



Reproducibility report for: Reproducibility study of the Fabbri et al. 2017 model of the human sinus node action

potential.

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Curation outcome summary: Successfully reproduced all 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:
$\hfill\Box$ The model is executable without source (e.g. desktop application, compiled code, online service)
$\hfill\Box$ There are sufficient details to repeat the required simulation experiments
■ The model is described mathematically in the article(s):
Equations representing the biological system
■ There are tables or lists of parameter values
☐ There are tables or lists of initial conditions
Machine-readable tables of parameter values
Machine-readable tables of initial conditions
■ 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
■ The paper indicates how the algorithms yield the desired output

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## 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
- 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: Extensive
- 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

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

**Summary comments:** Model and source code are available in the associated OMEX archive. This was used in our attempt to reproduce the results presented in the paper. We successfully ran the python scripts provided to reproduce Figure 1 - Figure 7 as presented in this manuscript.

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