Author Contributions Checklist Form

The purpose of the Author Contributions Checklist (ACC) Form is to document the artifacts associated with a manuscript (i.e., the data and code supporting the computational findings) and describe how to reproduce the findings. The final version of this document is included as online supplemental material with the published paper and on the JASA Github site (<http://github.com/JASA-ACS/>) and is referenced in the abstract.

# 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

## Availability

Data **are** publicly available

Data **cannot be made** publicly available

### 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:

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

### Non-publicly available data

Discussion of lack of publicly available data:

## Description

### File format(s) (check all that apply):

CSV or other plain text:

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

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

Other (please specify below)

# Part 2: Code

## Description

### Code format(s) (check all that apply)

Script files

R  Python  Matlab

Other:

Package

R  Python  MATLAB toolbox

Other:

Reproducible report

R Markdown  Jupyter notebook

Other:

Shell script

Other (please specify below)

### **Supporting software requirements**

Version of primary software used (e.g., R version 3.6.0):

Libraries and dependencies used by the code, including version numbers

### Supporting hardware requirements (optional)

### Parallelization used

No parallel code used

Multi-core parallelization on a single machine/node

Number of cores used:

Multi-machine/multi-node parallelization

Number of nodes and cores used:

### License

**\*\*\* Click to select a license type \*\*\***

### Additional information (optional)

Code is also available online at:

Unique identifier / DOI / version information:

# Part 3: Workflow

## Scope

The provided workflow reproduces:

**\*\*\* Click to select \*\*\***

## Workflow

The materials provided should provide a straightforward way for reviewers and readers to reproduce analyses with as few steps as possible.

### Format(s) (check all that apply):

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

### Expected run-time

**\*\*\* Click to select duration \*\*\***

### Additional documentation (optional)

# Notes (optional)