

Introduction

17:00- 17.10

Prof. Dr. Carmen Faso, MCID Co-Chair Prof. Dr. med. Nicola Low (Moderator)

Speaker presentation

- **17:15- Dr. Emma Hodcroft,** Institute of Social and Preventive Medicine, UniBern "What is Omicron and how do we identify it?"
- 17:30- Prof. Dr. Volker Thiel, Institute for Virology and Immunology, Vetsuisse, UniBern "How is Omicron different from other variants?"
- **17:45- PD Dr. Christian Althaus**, Institute of Social and Preventive Medicine, UniBern "How well does Omicron spread compared to other variants?"
- **18:00- Prof. Dr. med. Annelies Wilder-Smith,** Institute of Social and Preventive Medicine, UniBern "What does Omicron mean for international travel?"
- 18:15- Prof. Dr. Joseph Francois, World Trade Institute, UniBern"Why and how does Omicron cause an international economic shock?"

Panel discussion

Including the speakers above, together with:

Prof. Dr. Dr. Claus Beisbart Institute of Philosophy, UniBern

PD Dr. med. Manuela Funke-Chambour Department for Biomedical Research, Inselspital



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Multidisciplinary Center for Infectious Diseases

MCID Bern

www.mcid.unibe.ch





How well does Omicron spread compared to other variants?

Multidisciplinary Center for Infectious Diseases (MCID) Event: The SARS-CoV-2 variant Omicron: a snapshot of where we are 8 December 2021

Christian L. Althaus, Ph.D.

Institute of Social and Preventive Medicine University of Bern Mittelstrasse 43 3012 Bern

Image: Nextstrain

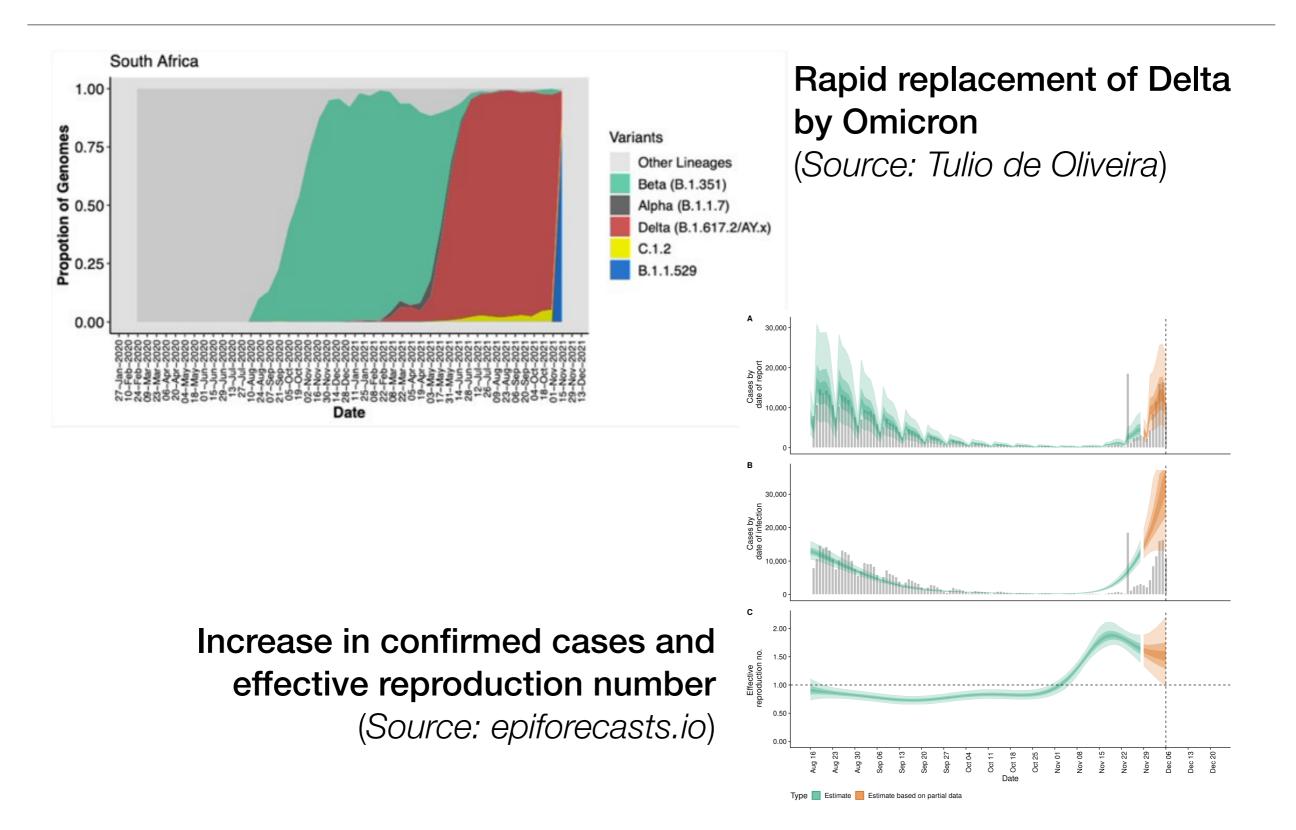




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Fonds national suisse Schweizerischer Nationalfonds Fondo nazionale svizzero Swiss National Science Foundation

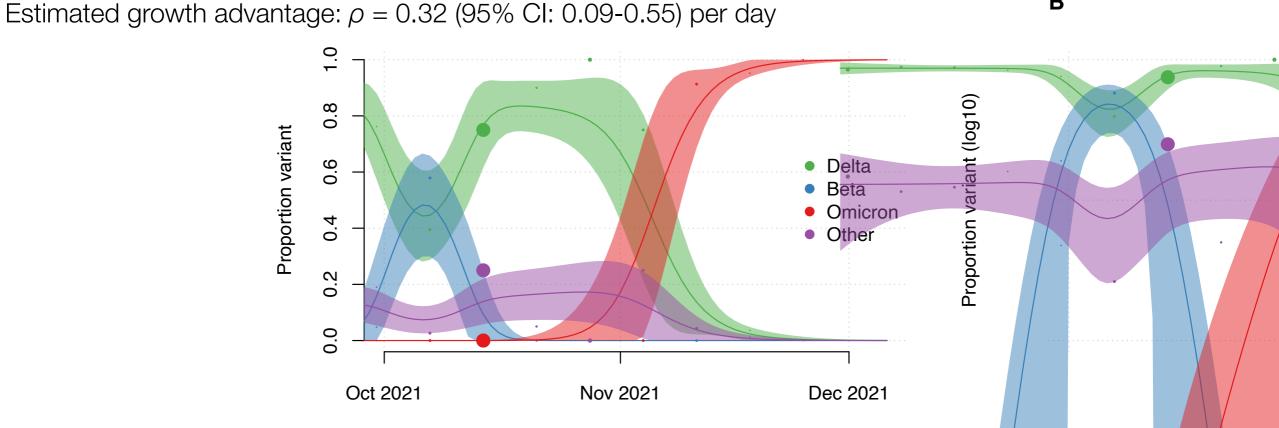
Growth of Omicron in South Africa



Omicron vs. Delta in Gauteng, South Africa

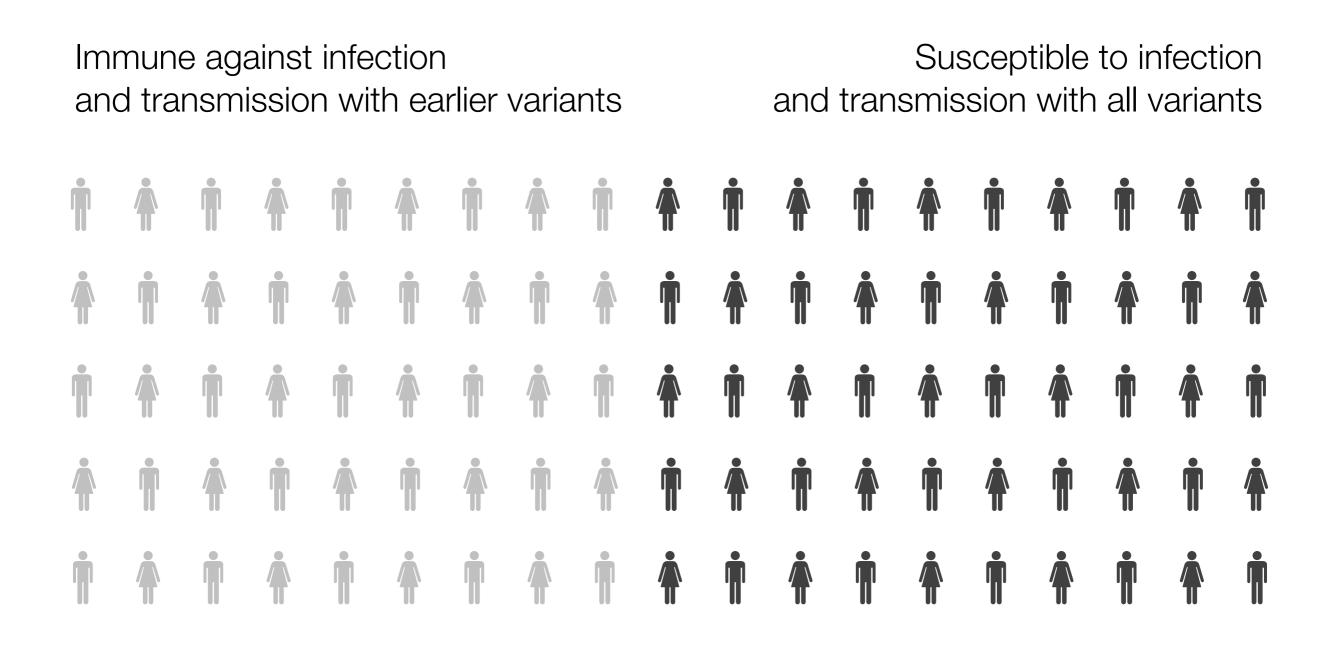
• GISAID: 569 sequences from Gauteng, South Africa (1 Sep - 22 Nov 2021)

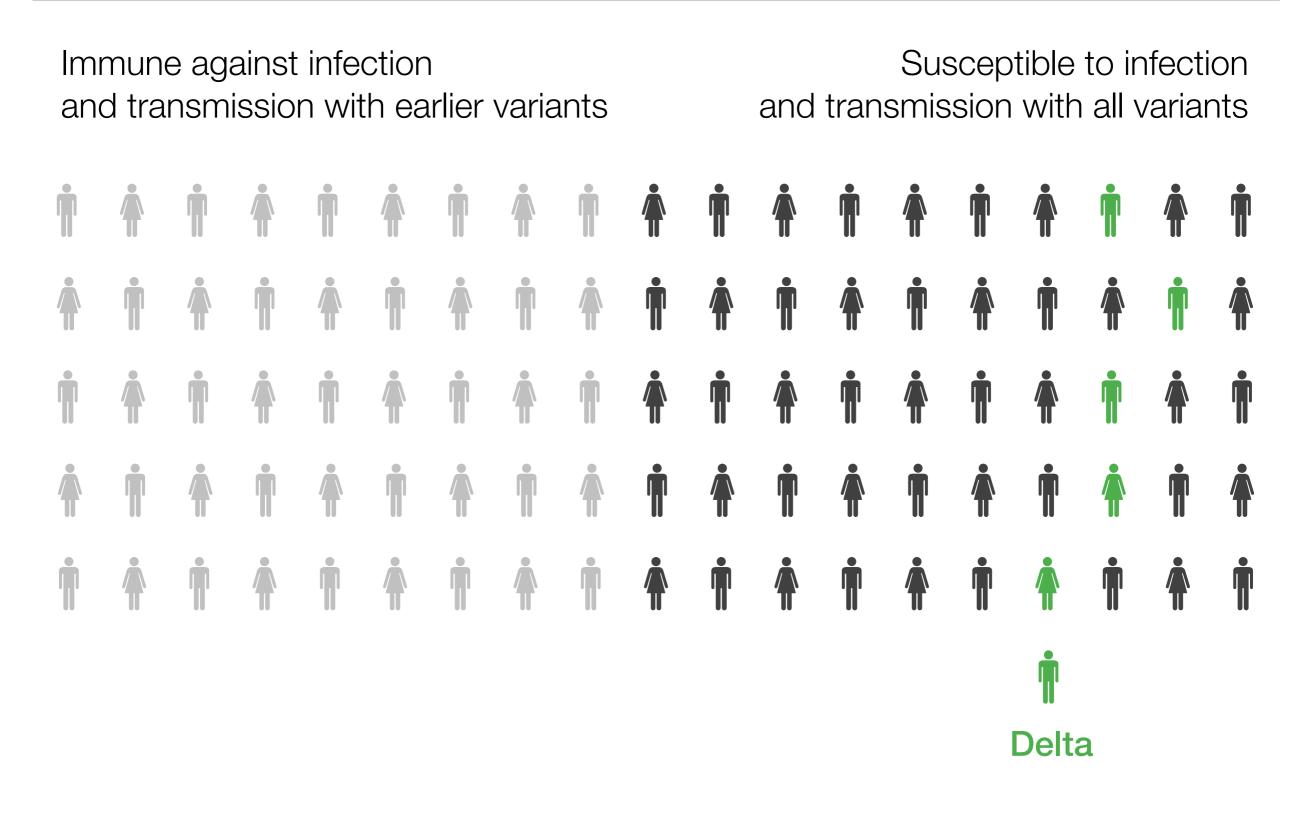
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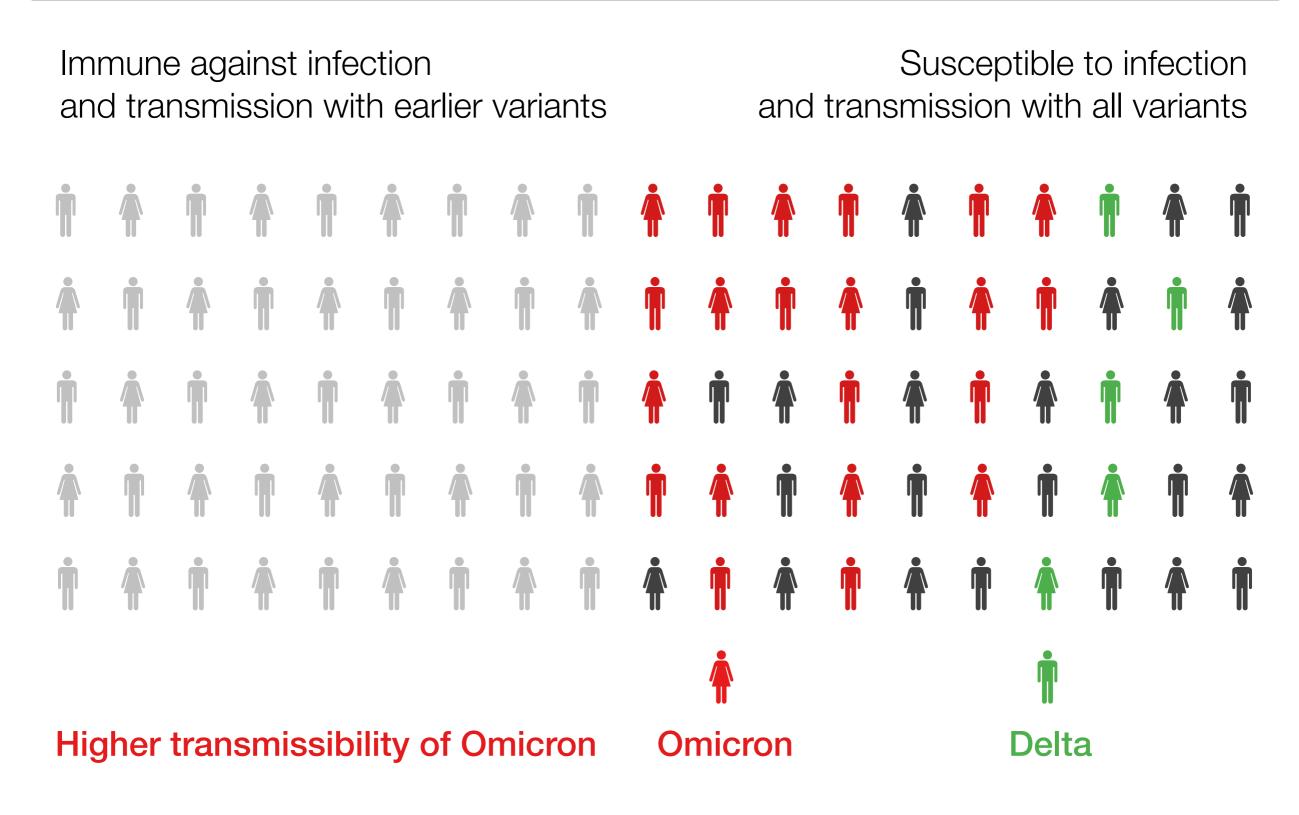


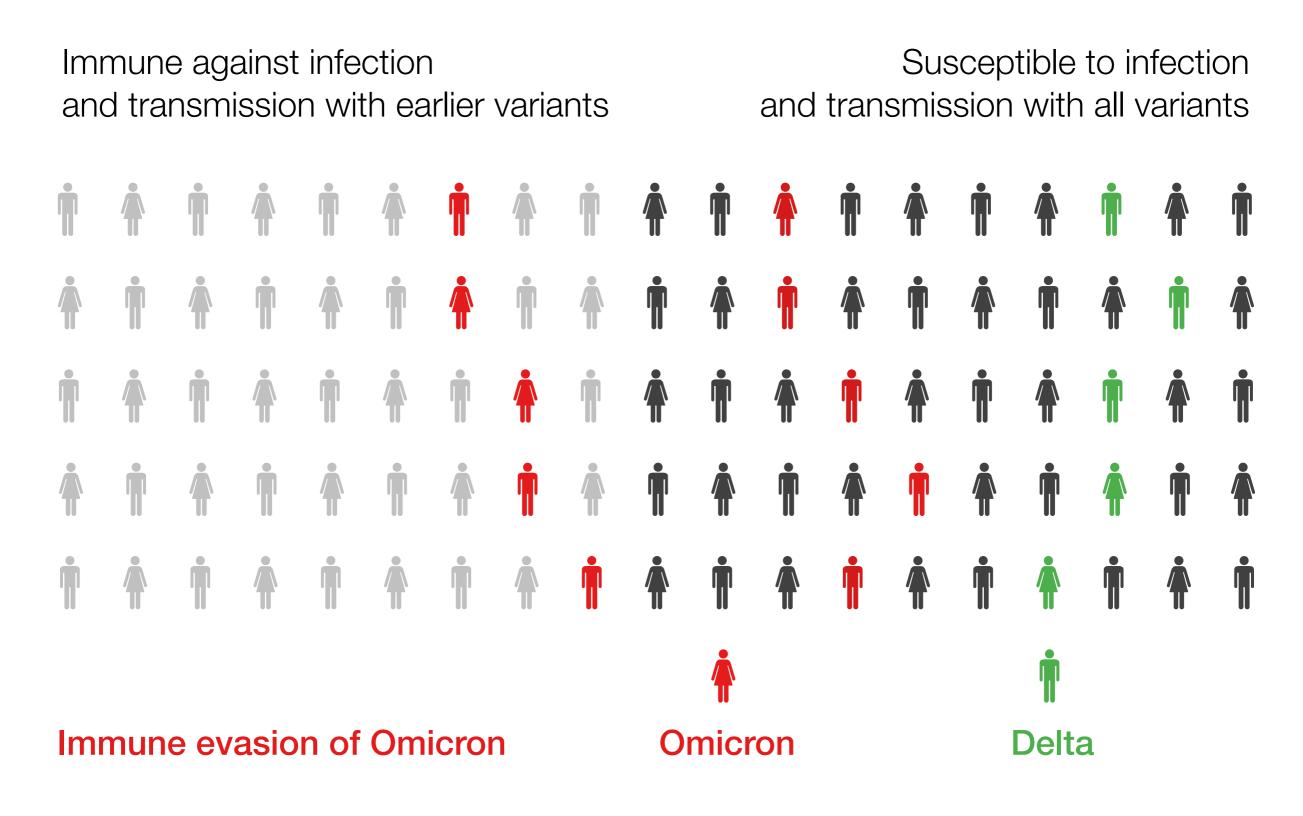
- Caution: Targeted sequencing and stochastic effects (e.g., superspreading) can lead to overestimates of the growth advantage.
- But: Consistent with difference in rate (or R_e) of epidemic decline (Delta, Sep-Oct) and growth (Omicron, Nov-Dec)

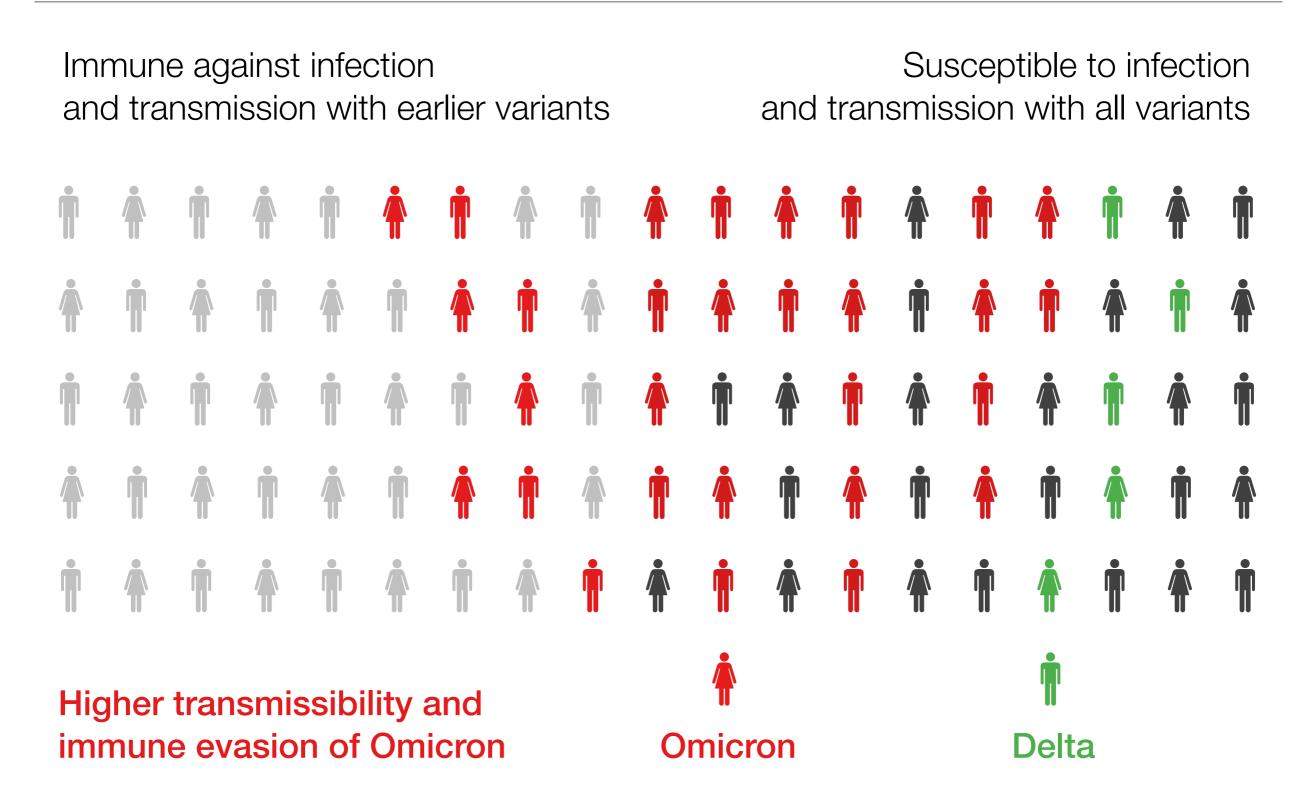
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Seroprevalence in South Africa

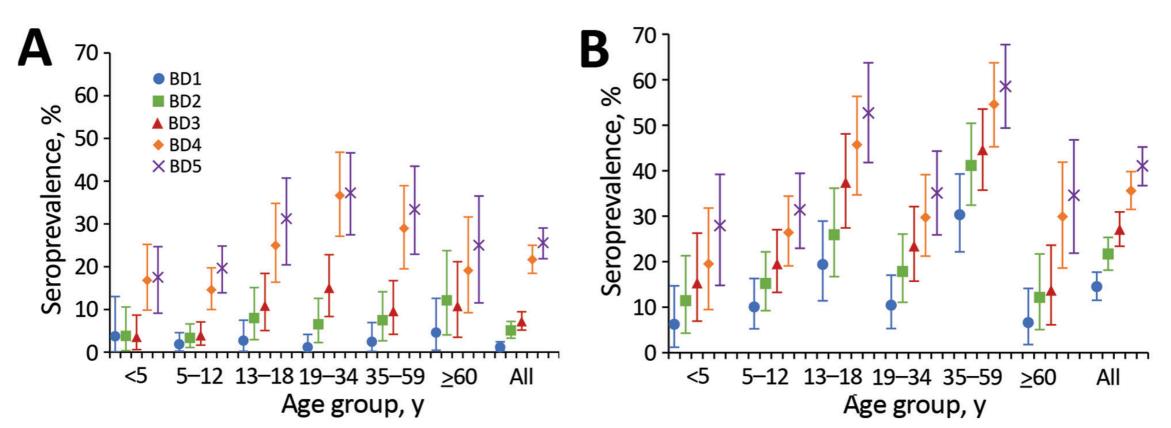
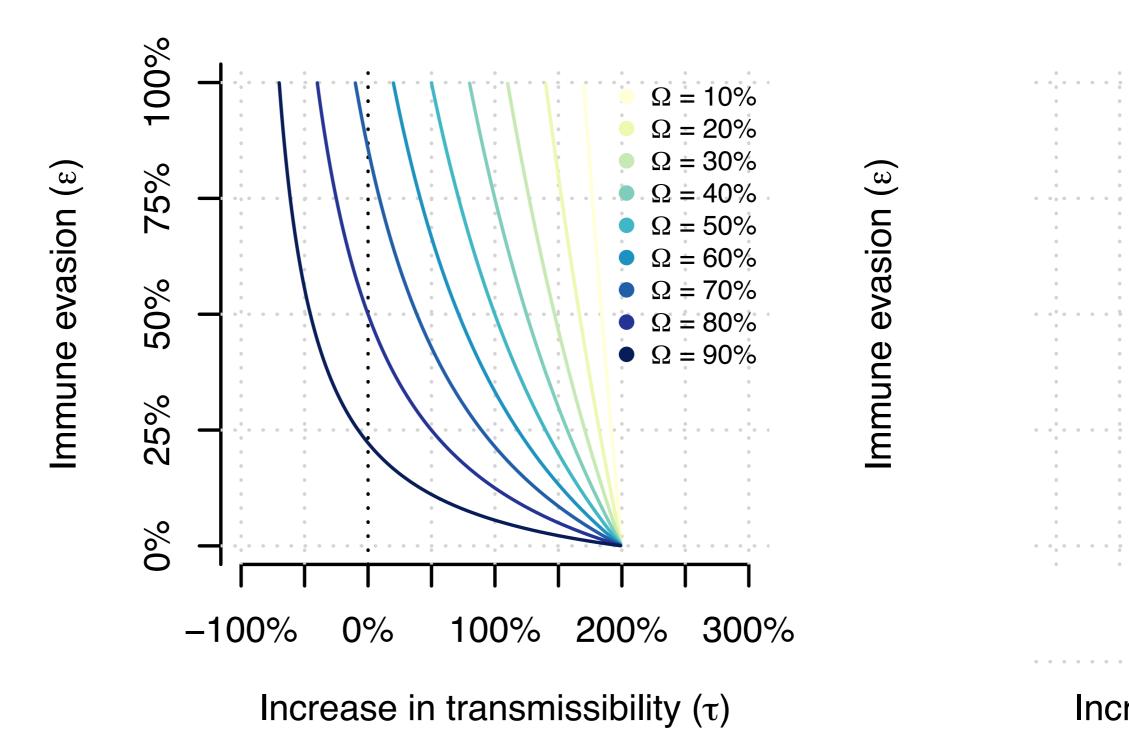


Figure 2. Seroprevalence of severe acute respiratory syndrome coronavirus 2 at each blood collection, by age group, in a rural community (A) and an urban community (B), South Africa, March 2020–March 2021.

Seroprevalence (and level of protective immunity against infection and subsequent transmission) arguably higher after Delta wave in South Africa. In addition, almost 30% of the population has been vaccinated.

Reference: Kleynhans et al. (2021, Emerg Infect Dis)

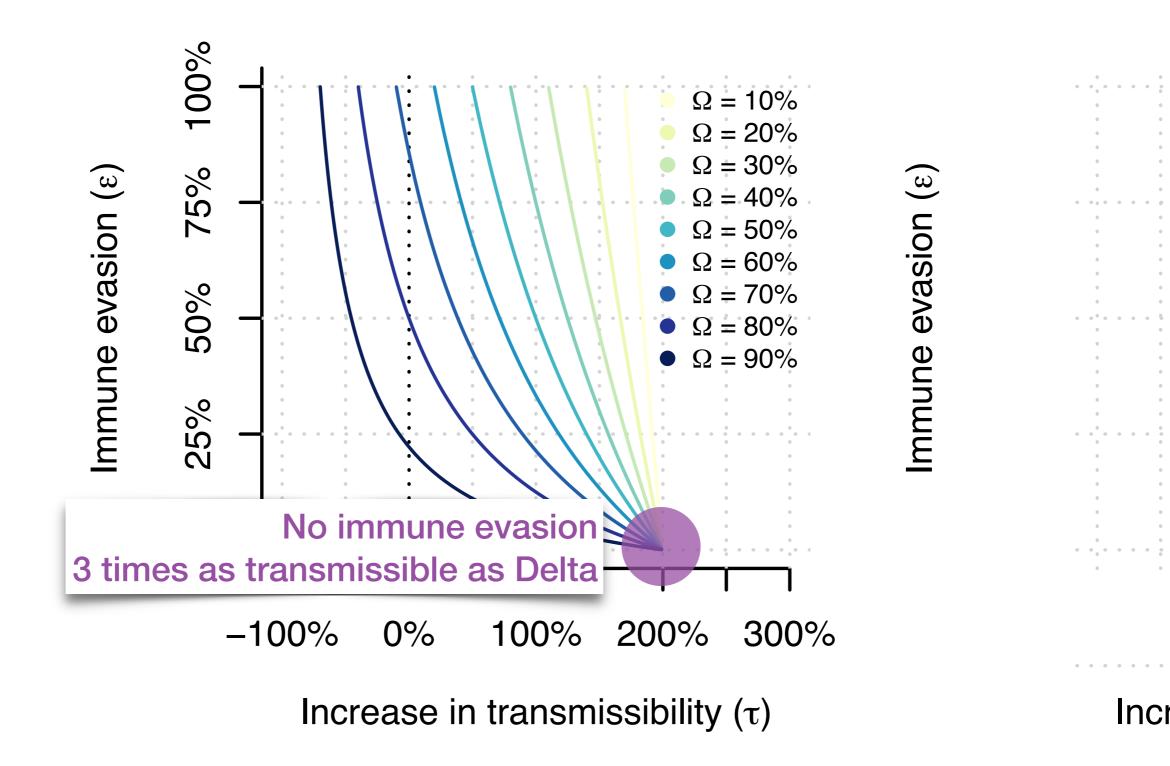
Relationship between potential increase in transmissibility and immune evasion for Omicron in Gauteng, South Africa



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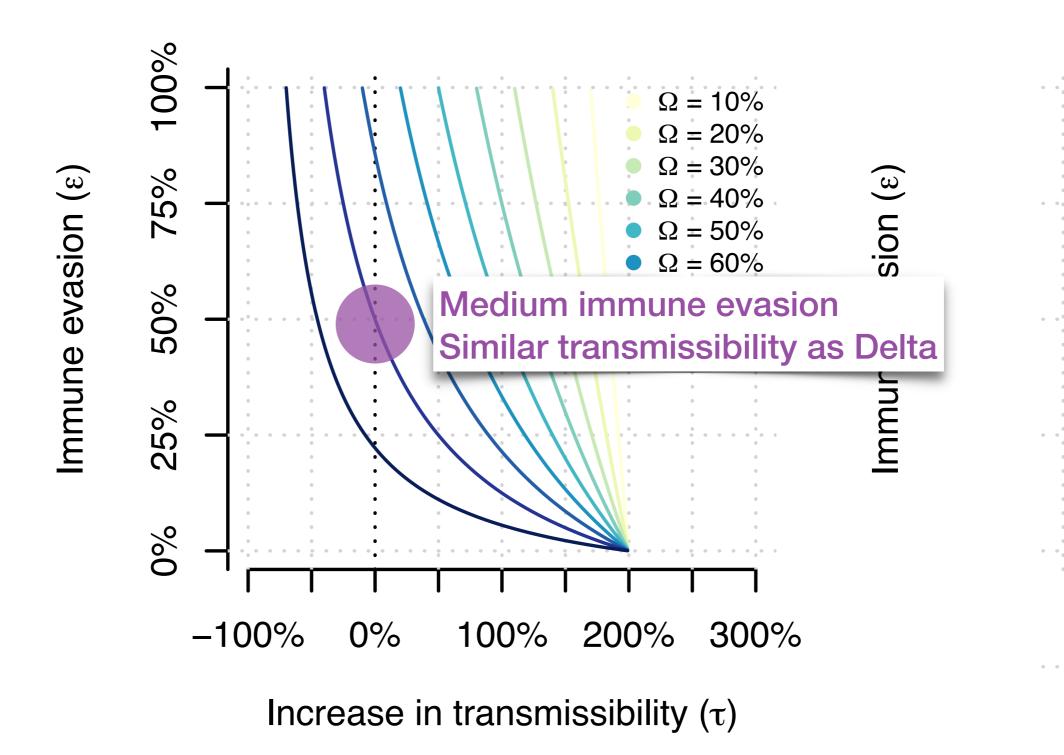
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Relationship between potential increase in transmissibility and immune evasion for Omicron in Gauteng, South Africa



В

Relationship between potential increase in transmissibility and immune evasion for Omicron in Gauteng, South Africa

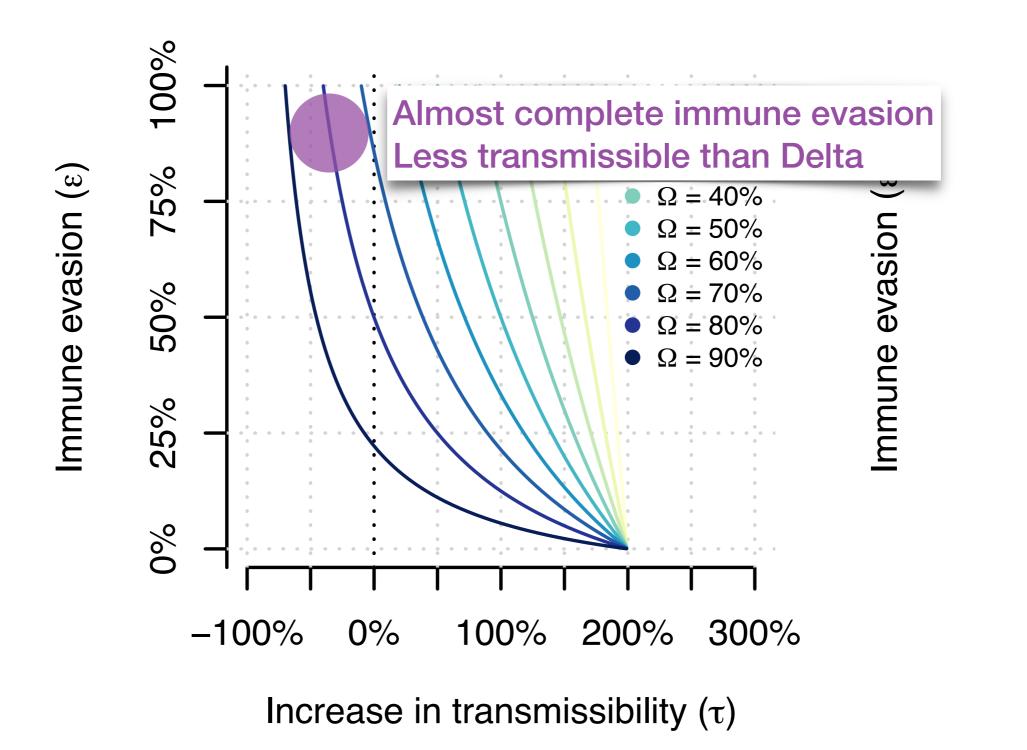


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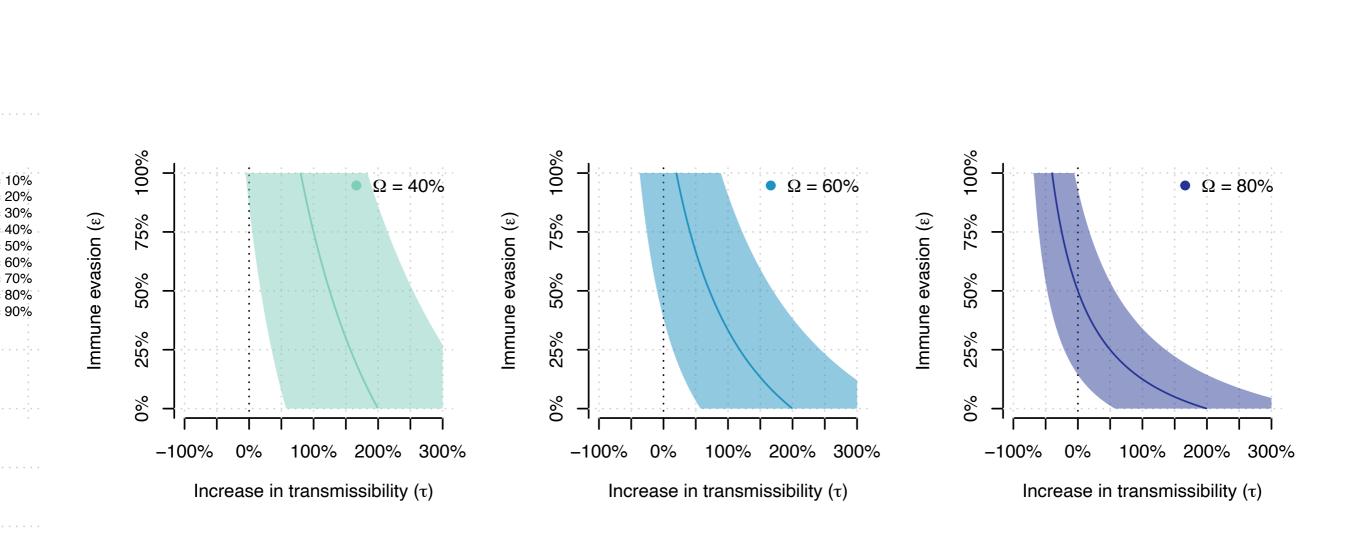
Relationship between potential increase in transmissibility and immune evasion for Omicron in Gauteng, South Africa:



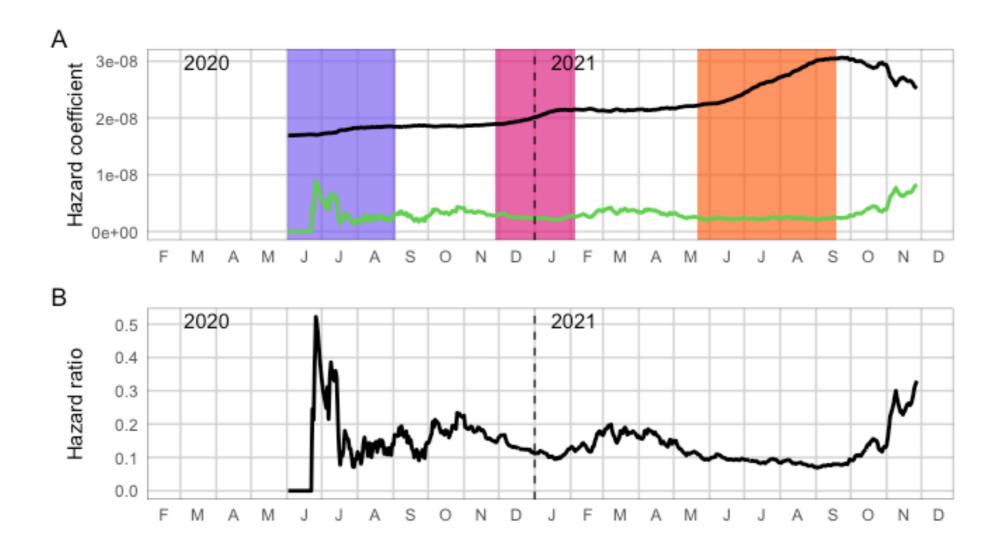
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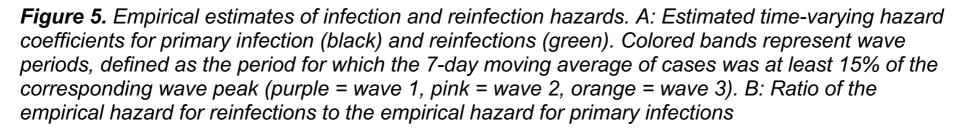
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Considerable uncertainty around the estimates remain



Further evidence for immune evasion of Omicron: Increased reinfection in South Africa





Summary

- Early evidence suggests that the SARS-CoV-2 variant Omicron has a considerable growth advantage compared to Delta in South Africa (~ 0.3 per day).
- Partial evasion of protective immunity against infection and transmission with previous variants can explain the observed dynamics, particularly when population immunity is high.
- An increase or decrease of the intrinsic transmissibility of Omicron compared to Delta cannot be excluded, however.
- Close monitoring of the spread of Omicron in countries outside South Africa will be necessary to better understand the potential of this variant to evade naturally acquired and vaccine-elicited immunity.

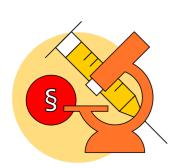
Thanks

- Emma Hodcroft, Martina Reichmuth Institute of Social and Preventive Medicine, University of Bern, Bern, Switzerland
- Houriiyah Tegally, Richard J. Lessells, Tulio de Oliveira KwaZulu-Natal Research Innovation and Sequencing Platform (KRISP), Department of Laboratory Medicine & Medical Sciences, University of KwaZulu-Natal, Durban, South Africa
- Richard Neher University of Basel, Basel, Switzerland









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