Using R for mathematical modelling of SARS-CoV-2

Dr Alexandra Hogan 5 May 2022



Throughout the pandemic, ongoing questions about who should be prioritised for vaccine delivery, e.g.:

- ➤ Within countries (e.g. healthcare workers, the elderly, working-age, priority groups)?
- ➤ Equitable and efficient global distribution of doses?
- ➤ How to allocate doses once program is underway?

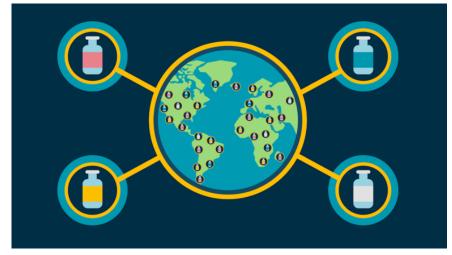
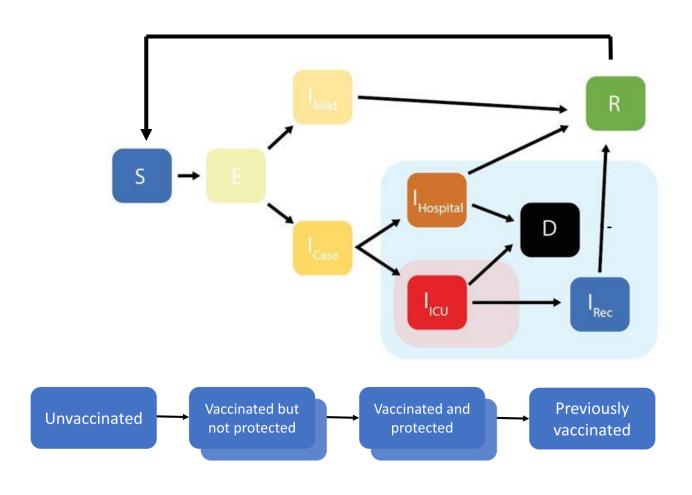


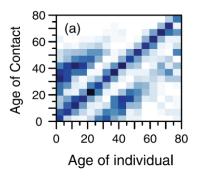
Image credit: wellcome.org

SARS-CoV-2 transmission model

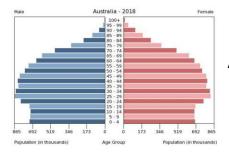
- Age-structured deterministic SEIRS model (17 five-year age groups) with expanded healthcare component
- Age-dependent disease severity
- Setting-specific healthcare capacity, contact patterns, and demography
- Vaccination incorporated by replicating compartments across vaccine states
- Vaccines:
 - modes of action (infection, disease, and transmission)
 - age-targeting and prioritisation
- Fully open source as an R package



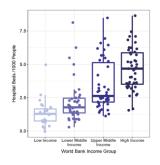
https://github.com/mrc-ide/nimue



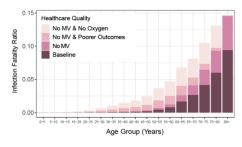
Patterns of mixing between age-groups.



Age structure (UN World Population Prospects)



Setting specific healthcare capacity –for both general hospital and ICU beds.



Age- and healthcarecapacity-dependent COVID-19 mortality

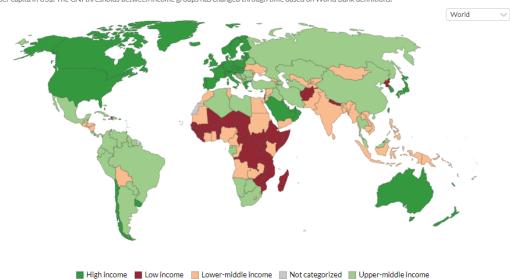


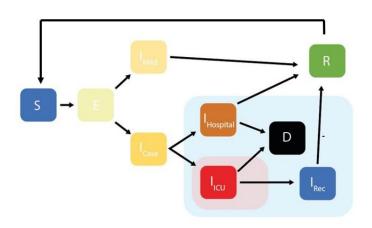
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> Our World in Data



The World Bank's income classifications split countries into one of four categories determined by the country's gross national income (GNI) per capita in US\$. The GNI thresholds between income groups has changed through time based on World Bank definitions.





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Dr Peter Winskill



Professor Azra Ghani





Imperial College London



Combating disease threats worldwide

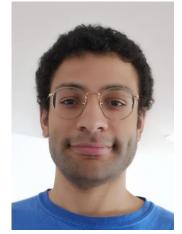




Medical Research Council







Giovanni Charles



Dr Oliver Watson

Resources

Code for today's session:

https://github.com/abhogan/covid_model_workshop

Paper on which today's modelling is based:

https://www.sciencedirect.com/science/article/pii/S0264410X21004278?via%3Dihub

Code base for the paper above:

https://github.com/mrc-ide/covid_vaccine_allocation

Documentation for the package nimue:

https://mrc-ide.github.io/nimue/

