Current Status of the Most Dominant SARS-CoV-2 Variant





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Session 14 December 2022



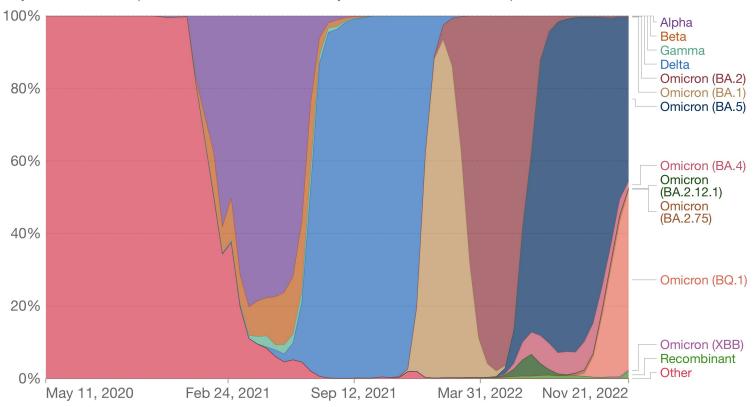




SARS-CoV-2 variants in analyzed sequences



The number of analyzed sequences in the preceding two weeks that correspond to each variant group. This number may not reflect the complete breakdown of cases since only a fraction of all cases are sequenced.

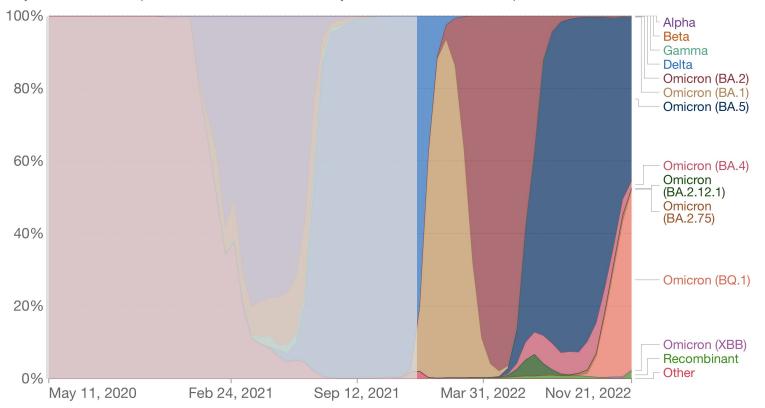


Source: GISAID, via CoVariants.org – Last updated 25 November 2022 OurWorldInData.org/coronavirus • CC BY Note: Recently-discovered or actively-monitored variants may be overrepresented, as suspected cases of these variants are likely to be sequenced preferentially or faster than other cases.

SARS-CoV-2 variants in analyzed sequences



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Omicron BA.1
Omicron BA.2
Omicron BA.5

Omicron XBB

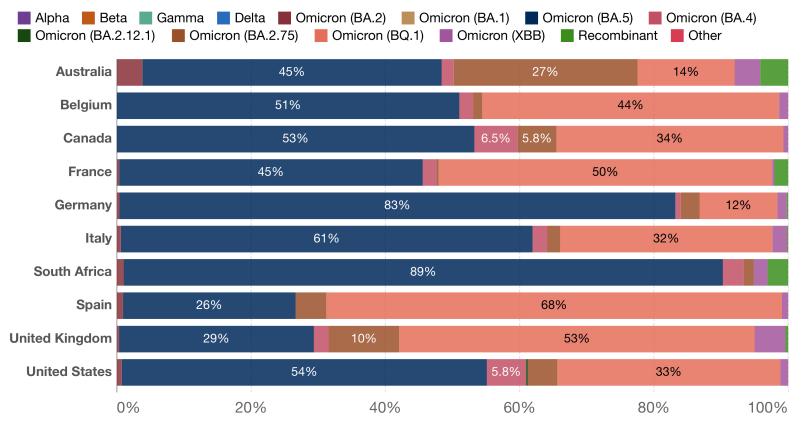
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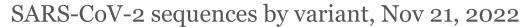




The share of analyzed sequences in the preceding two weeks that correspond to each variant group. This share may not reflect the complete breakdown of cases since only a fraction of all cases are sequenced.

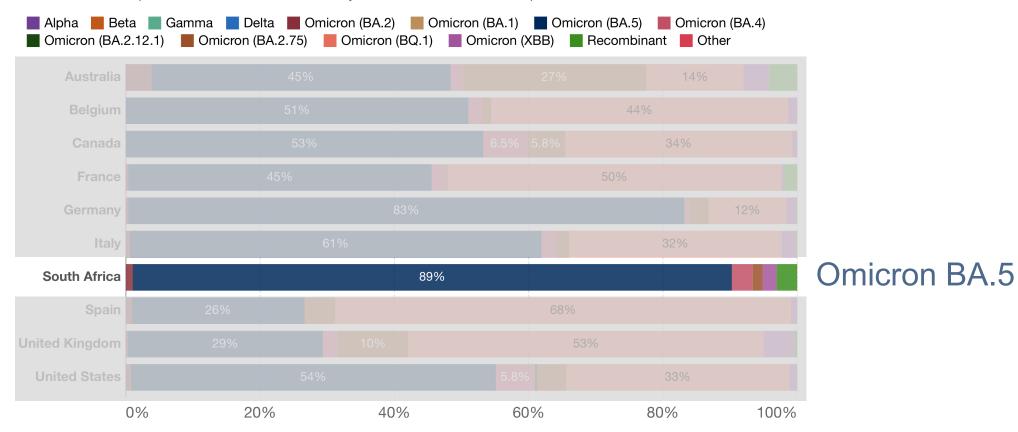


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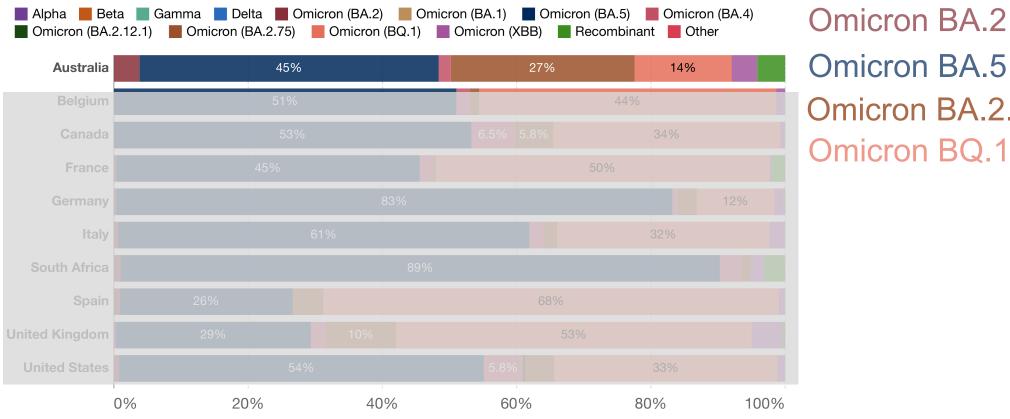
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SARS-CoV-2 sequences by variant, Nov 21, 2022



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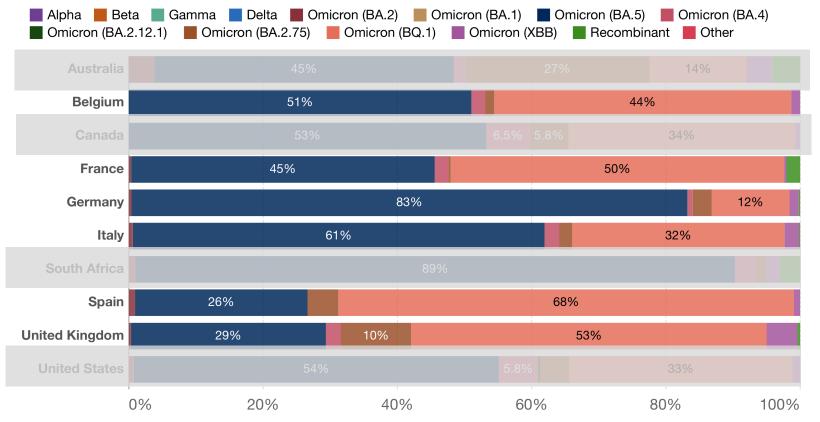
Omicron BA.2 Omicron BA.5 Omicron BA.2.75

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SARS-CoV-2 sequences by variant, Nov 21, 2022



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Omicron BA.2

Omicron BA.5

Omicron BA.2.75

Omicron BQ.1

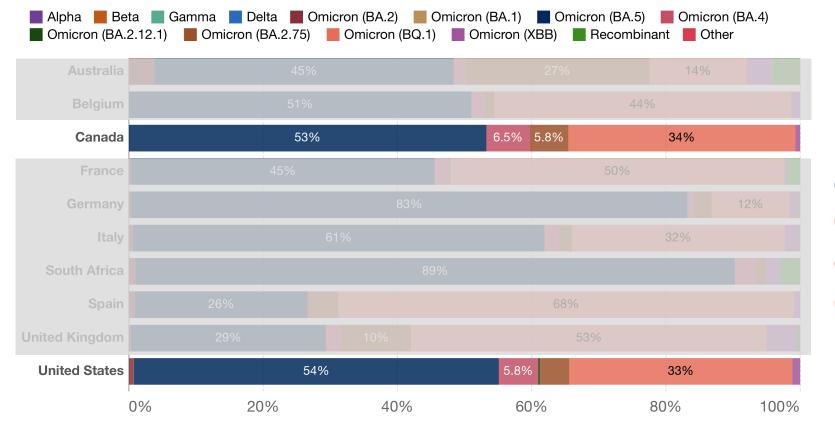
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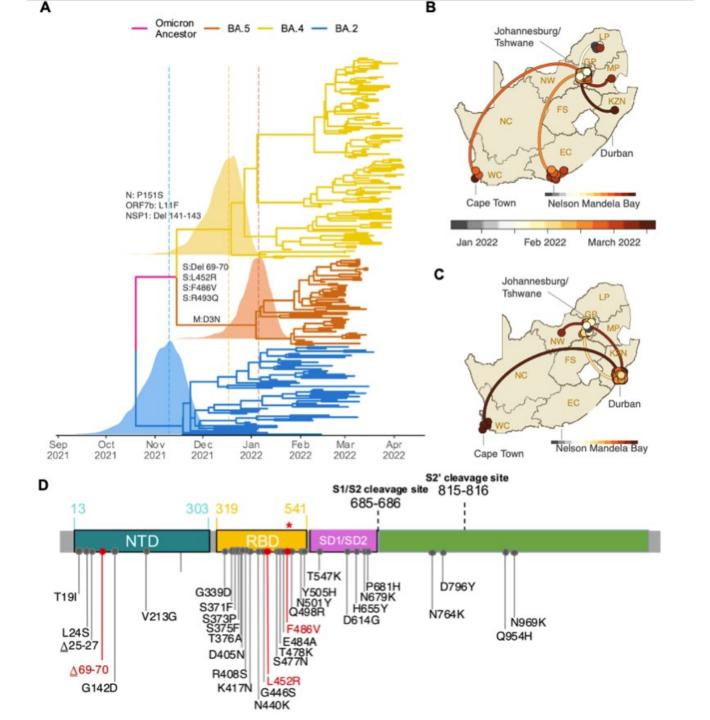


Omicron BA.5 Omicron BA.4** Omicron BA2.75 Omicron BQ.1

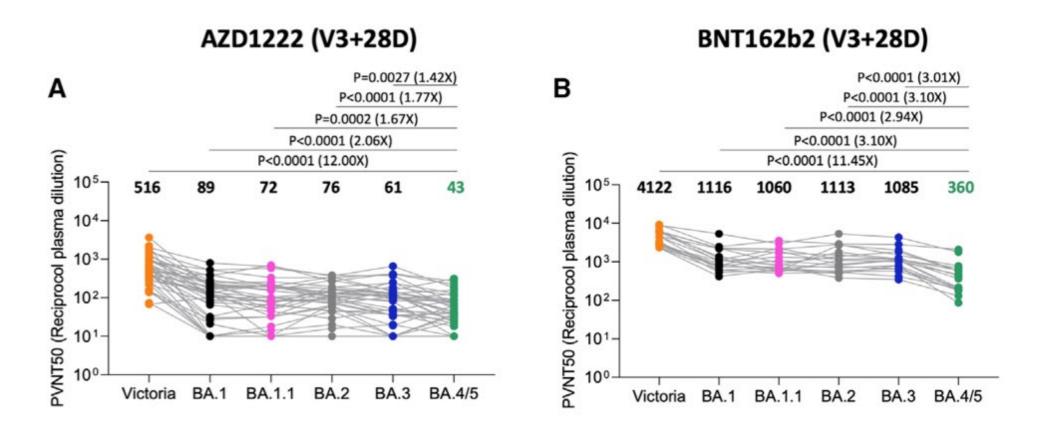
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How did BA.4/5 sweep across the world so fast?



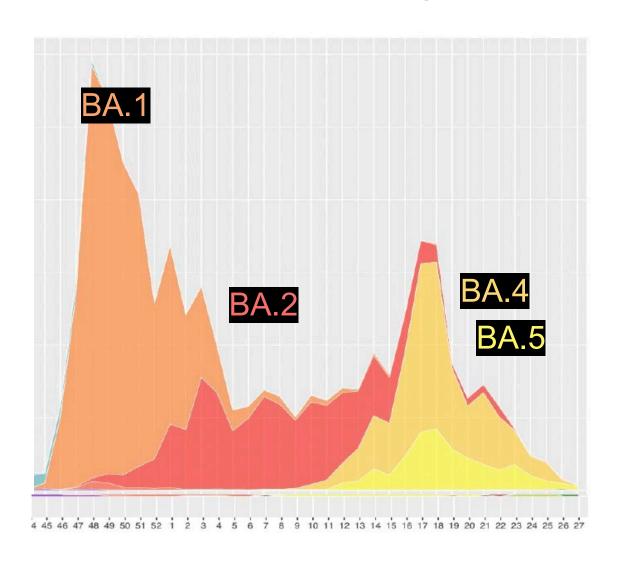
High level resistance of BA.4/5 to vaccine sera

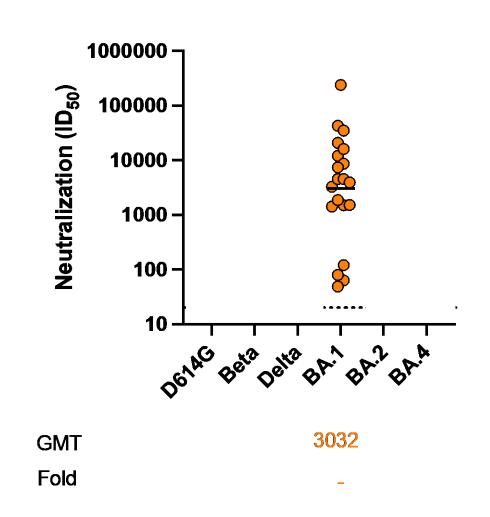


High level resistance of BA.4/5 to therapeutically relevant mAbs

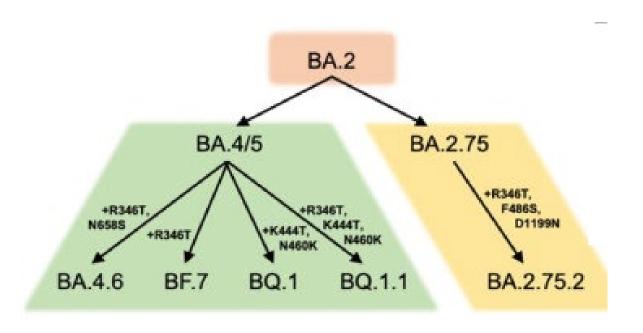
	IC50 (μg/mL) Pseudoviruses					
mAbs	Victoria	BA.1	BA.1.1	BA.2	BA.3	BA.4/5
REGN10987	0.002	10	10	0.616	10	10
REGN10933	0.001	10	10	10	10	10
AZD1061	0.002	0.308	10	0.008	0.019	0.015
AZD8895	0.001	0.246	0.1	1.333	10	10
AZD7442	0.001	0.232	0.806	0.008	0.065	0.065
ADG10	0.007	10	10	10	10	10
ADG20	0.003	0.348	0.253	10	10	10
ADG30	0.014	10	10	10	10	10
Ly-CoV-555	0.002	10	10	10	10	10
Ly-CoV16	0.014	10	10	10	10	10
S309	0.13	0.094	0.138	0.638	0.228	1.041

Omicron BA.1 triggered strain-specific responses, and was rapidly followed by a BA.4/5 wave





Neutralization profile of Omicron BQ.1 and Omicron BA2.75, and their offspring



Panke Qu, Shan-Lu Liu et al, BioRxiv, 2022



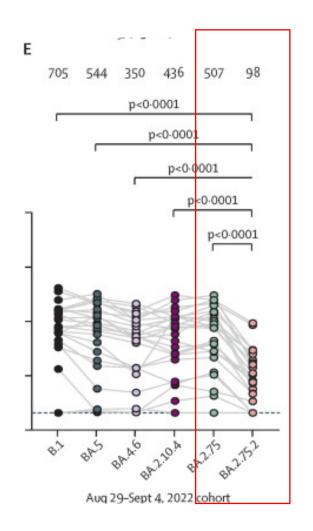


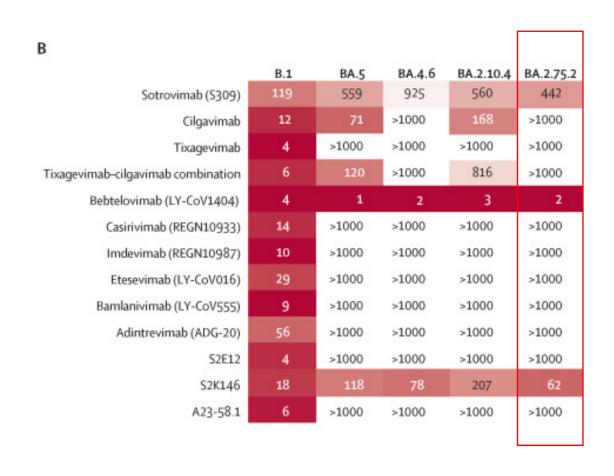
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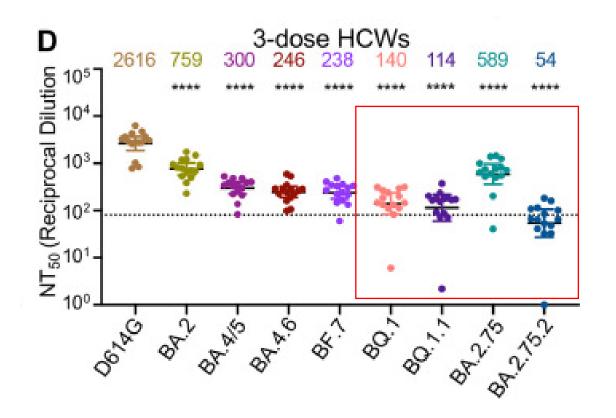
Omicron BA.5
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Omicron BA2.75
Omicron BQ.1

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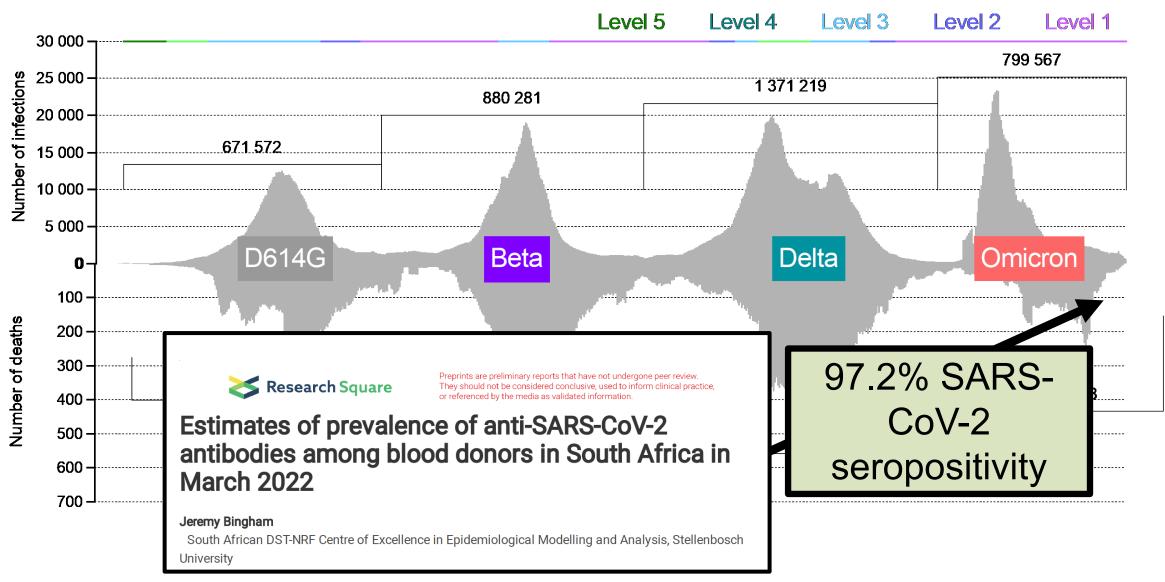




Neutralization profile of Omicron BQ.1 and Omicron BA2.75, and their offspring

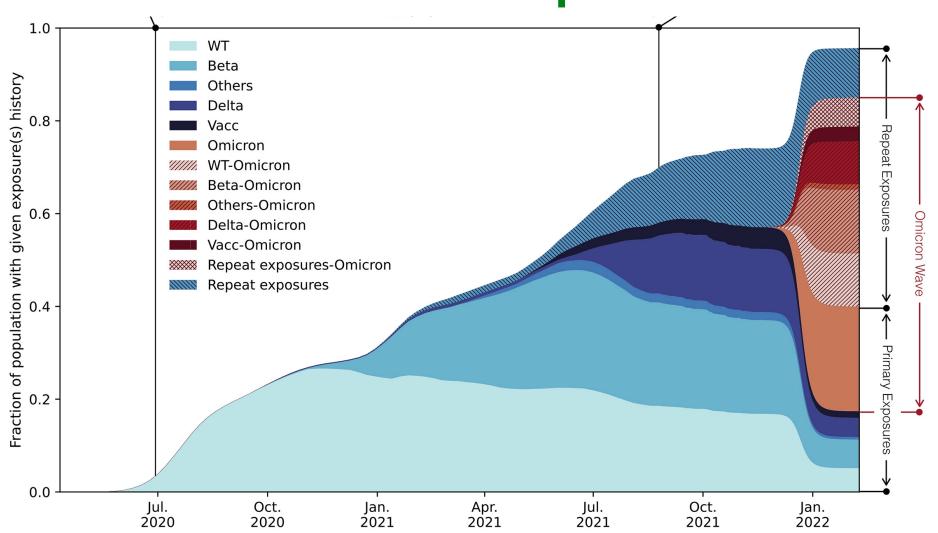


South Africa's epidemic waves and variants...



Bingham et al, Research Square

However, its a very complex hybrid immune landscape



Decoupling of infections and hospitalizations

ORIGINAL ARTICLE

Population Immunity and Covid-19 Severity
with Omicron Variant in South Africa

Shabir A. Madhi, Ph.D., Gaurav Kwatra, Ph.D., Jonathan E. Myers, M.D.,
Waasila Jassat, M.Med., Nisha Dhar, Ph.D., Christian K. Mukendi, M.Sc.,
Amit J. Nana, B.Sc., Lucille Blumberg, M.Med., Richard Welch, B.Sc.,

Few deaths in SA in the Omicron waves, due to high population immunity

Nicoletta Ngorima-Mabhena, M.B., Ch.B., and Portia C. Mutevedzi, Ph.D.

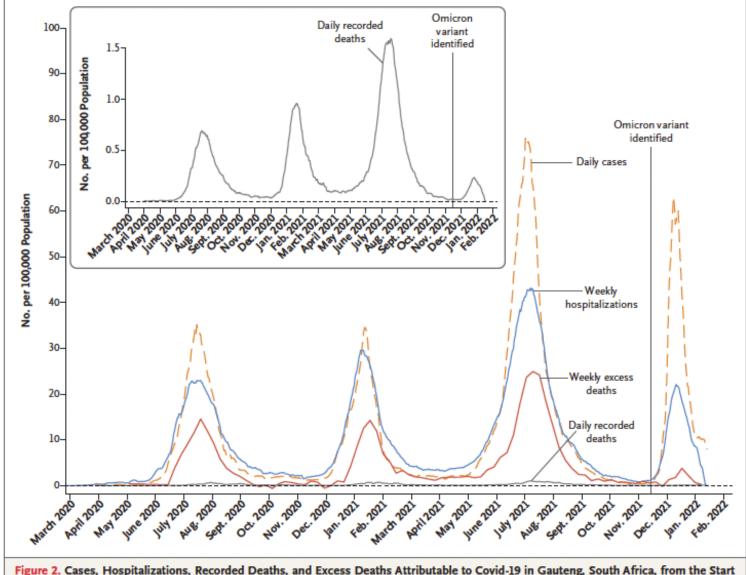


Figure 2. Cases, Hospitalizations, Recorded Deaths, and Excess Deaths Attributable to Covid-19 in Gauteng, South Africa, from the Start of the Pandemic through January 12, 2022.

Summary

- New variants continue to emerge, and there is local variation in which variant dominates
- This may be related to prior immune histories, as well as intrinsic virologic factors
- Emerging variants show high levels of neutralization escape, compromising mAb treatments
- However, population immunity is extremely high (mostly) and T cells and non-neutralizing antibodies remain resilient against emerging VOCs, reducing severity of clinical outcomes

Acknowledgements





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NICD

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Duke

David Montefiori

WITS

Shabir Madhi Patrick Arbuthnot

MRC

Glenda Gray Linda-Gail Bekker

Malawi

Kondwani Jambo

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Marion Vermeulen Karin vd Berg

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BILL&MELINDA GATES foundation

