

Virtual Childcare Centers: Predicting Optimal Strategies against the Spread of SARS-CoV-2 Variants

Philipp Städter^{1,2}, Paul Rudolph^{1,2}, Sandra Timme¹, Johannes Forster³, Oliver Kurzai³, Marc Thilo Figge^{1,4}

- ¹ Applied Systems Biology, Leibniz Institute for Natural Product Research and Infection Biology Hans Knöll Institute, Jena, Germany
- ² Faculty of Biological Sciences, Friedrich Schiller University Jena, Jena, Germany
- ³ Institute for Hygiene and Microbiology, University of Würzburg, Würzburg, Germany
- ⁴ Institute of Microbiology, Faculty of Biological Sciences, Friedrich Schiller University Jena, Jena, Germany

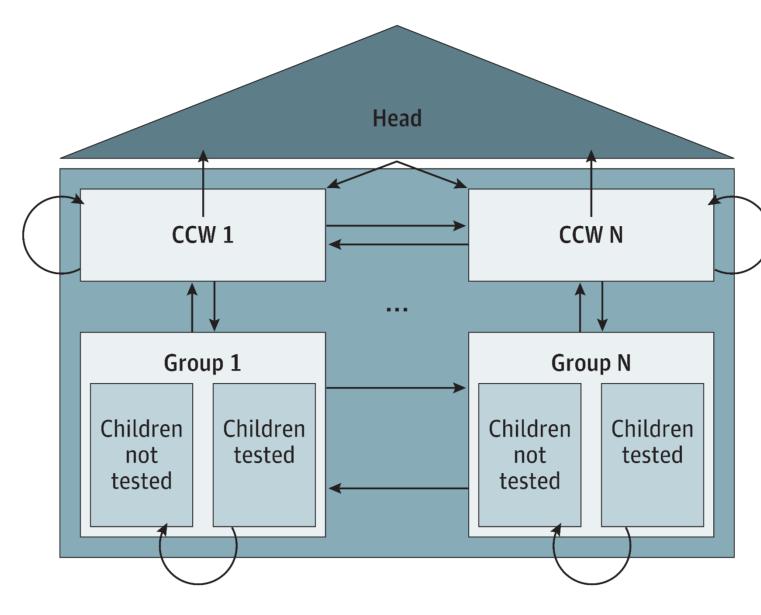
Background

- global closure of day care centers to prevent viral spread
- results include negative effects on children's well being
- viral spread through interactions inside day care centers [1]
- temporal viral load governs transmission dynamics [2]

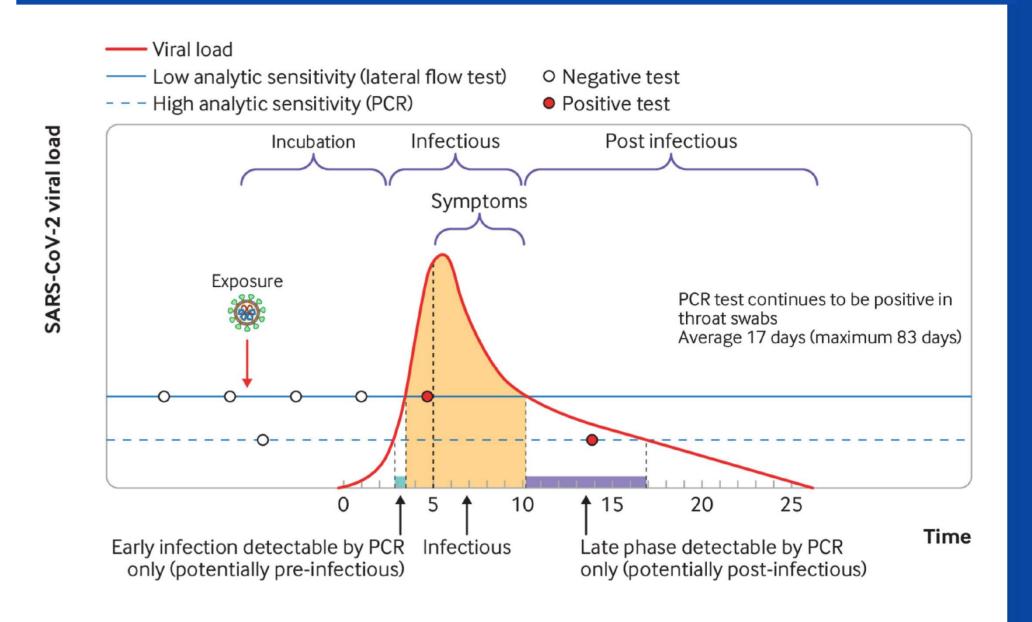
Objective:

Prediction of optimal surveillance strategies to prevent viral spread while keeping day care centers open by computational modeling

Day care center (DCC) structure

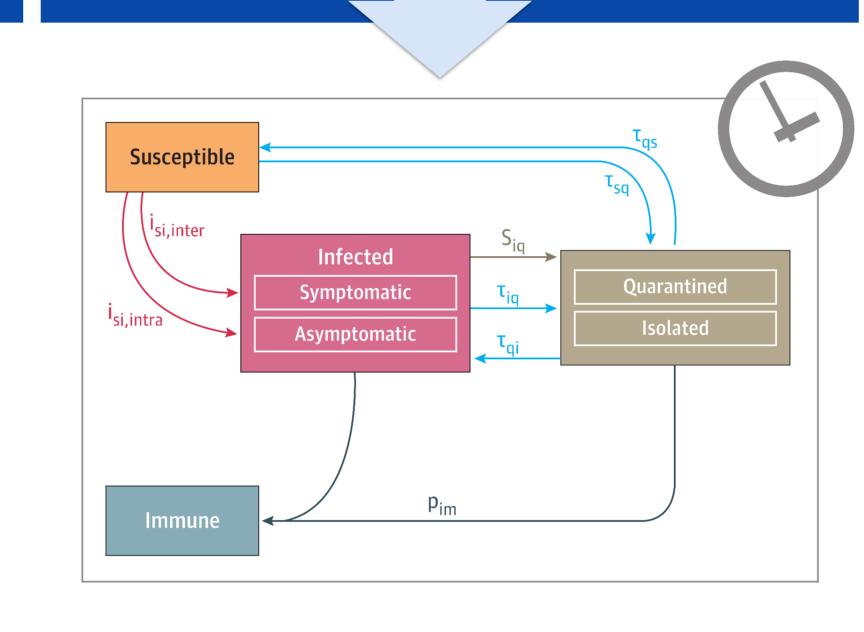


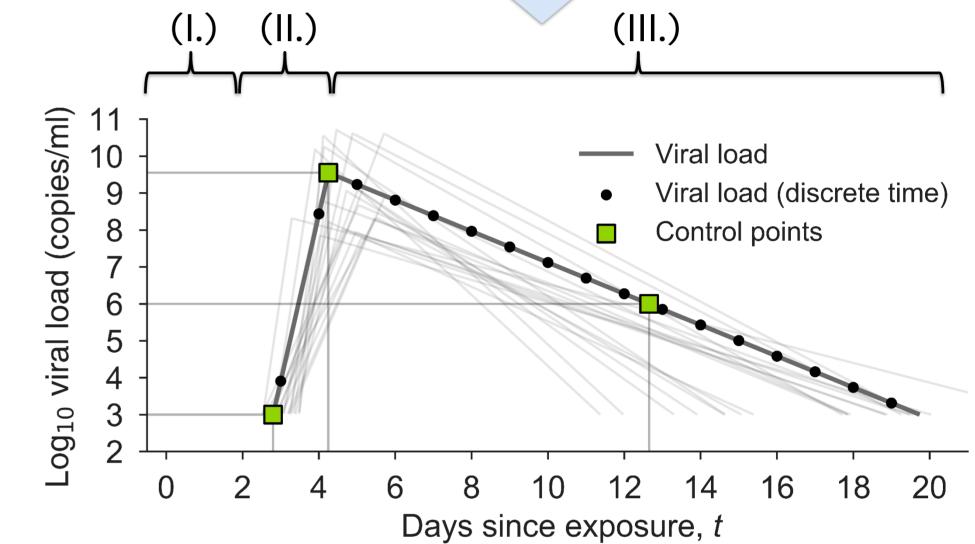
Viral load dynamics



Individual-based model

- day care centers are described by a modified **stochastic** individual-based model [3] based on an SIR model [4]
- through repeated simulations we can capture
 - random events of a small population size
 - effects of surveillance strategies
- the probability of a state transition depends on viral load
- the viral load kinetic model [5] is designed through a piecewise linear function with a
 - variable **latent** phase
 - II. rapid **growth** phase
 - III. slow **decay** phase



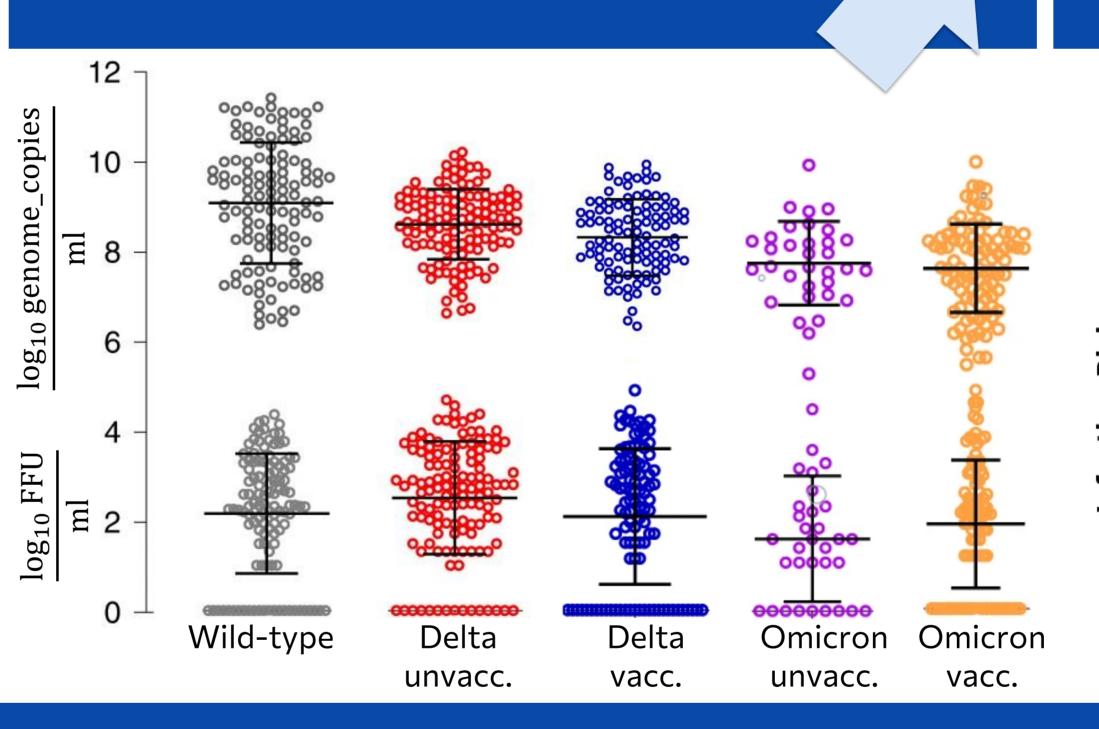


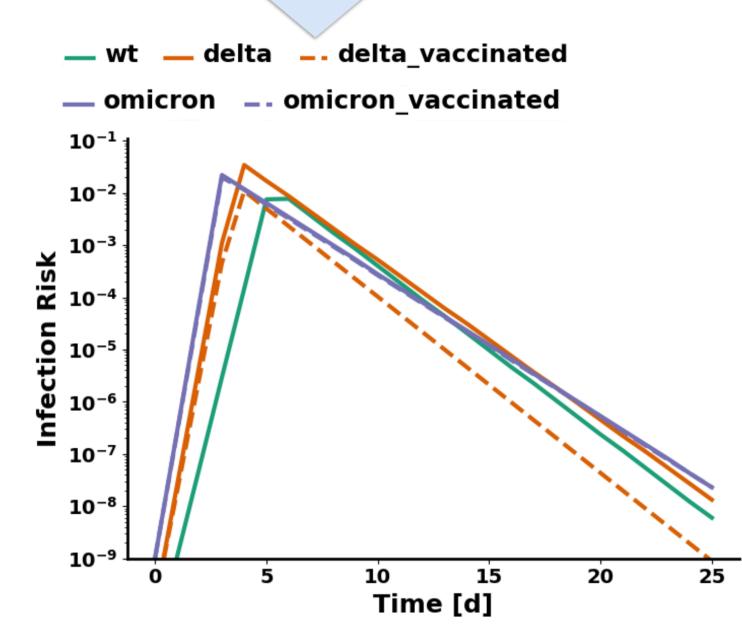
Experimental data & Infection Risk

- viral load represents the total amount of viral particles
- viral titre represents the amount of infectious viral particles
- 1) piecewise linear functions are generated by sampling from viral load posterior distributions for key parameters [6]
- 2) the infectivity is described by the D50 value that is inferred from the viral titre posterior distributions
- 3) the infection risk probability is calculated using the infection risk calculator [7]



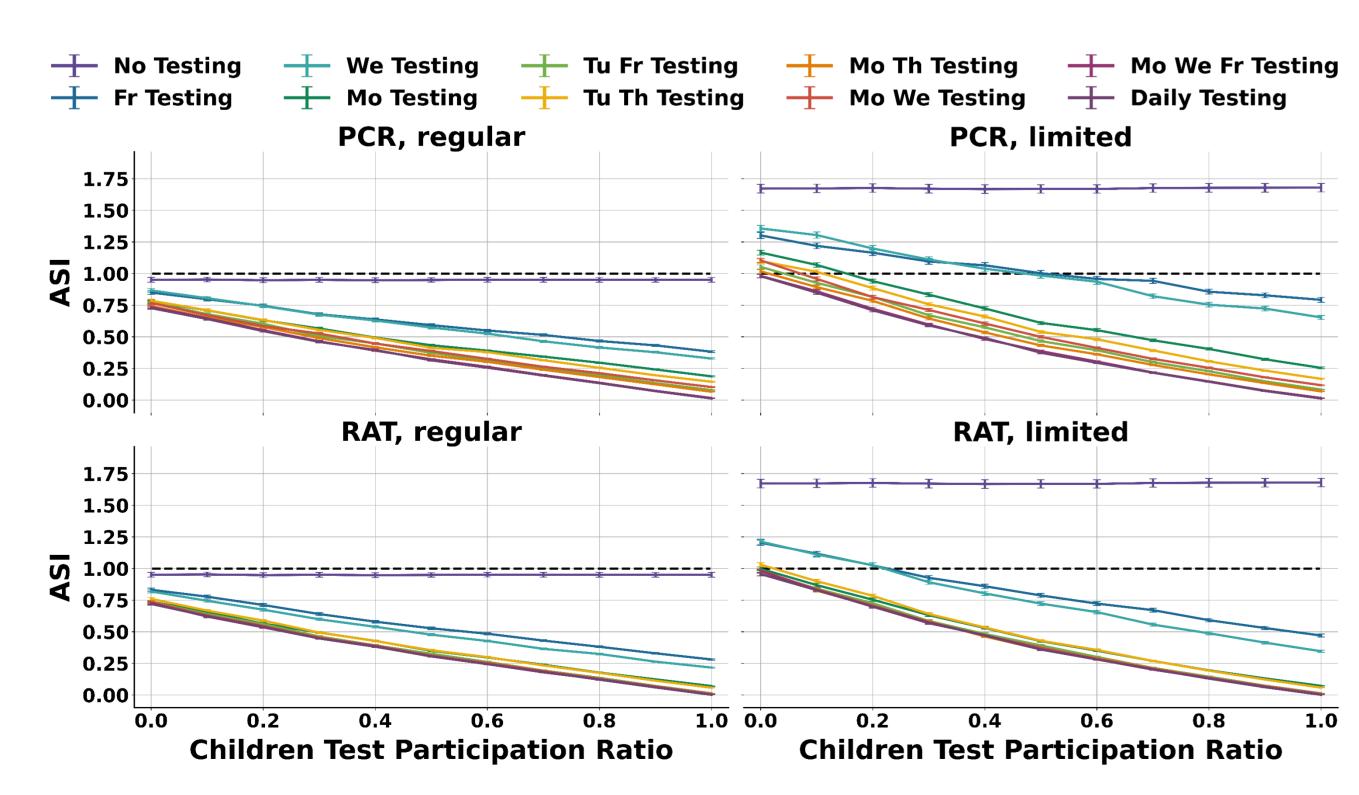
genome_copies $P_{infection_risk} = 1 - \langle 10^{\overline{D}_{50}} \rangle$



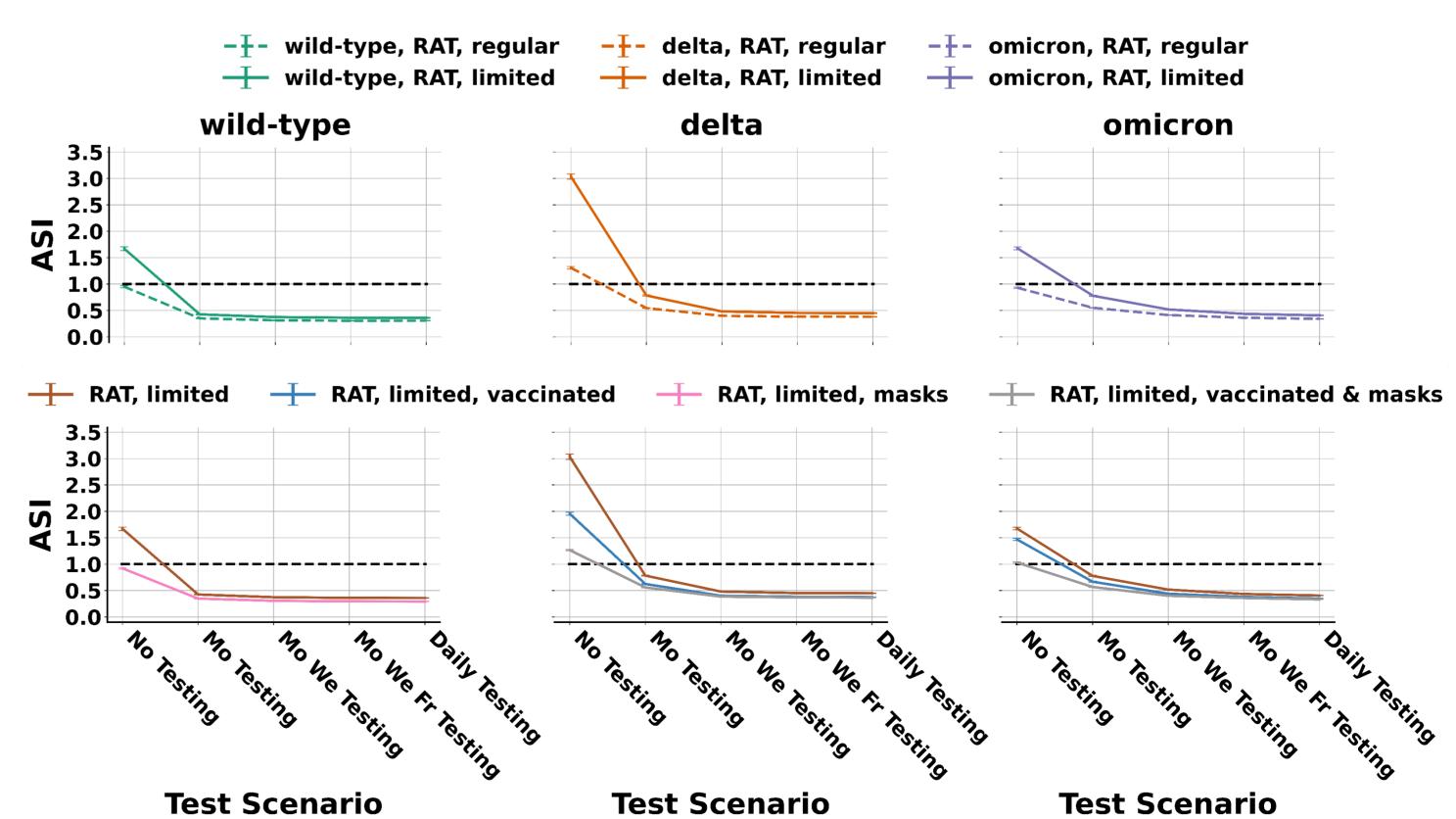


Results

- rapid antigen test outperforms PCR test due to receiving test result fast
- **limited** policy **as efficient** as regular policy with at least **1x/week** testing
- monday is the best 1x/week test scenario



- delta and omicron variants lead to an increase in average secondary infections (ASI)
- masks are the most efficient infection prevention measure



Philipp.Staedter@leibniz-hki.de

www.leibniz-hki.de

References

- ¹ Forster *et al.* 2022 *JAMA Network Open* 5(1)
- ² Crozier *et al.* 2021 *BMJ* 372
- ³ Timme et al. 2018 Front Immunol. 9 ⁴Groendyke et al. 2021 Epidemiol Met. 10(1)
- ⁵ Larremore *et al.* 2021 *Science Advances* 7(1) ⁶ Puhach et al. 2022 Nature Medicine 28(7)

⁷Lelieveld et al. 2020 Int. J. Environ. Res. Public Health 17(21)





