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
Part	1:	ыага

- ☐ 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

The OhioT1DM data set(2018) contains glucose monitoring, insulin, physiological sensor, and self-reported life-event data for six patents with type 1 diabetes over eight weeks.

### Availability

- $\boxtimes$  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 data* section, below.

#### Publicly available data

- ☐ Data are available online at:http://smarthealth.cs.ohio.edu/OhioT1DM-dataset.html
- ☐ 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 here:

#### Non-publicly available data

#### Description

#### File format(s)

- $\boxtimes$  CSV or other plain text.
- □ Software-specific binary format (.Rda, Python pickle, etc.): pkcle
- □ Standardized binary format (e.g., netCDF, HDF5, etc.):
- $\square$  Other (please specify):

## Data dictionary

- ☑ Provided by authors in the following file(s):./real\_data/code\_data/code\_data/Data\_Ohio.csv in VEPO repo.
- □ Data file(s) is(are) self-describing (e.g., netCDF files)
- ✓ Available at the following URL: http://smarthealth.cs.ohio.edu/OhioT1DM-dataset.html

## Additional Information (optional)

# Part 2: Code

## Abstract

The code for VEPO(URL:https://anonymous.4open.science/r/VEPO-B93E) calculations is consist of two parts, experiments in Section 5.1 of text and Supplement D.1 can be found in fold toy , while experiments in Section 5.2 and Supplement D.2 can be found in fold real\_data . The source code of manuscript and ACC form is attached in this repo for the convenience of reviewing.

Description	
$\operatorname{Code} \ \operatorname{format}(\operatorname{s})$	
<ul> <li>□ Script files</li> <li>□ R</li> <li>⋈ Python</li> <li>□ Matlab</li> <li>□ Other:</li> </ul>	
<ul> <li>□ Package</li> <li>⋈ R</li> <li>⋈ Python</li> <li>□ MATLAB toolbox</li> <li>□ Other:</li> </ul>	
<ul> <li>□ Reproducible report</li> <li>□ R Markdown</li> <li>□ Jupyter notebook</li> <li>□ Other:</li> </ul>	
☐ Shell script ☐ Other (please specify):	
Supporting software requirements	
Version of primary software used Python 3.6.9	
Libraries and dependencies used by the code README.md in our repository for detail.	Seeing requirements.txt or "Installation" section i
Supporting system/hardware requirements (opt	tional)
Parallelization used	
<ul> <li>□ No parallel code used</li> <li>⋈ Multi-core parallelization on a single machine/no</li> <li>− Number of cores used: 10</li> <li>□ Multi-machine/multi-node parallelization</li> <li>− Number of nodes and cores used:</li> </ul>	ode
License	
<ul> <li></li></ul>	

# Additional information (optional)

# Part 3: Reproducibility workflow

# Scope

The provided workflow reproduces:
<ul> <li>□ Any numbers provided in text in the paper</li> <li>□ The computational method(s) presented in the paper (i.e., code is provided that implements the method(s))</li> <li>☑ All tables and figures in the paper</li> <li>□ Selected tables and figures in the paper, as explained and justified below:</li> </ul>
Workflow
Location
The workflow is available:
<ul> <li>□ As part of the paper's supplementary material.</li> <li>□ In this Git repository:</li> <li>□ Other (please specify):</li> </ul>
Format(s)
<ul> <li>□ Single master code file</li> <li>□ Wrapper (shell) script(s)</li> <li>□ Self-contained R Markdown file, Jupyter notebook, or other literate programming approach</li> <li>□ Text file (e.g., a readme-style file) that documents workflow</li> <li>□ Makefile</li> <li>□ Other (more detail in <i>Instructions</i> below)</li> </ul>
Instructions
The detail of how to use code and reproduce our results can be found in README.md in our VEPC repository.
Expected run-time
Approximate time needed to reproduce the analyses on a standard desktop machine:
<ul> <li>□ &lt; 1 minute</li> <li>□ 1-10 minutes</li> <li>□ 10-60 minutes</li> <li>□ 1-8 hours</li> <li>□ &gt; 8 hours</li> <li>□ Not feasible to run on a desktop machine, as described here:</li> </ul>
Additional information (optional)
Notes (optional)