Author Contributions Checklist Form

This form documents the artifacts associated with the article (i.e., the data and code supporting the computational findings) and describes how to reproduce the findings.

# Part 1: Data

This paper **does not** involve analysis of external data (i.e., no data are used or the only data are generated by the authors via simulation in their code).

I certify that the author(s) of the manuscript have legitimate access to and permission to use the data used in this manuscript.

## Abstract

An earthquake data set for the period between 1885 and 1980 for a region close to Japan is used in Section 7.1; this data is taken from the cited paper from Ogata.

An Rocky Mountain Spotted Fever data set for California and Florida, raning from 1960 to 2011 is used; this data is also ued by the cited Schoenberg paper. The data is available from Project Tycho (open source).

## Availability

Data **are** publicly available

Data **cannot be made** publicly available

If the data are publicly available, see the *Publicly available data* section. Otherwise, see the *Non-publicly available dat*a section, below.

### Publicly available data

Data are available online at:

Data are available as part of the paper’s supplementary material.

Data are publicly available by request, following the process described here:

Simply make a free account on <https://www.tycho.pitt.edu/accounts/login/>. After this, the data can be downloaded.

Data are or will be made available through some other mechanism, described here:

### Non-publicly available data

Discussion of lack of publicly available data:

## Description

### File format(s)

CSV or other plain text:

Software-specific binary format (.Rda, Python pickle, etc.):

Standardized binary format (e.g., netCDF, HDF5, etc.):

Other (described here):

.xlsx

### Data dictionary

Provided by the authors in the following file(s):

Data file(s) is (are) self-describiing (e.g., netCDF files)

Available at the following URL:

### Additional information (optional)

Data is described in Statistical models for earthquake occurrences and residual analysis for point processes, Ogata (1988) and A recursive point process model for infectious diseases, Schoenberg (2019). Apart from that, it is selfexplanatory, and instructions are given in TestingOgata.m and testingSchoenberg.m

# Part 2: Code

## Abstract

For simulations, we simulate sample paths, and using the simulated paths we perform Algorithm 1 from our paper. We do this for a large number of sample paths to find rejection rates and evaluate performance of our method under H0 and H1.

For the testing procedures, we import the data, preprocess it. For Ogata, we take their estimates, and compare tests based on random time change and our tests. For Schoenberg, we estimate the parameters and compare our tests to random time change based tests.

## Description

### Code format(s)

Script files

R  Python  Matlab

Other:

Package

R  Python  MATLAB toolbox

Other:

Reproducible report

R Markdown  Jupyter notebook

Other:

Shell script

Other (described here):

### **Supporting software requirements**

Version of primary software used

Matlab R2023a

Libraries and dependencies used by the code

cmtest.m, Cramer-von Mises test

[Version 1.1.0.0](https://nl.mathworks.com/matlabcentral/fileexchange/50157-cramer-von-mises-test" \l "version_history_tab) (17.5 KB) by [Ahmed BenSaïda](https://nl.mathworks.com/matlabcentral/profile/authors/135854)

### Supporting system/hardware requirements (optional)

### Parallelization used

No parallel code used

Multi-core parallelization on a single machine/node

Number of cores used: 8

Multi-machine/multi-node parallelization

Number of nodes and cores used:

### License

MIT License (default)

BSD

GPL v3.0

Creative Commons

Other (described here):

### Additional information (optional)

# Part 3: Reproducibility workflow

## Scope

The provided workflow reproduces:

Any numbers provided in text in the paper

The computational method(s) presented in the paper (i.e., code is provided that implements the method(s))

All tables and figures in the paper

Selected tables and figures in the paper, as explained and justified here:

Table1seeded.m can be run directly to reproduce the results from Table1 exactly. AAA\_Replication\_instructions.txt outlines how the remaining tables and figures can be reproduced; which files are needed and what needs to plugged in.

## Workflow details

### Location

The workflow is available:

As part of the paper’s supplementary material

In this Git repository:

Other:

### Format(s)

Single master code file

Wrapper (shell) script(s)

Self-contained R Markdown file, Jupyter notebook, or other literate programming approach

Text file (e.g., a readme-style file) that documents workflow

Makefile

Other (more detail in 'Instructions' below)

### Instructions

AAA\_Replication\_instructions.txt

Expected run-time

Approximate time needed to reproduce the analyses on a standard desktop machine:

<1 minute

1-10 minutes

10-60 minutes

1-8 hours

>8 hours

Not feasible to run on a desktop machine, as described here:

Table 1, 4 6 will take <30 minutes each, 2, 3 and 5 will take at most 3 days each. Results of Section 7 can be retrieved by less than 10 minutes computation time.

### Additional documentation (optional)

# Notes (optional)