

Reproducibility report for: Projecting contact matrices in 177 geographical regions: an update and comparison with empirical data for the COVID-19 era. **Submitted to:** PLoS Computational Biology **Manuscript number/identifier:** PCOMPBIOL-D-20-01307

Curation outcome summary: Successfully reproduced the figures presented in this manuscript.

Box 1: Criteria for repeatability and reproducibility
Model source code provided:
Source code: a standard procedural language is used (e.g. MATLAB, Python, C)
 There are details/documentation on how the source code was compiled There are details on how to run the code in the provided documentation The initial conditions are provided for each of the simulations Details for creating reported graphical results from the simulation results
Source code: a declarative language is used (e.g. SBML, CellML, NeuroML)
 The algorithms used are defined or cited in previous articles The algorithm parameters are defined Post-processing of the results are described in sufficient detail
Executable model provided:
\Box The model is executable without source (e.g. desktop application, compiled code, online service)
\Box There are sufficient details to repeat the required simulation experiments
The model is described mathematically in the article(s):
Equations representing the biological system
\Box There are tables or lists of parameter values
\Box There are tables or lists of initial conditions
\Box Machine-readable tables of parameter values
□ Machine-readable tables of initial conditions
\Box The simulation experiments using the model are described mathematically in the article:
Integration algorithms used are defined
Stochastic algorithms used are defined
Random number generator algorithms used are defined
Parameter fitting algorithms are defined
\Box The paper indicates how the algorithms yield the desired output



Box 2: Criteria for accessibility

Model/source code is available at a public repository or researcher's web site

- □ Prohibitive license provided
- □ Permissive license provided
- Open-source license provided
- $\hfill\square$ All initial conditions and parameters are provided
- □ All simulation experiments are fully defined (events listed, collection times and measurements specified, algorithms provided, simulator specified, etc.)

Box 3: Rules for Credible practice of Modeling and Simulation^a

^aModel credibility is assessed using the Interagency Modeling and Ananlysis Group conformance rubric: https://www.imagwiki.nibib.nih.gov/content/10-simple-rules-conformance-rubric

- Define context clearly: Extensive
- Use appropriate data: Extensive
- Evaluate within context: Extensive
- □ List limitations explicitly: Insufficient
- Use version control: Extensive
- Document adequately: Partial
- □ Conform to standards: Insufficient

Box 4: Evaluation

- Model and its simulations could be repeated using provided declarative or procedural code
- Model and its simulations could be reproduced



Director: Professor Herbert M. Sauro University of Washington, Seattle, WA https://reproduciblebiomodels.org

Summary comments: Model source code and data were made available for download via a Github link within the manuscript (https://github.com/kieshaprem/synthetic-contact-matrices). This was used in our attempt to reproduce the results presented in the paper. We successfully ran the necessary R scripts to reproduce the figures presented in the manuscript.

And Ragadanth

Anand K. Rampadarath¹, PhD Curator Center for Reproducible Biomedical Modeling

X 1. P.

David P. Nickerson, PhD Curation Service Director Center for Reproducible Biomedical Modeling

Auckland Bioengineering Institute, University of Auckland

¹Email: a.rampadarath@auckland.ac.nz