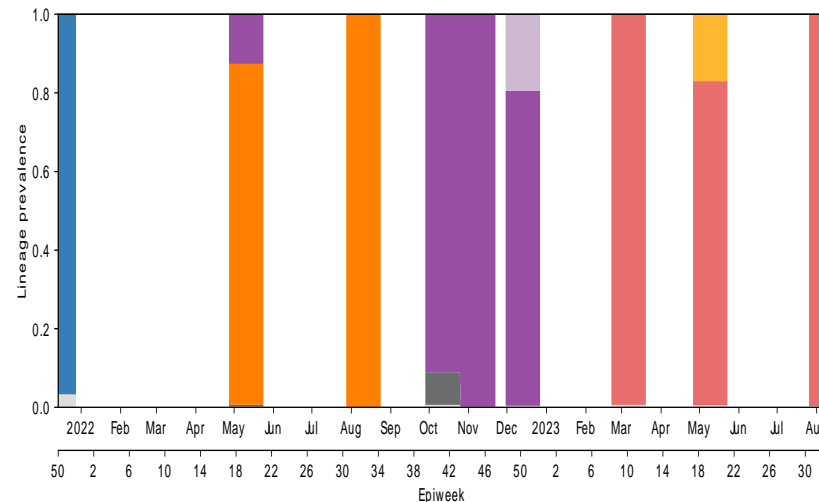
- SNP analysis could not be performed as the SARS-CoV-2 sequencing coverage in the Brickfield samples collected during Epi weeks 30-39 were too low for meaningful interpretation.
- A combination of mutations (V127F, L212I, V213G, L216F, H245N, A264D, I332V, K356T) associated with lineage BA.2.86 were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

## Brickfield Pre-treatment works



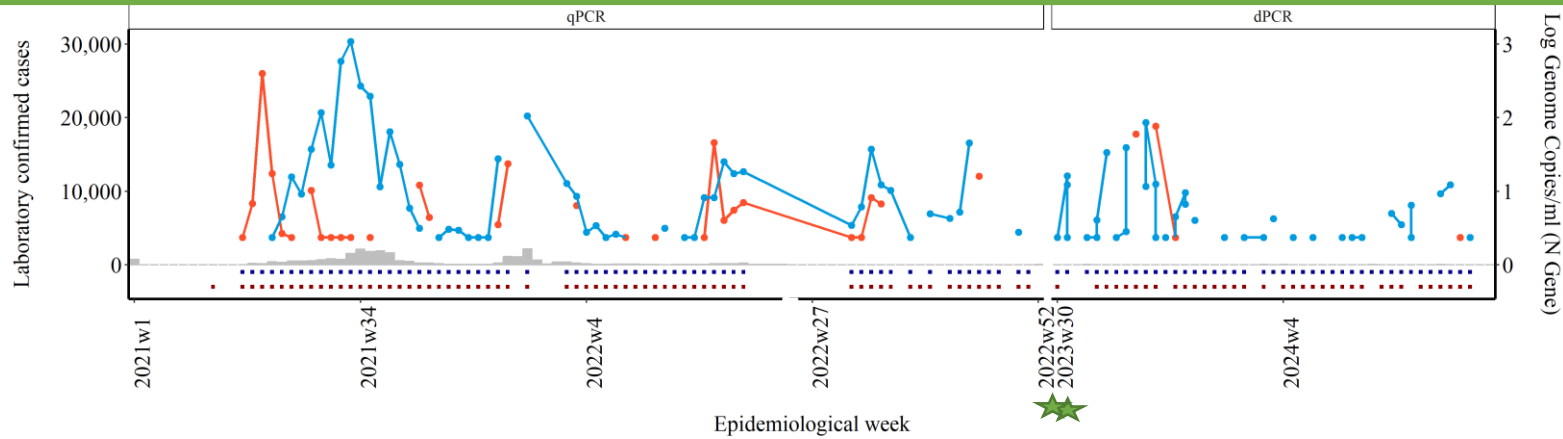
■ BA.2.86.X ■ XBB.1.5.X ■ BA.5.X ■ BA.2.12.X ■ BA.1.X ■ Recombinants ■ Other  
 ■ XBB.1.9.X ■ XBB.X ■ BA.4.X ■ BA.2.X

## Kwanobuhle Wastewater Treatment Works



■ XBB.1.9.X ■ XBB.X ■ BA.5.X ■ BA.4.X ■ BA.1 ■ Recombinants ■ Other  
 ■ XBB.1.5.X ■ BQ.1.1.X

# Eastern Cape – Buffalo City



## SARS-CoV-2 levels and Genomic Results in Epi week 23:

- 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 increase to moderate in Epi week 21 and decrease to low in Epi week 23.
- SARS-CoV-2 levels in East Bank WWTW in Epi week 44 remain low after decrease was observed from Epi week 38. Levels remain low in Epi week 22. No new results for Epi week 23 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 Recombinants were 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.

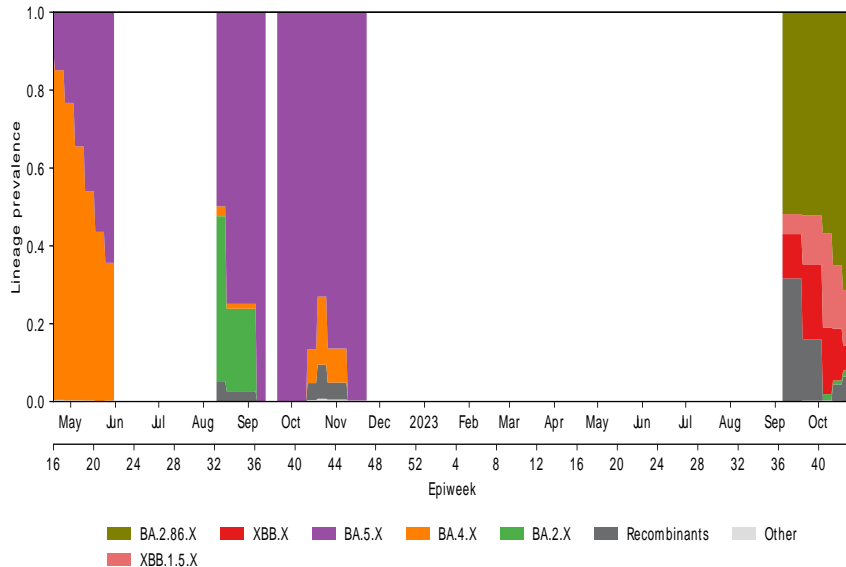
## SNP Analysis:

- 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.

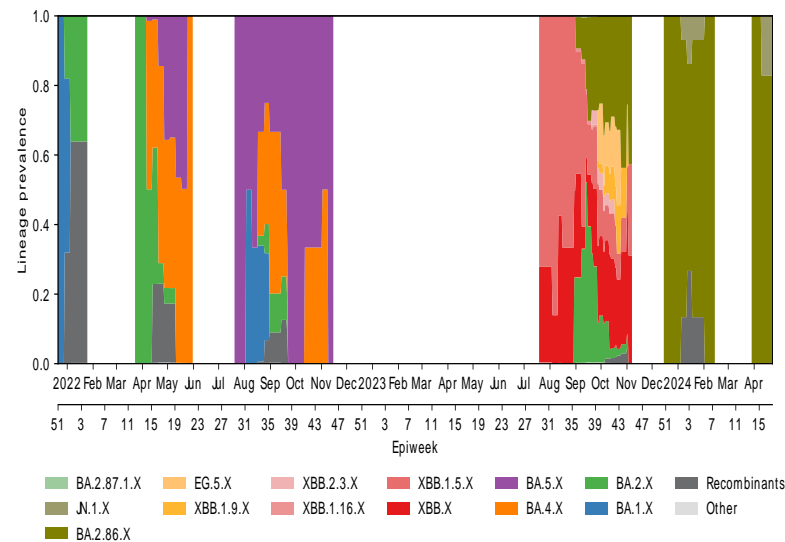
★ Chloroform start and end date

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.

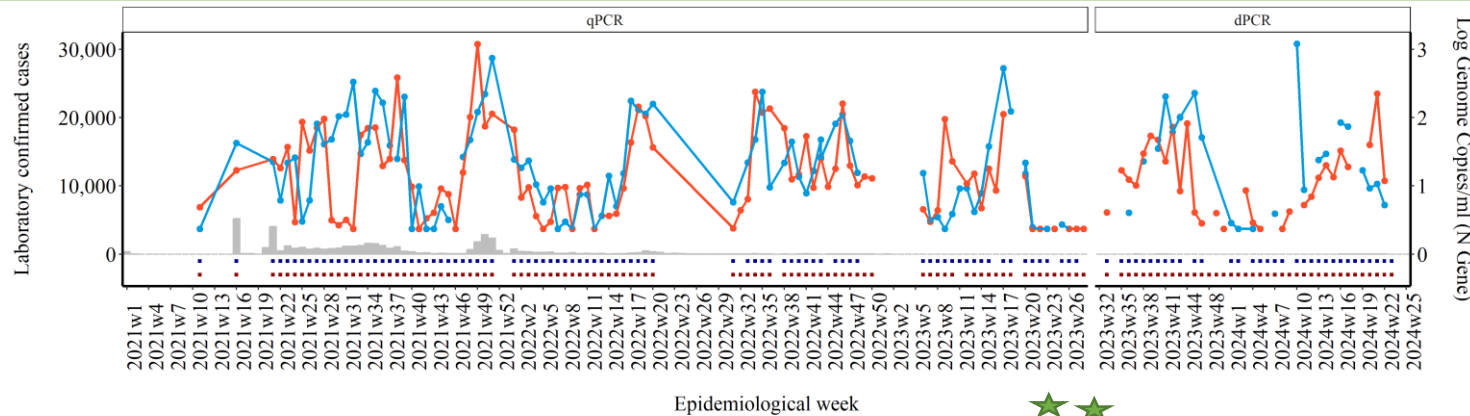
## East Bank Wastewater Treatment Works



## Mdantsane Wastewater Treatment Works



# Free State – Mangaung



★ Chloroform start and end date

• Bloemspruit Sample Collection • Bloemspruit Wastewater Treatment Works • Sterkwater Sample Collection • Sterkwater 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 Mangaung (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 23:

- 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. Levels increase to high in Epi week 21 and decrease to moderate in Epi week 23.
- 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. Levels remain low in Epi week 23.

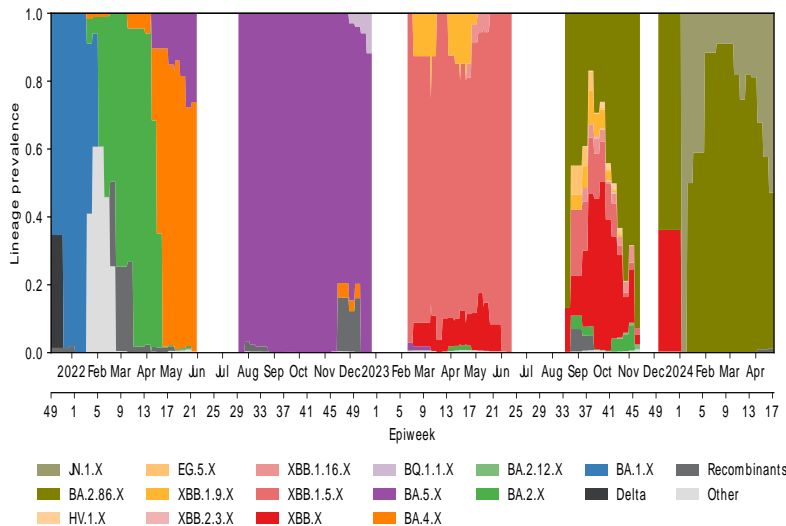
**\* Sequencing data ending in Epi week 17 in Bloemspruit and Epi week 13 in Sterkwater Treatment Works.**

- 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.

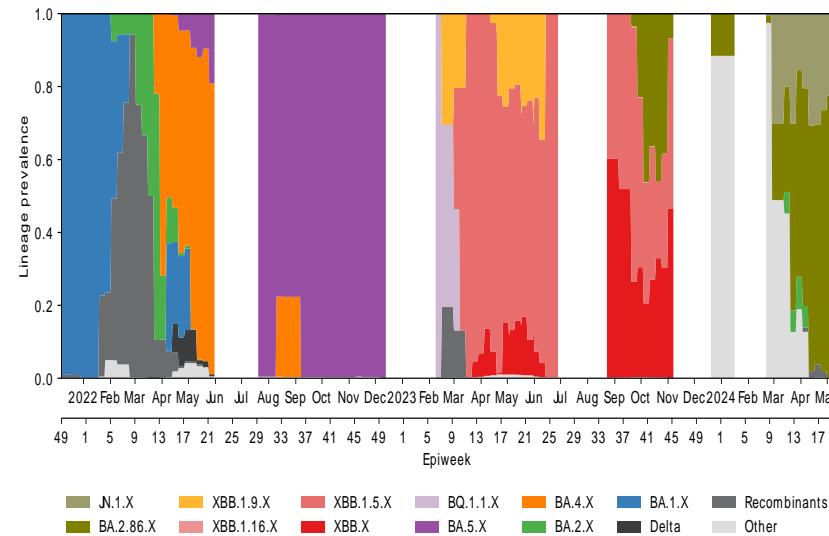
## SNP Analysis:

- 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.

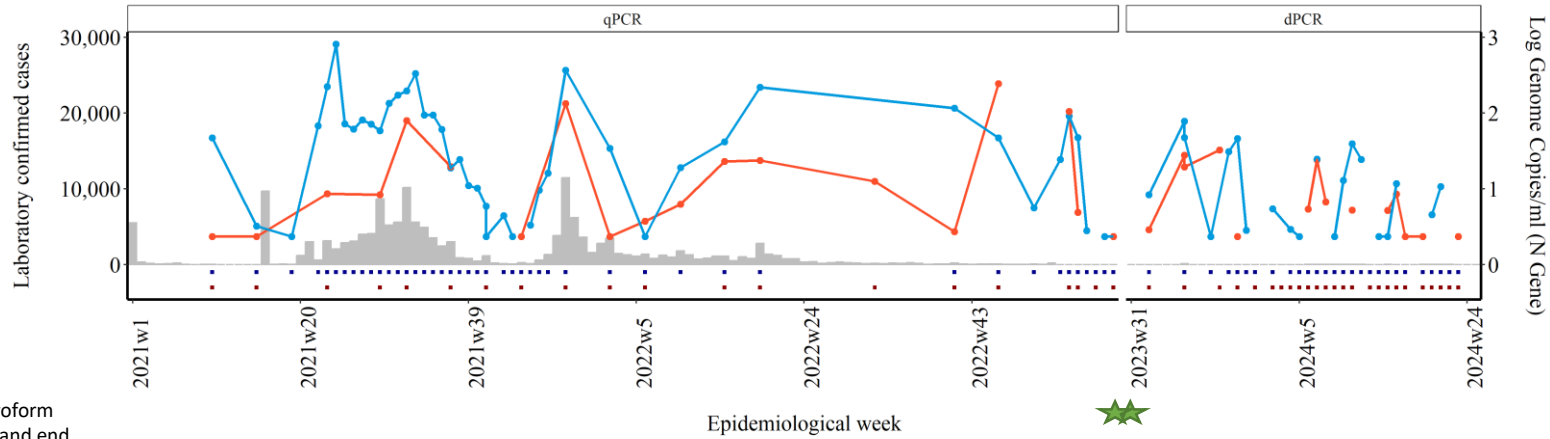
## Bloemspruit Wastewater Treatment Works



## Sterkwater Wastewater Treatment Works



# Western Cape – City of Cape Town



★ Chloroform start and end date

• Borcheds Quarry Sample Collection • Borcheds Quarry Wastewater Treatment Works • Zandvleit Sample Collection • Zandvleit Wastewater Treatment Works

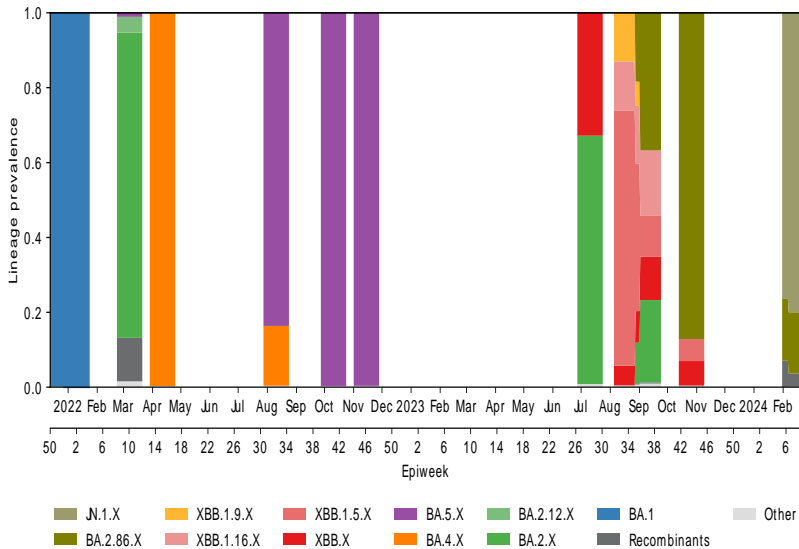
Wastewater levels of SARS-CoV-2 in log-transformed genome copies/ml (right vertical axis, line graph) and laboratory confirmed cases of SARS-CoV-2 for Cape Town (left vertical axis, grey bars) by epidemiological week. Coloured dots below the horizontal axis signify if a sample was collected. SARS-CoV-2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

## SARS-CoV-2 levels and Genomic Results in Epi week 23:

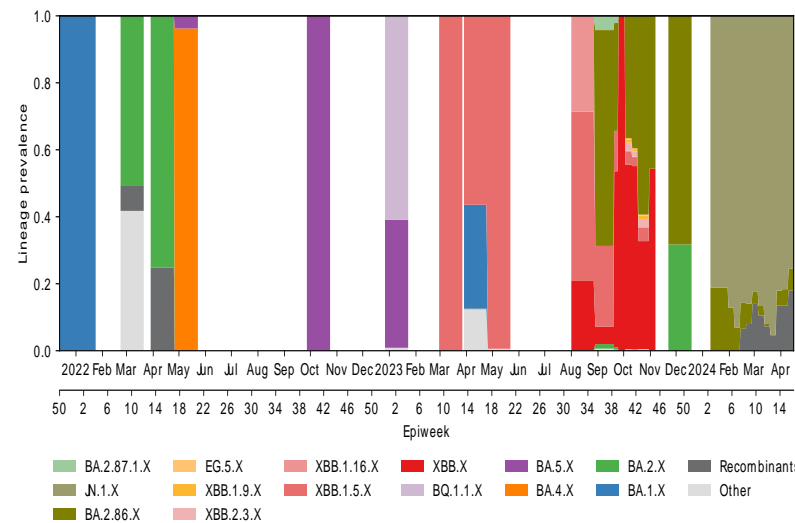
- After a sharp increase in SARS-CoV-2 levels was seen in Epi week 37, a subsequent decrease in SARS-CoV-2 levels in Borcheds Quarry WWTW was seen. Levels are low in Epi week 6 and increase to moderate in Epi week 7. Levels decrease in Epi week 8 to low and remain low in Epi week 23.
- In Epi week 4, 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 and remain moderate in Epi week 21. No new results for Epi week 23 are available.

**\* Sequencing data ending in Epi week 6 in Borcheds Quarry and Epi week 10 in Zandvleit.**

## Borcheds Quarry Wastewater Treatment Works



## Zandvleit Wastewater Treatment Works

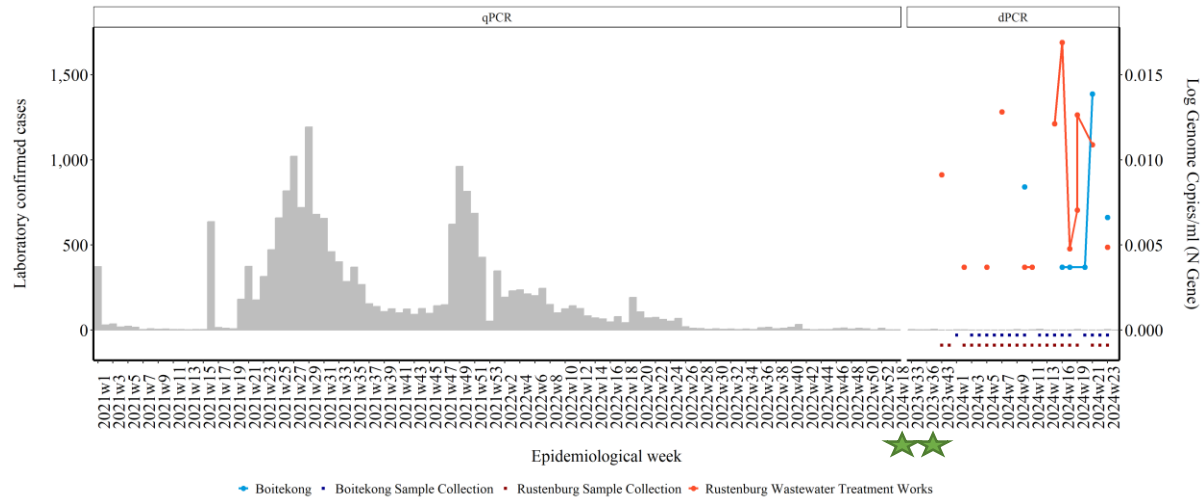


- During Epi week 6, JN.1.X sub-lineages were dominantly circulating in Borcheds. BA.2.86.X and Recombinants were also detected.
- During Epi week 2 - 6, lineages BA.2.86.X and JN.1.X were circulating in Zandvleit, with JN.1.X sub-lineages being dominant.
- In epi week 10 and 14 BA.2.86.X, JN.1.X and Recombinants lineages were circulating in Zandvleit, 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 Zandvleit.

# North West – Bojanala Platinum



★ Chloroform start and end date

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.

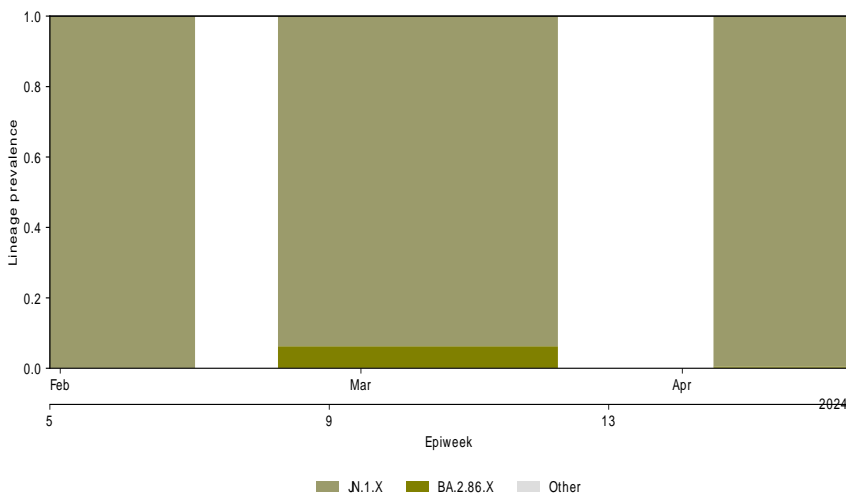
\* SARS-CoV-2 wastewater sample collection and testing at Boitekong began in Epi week 1, 2024.

SARS-CoV-2 levels and Genomic Results in Epi week 23:

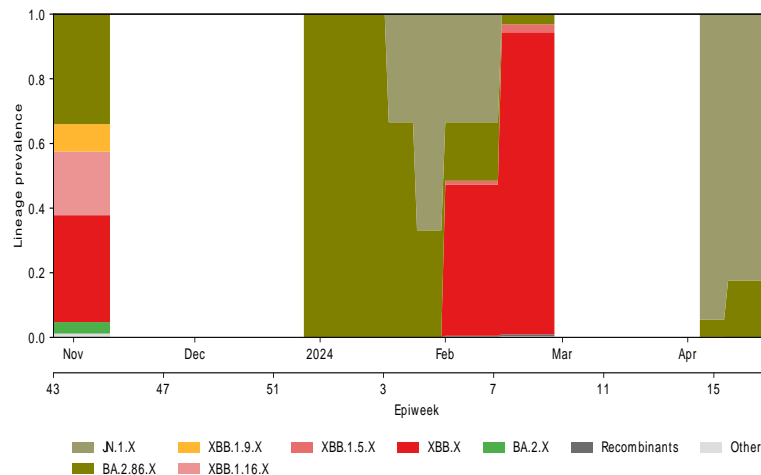
- At Rustenburg WWTW levels from Epi week 43 remain below 0.015 log genome copies/ml and remain low up until Epi week 23.
- At Boitekong wastewater results for SARS-CoV-2 were low in Epi week 10, 2024. Levels remain low in Epi week 16. No new results for Epi week 23 are available.

\* Sequencing data ending in Epi week 13 in Boitekong and Epi week 15 in Rustenburg.

## Boitekong



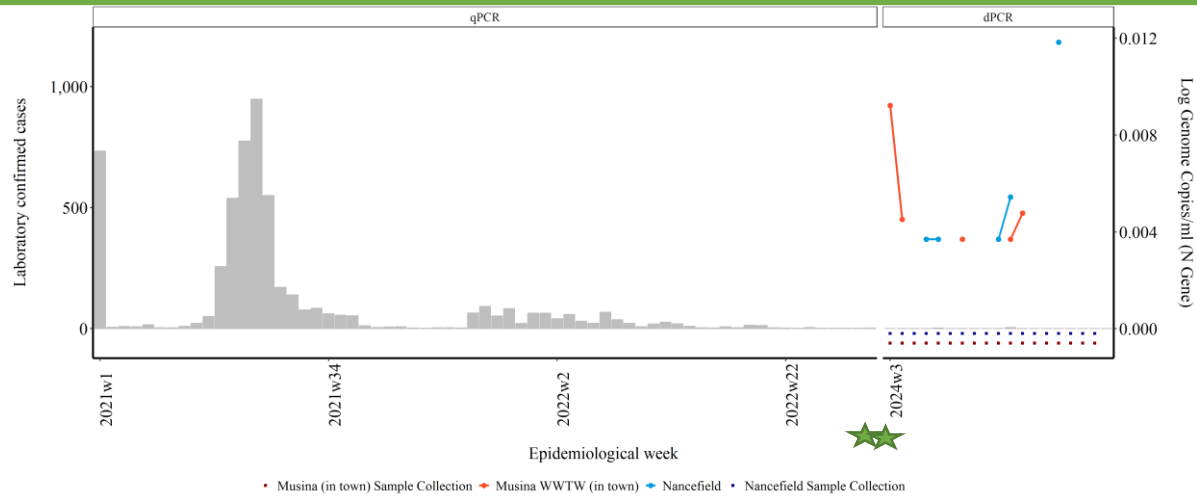
## Rustenburg



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



★ Chloroform start and end date

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 Vhembe (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 Musina WWTW began in Epi week 3, 2024.**

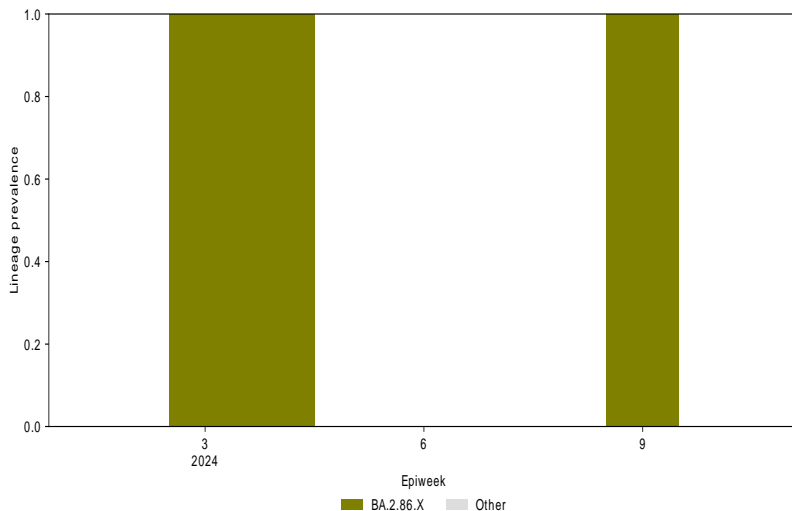
**\* SARS-CoV-2 wastewater sample collection and testing at Nancefield began in Epi week 3, 2024.**

SARS-CoV-2 levels and Genomic Results in Epi week 23:

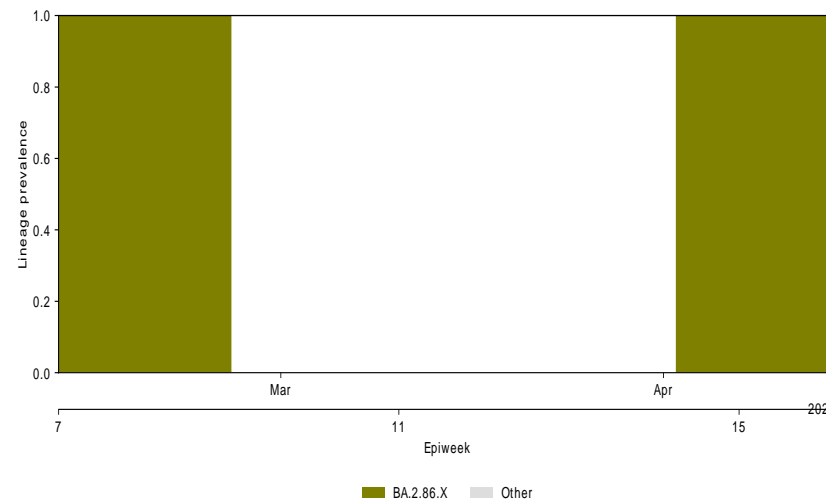
- At Musina WWTW levels from Epi week 3 remain below 0.01 log genome copies/ml and remain low up until Epi week 4, 2024. Levels remain low in Epi week 9. No new results for Epi week 23 are available.
- At Nancefield levels were low in Epi weeks 6 and 7, below 0.005 log genome copies/ml. They remain low in Epi week 12. No new results for Epi week 23 are available.

**\* Sequencing data ending in Epiweek 9 in Musina and Epi week 15 in Musina**

## Musina



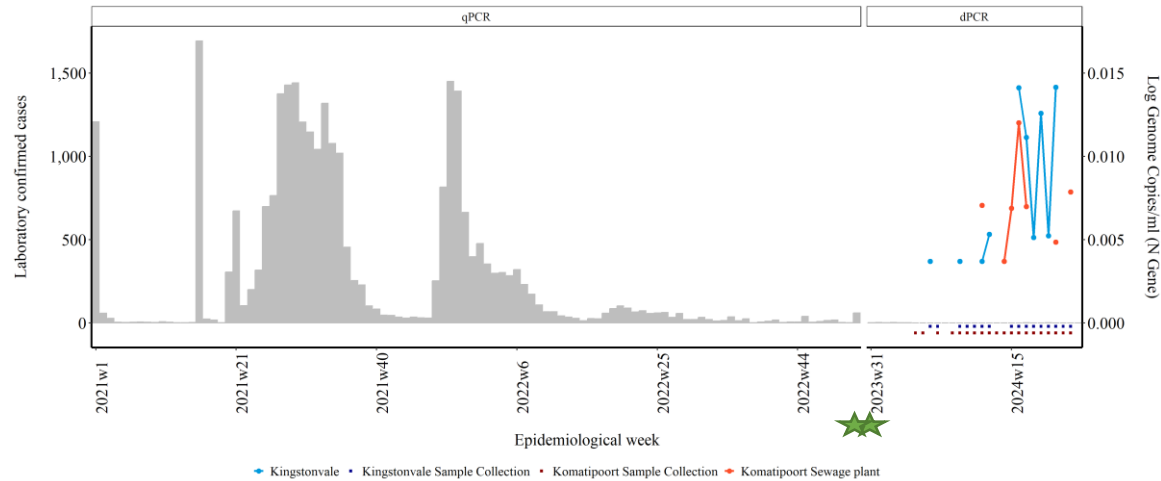
## Nancefield



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



★ Chloroform start and end date

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.

**\* 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.**

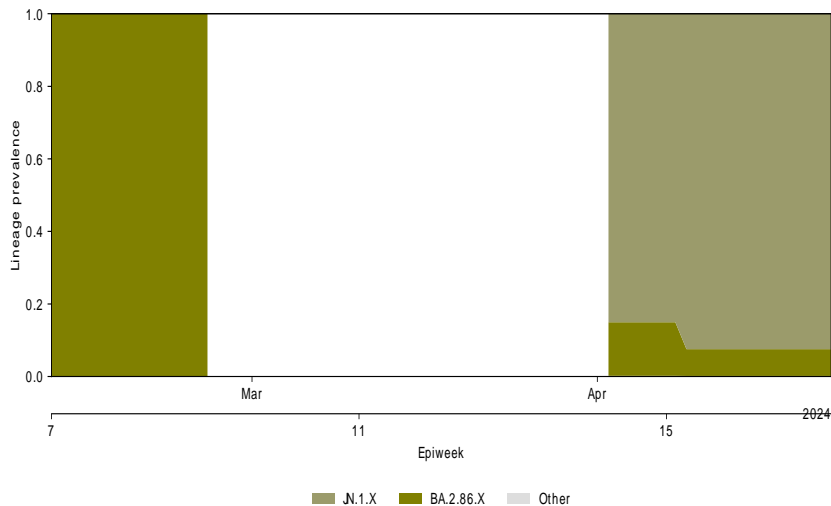
SARS-CoV-2 levels and Genomic Results in Epi week 23:

- 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 23.
- 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 21. No new result for Epi week 23 are available.

**\* 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.

## Kingstonvale



## Komatipoort

**\*No sequencing data currently available**

# COLLABORATORS



# FUNDERS



# TEAM

