

**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)
<ul> <li>There are details/documentation on how the source code was compiled</li> <li>There are details on how to run the code in the provided documentation</li> <li>The initial conditions are provided for each of the simulations</li> <li>Details for creating reported graphical results from the simulation results</li> </ul>
Source code: a declarative language is used (e.g. SBML, CellML, NeuroML)
<ul> <li>The algorithms used are defined or cited in previous articles</li> <li>The algorithm parameters are defined</li> <li>Post-processing of the results are described in sufficient detail</li> </ul>
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<sup>a</sup>

<sup>a</sup>Model 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.

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