Selection Bias

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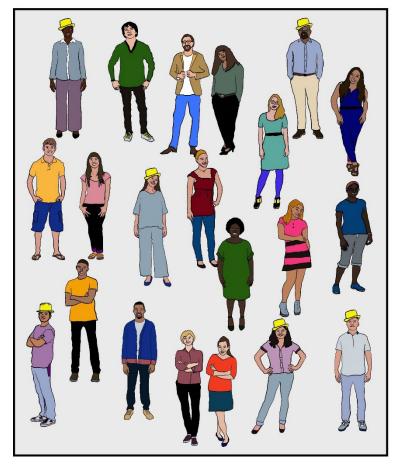


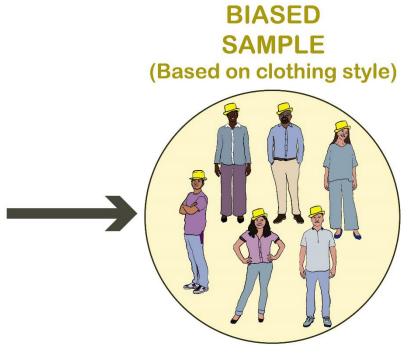


What is selection bias?

• A bias that is introduced when individuals selected into a study do not reflect the target population

POPULATION



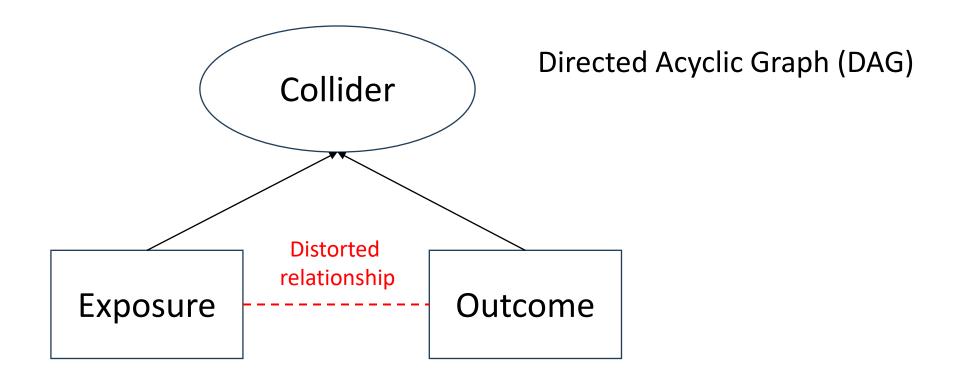


A consequence of selection bias

Can compromise internal validity

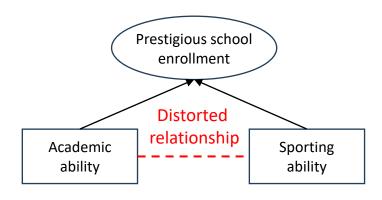
• Internal validity is where a study accurately measures the cause-and-effect relationship between an exposure and an outcome

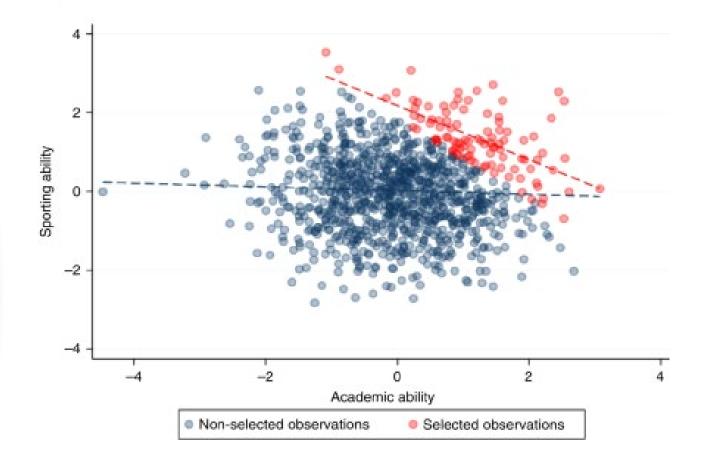
A selection bias can be a collider bias



Selecting for a collider that induces an association or distorts an association of interest is a selection bias

A simple collider bias scenario



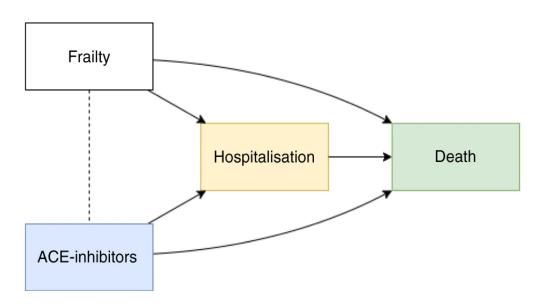


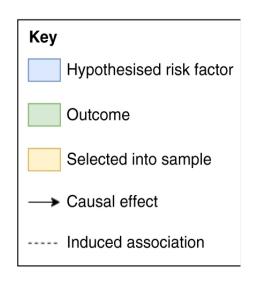
Griffith et al. 2020 (Nature Communications)



A more complicated collider bias scenario

Prognosis conditional on hospitalisation

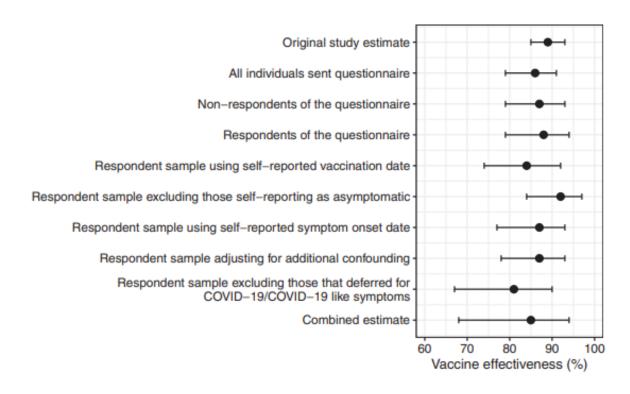




Directed Acyclic Graphs (DAGs) are useful tools to identify collider and other biases

Griffith et al. 2020 (Nature Communications)

Vaccine effectiveness (VE) study highlighting a potential for collider bias related to testing (Example #1)

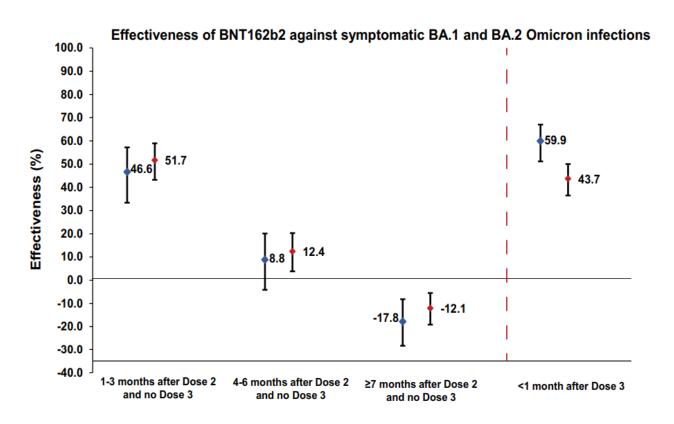


"We were unable to assess
whether collider bias was present ...
as the association between
health-seeking behaviour and testing
could not be assessed"

Adopted a test-negative design

Graham et al. 2023 (Nature Communications)

Vaccine effectiveness (VE) study highlighting a potential for collider bias related to testing (Example #2)



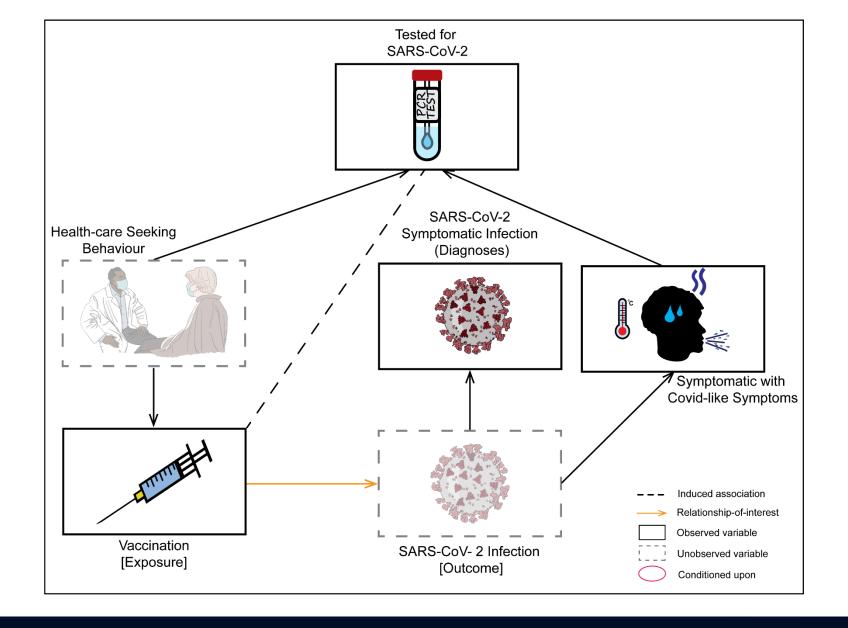
"Negative estimated effectiveness likely reflects an effect of bias

• • •

[which could arise due to]
differences in test-seeking
behaviour"

Adopted a test-negative design

Chemaitelly et al. 2022 (Nature Communications)

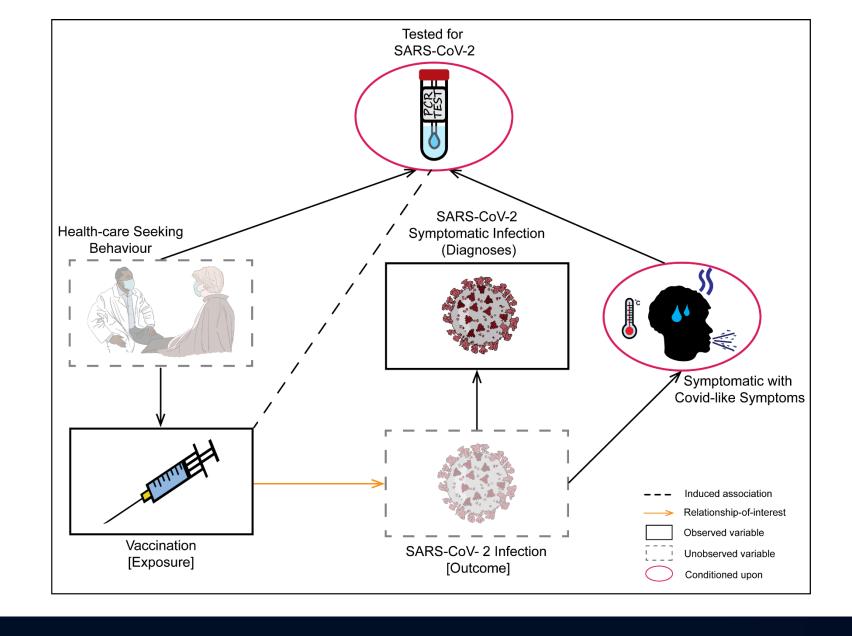


Bodner et al. (in prep)



Now using a test-negative study design to estimate symptomatic VE

Bodner et al. (in prep)





Identifying the potential for selection bias in the test-negative design (TND) is not new

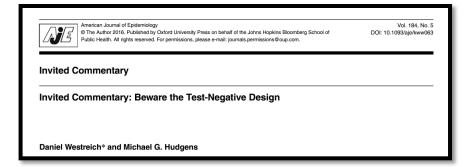
ORIGINAL ARTICLE

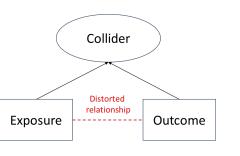
Epidemiology 2004;15: 615-625

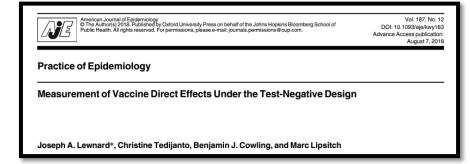
A Structural Approach to Selection Bias

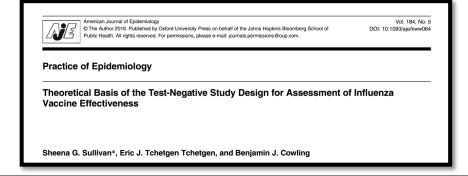
Miguel A. Hernán,* Sonia Hernández-Díaz,* and James M. Robins*





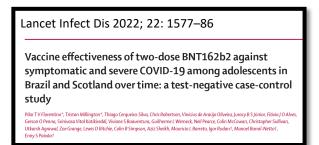


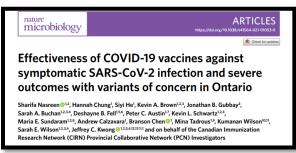




The test-negative design is commonly used when estimating VE

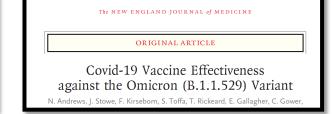


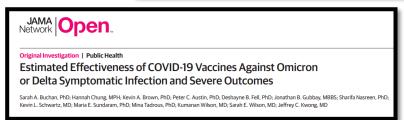






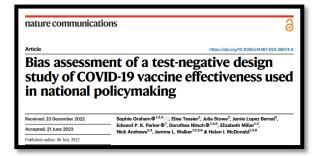






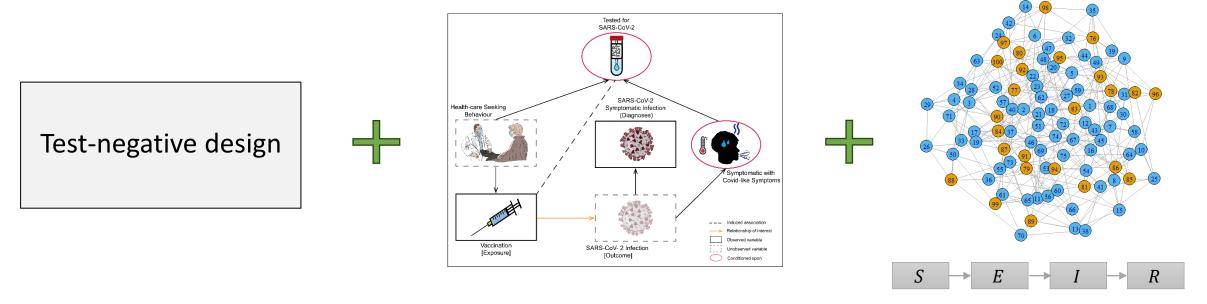






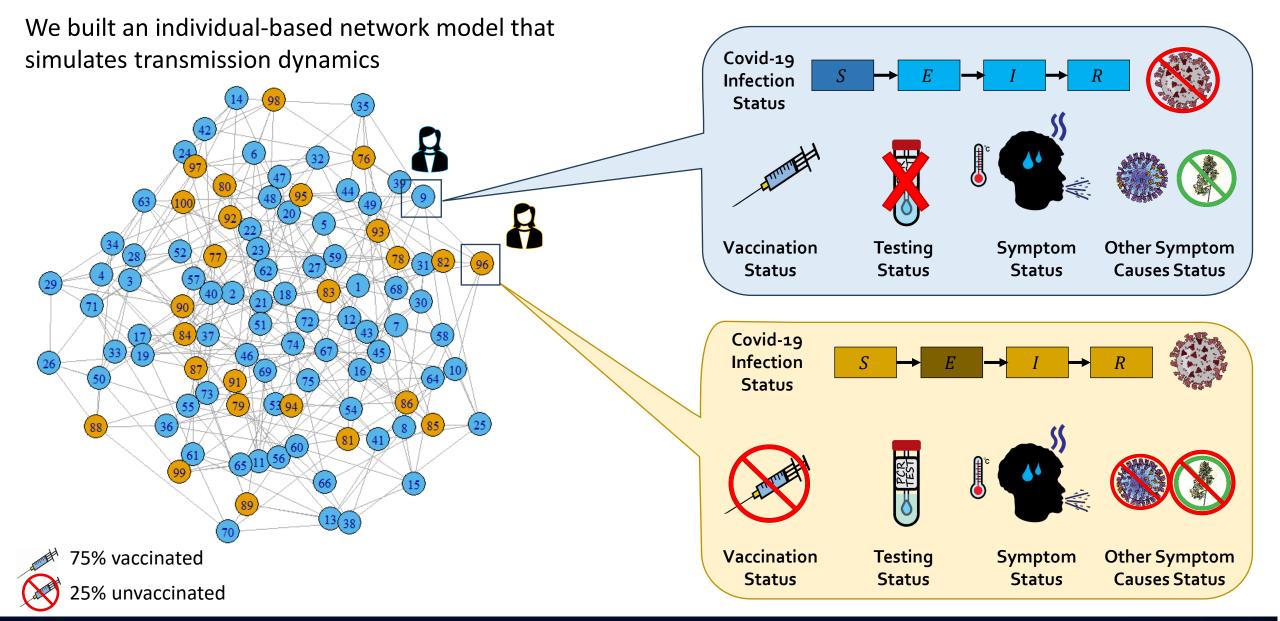


Understanding testing selection bias by combining the test-negative design, DAGs, and transmission models

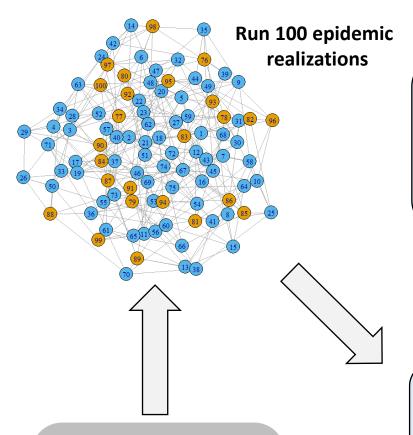


Today I will use this framework to assess:

- 1. How the presence of a testing collider influences estimates of VE
- 2. Whether other factors influence the effect of this selection bias on VE estimates







Target VE (VE)

VE = 1 - Odds Ratio[OR](t)

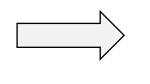
Simulated Full

Estimated VE (VE_E)

 $VE_E = 1 - Odds Ratio_E[OR_E](t)$

Test-Negative Design

Simulated Sample







Testing Scenarios

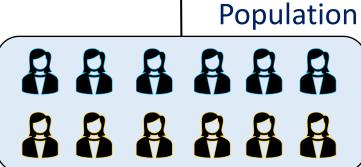
Equal Testing
Vac. 50% Higher

Vac. 100% Higher

Input

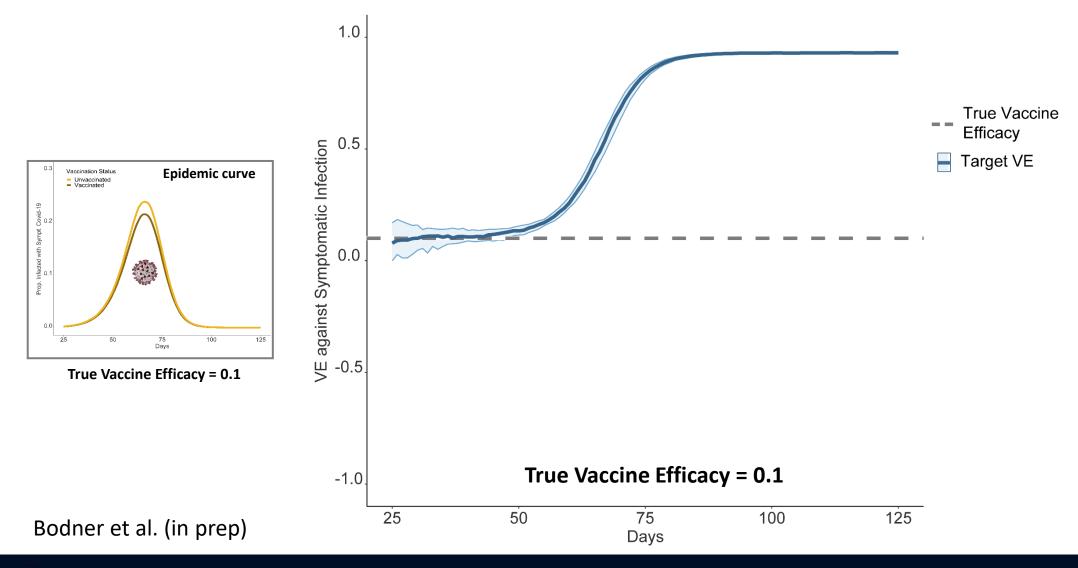
True Vaccine Efficacy

(i.e. Vaccine's true protective effect)



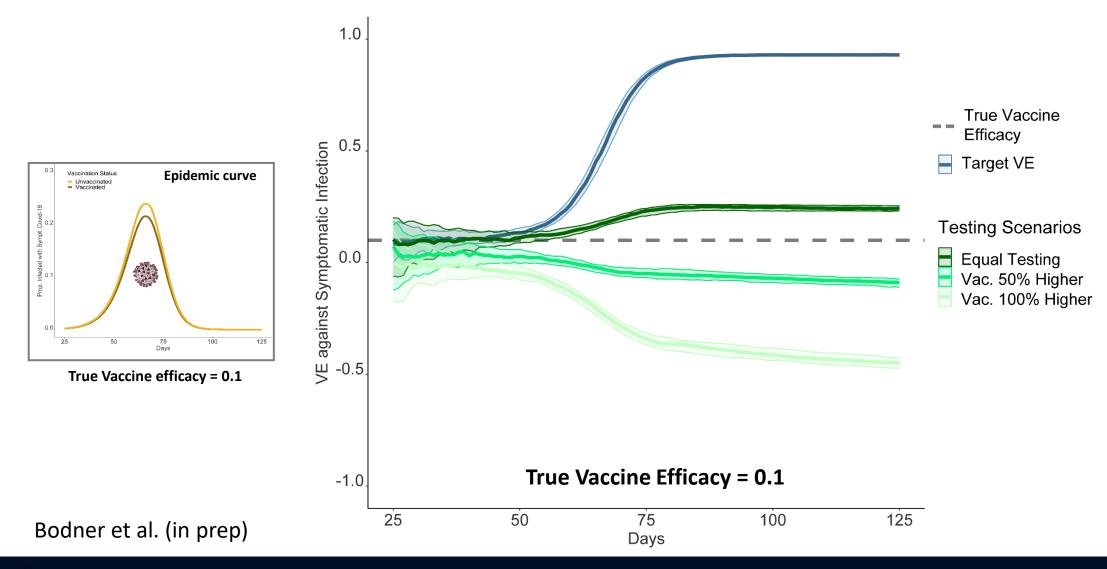


Target VE matches the True Vaccine Efficacy at the beginning of an epidemic





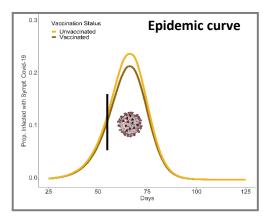
Higher testing for vaccinated causes VE to be increasingly underestimated over time



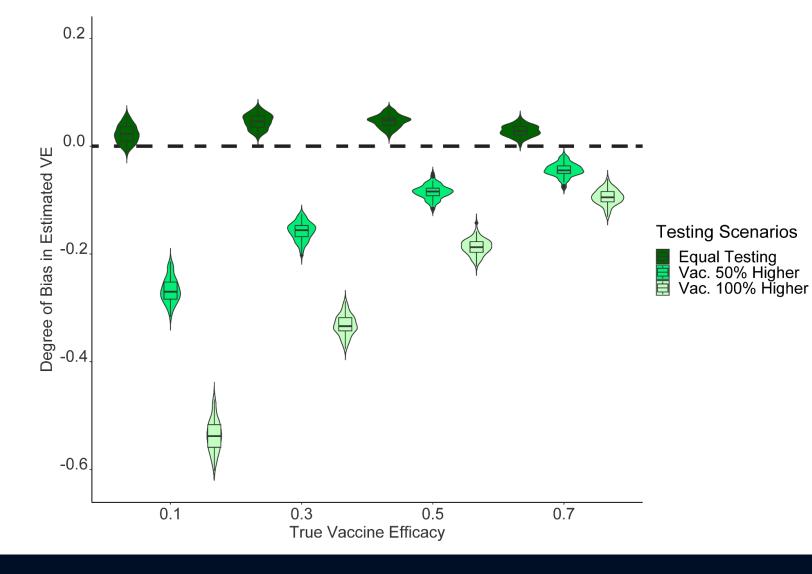


A higher true protective effect of the vaccine decreases the degree of bias

Measured at inflection point during epidemic growth



True Vaccine efficacy = 0.1



Bodner et al. (in prep)



Final thoughts

- Simulation models can help us to identify under which circumstances biases could have the greatest impact on VE estimates
 - E.g. In our simulations, we found that the amount of bias can change over time and across the true protective effect of a vaccine
- It is important to use DAGs when identifying potential selection biases when estimating VE
- Observational studies can be impacted by various types of selection bias (and colliders) – not only colliders related to testing
 - E.g. Colliders related to prior infection and symptoms

Acknowledgements



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