

Computation of Physiologic Closed-Loop Controller Performance Metrics

Tool Reference

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User Manual

Overview

This software package provides tools to compute physiologic closed-loop controller (PCLC) performance metrics from time-series data of a controlled physiologic variable. The package is available in both MATLAB and Python.

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MATLAB package files

- **evaluate_CL_metrics.m**: the PCLC performance metric computation function
- **test_data.mat**: an example dataset for demonstration purposes
- **run_example.m**: an example script demonstrating how to load data, call the function, and visualize results

Python package files

- **evaluate_CL_metrics.py**: the PCLC performance metric computation function
- **test_data.mat**: the same example dataset, loaded via `scipy.io.loadmat`
- **run_example.py**: an example script demonstrating how to load data, call the function, and visualize results

NOTE: *The Python implementation is an exact translation of the MATLAB function. All equations, thresholds, and logic are identical. `ind_CL_start` and `ind_CL_end` are passed as 1-based integers in both implementations, matching MATLAB convention.*

Data Requirements

Before calling the function, format input data as follows. While variable names below use MAP-specific terminology consistent with the provided example dataset, the function can be applied to other physiologic variables. Users applying the tool to other applications can substitute the appropriate controlled variable and disturbance signal in place of `MAP_Values` and `Hemorrhage_Values`, respectively.

- **MAP_Values** — full time-series of the controlled physiologic variable (e.g., MAP in mmHg)
- **MAP_Times** — corresponding time vector in minutes, same length as `MAP_Values`
- **Hemorrhage_Values** — full time-series of the disturbance signal (e.g., hemorrhage rate in l/min); set to a zero vector if no disturbance was applied
- **ind_CL_start** — scalar integer index (1-based) at which controller engagement begins
- **ind_CL_end** — scalar integer index (1-based) at which controller engagement ends
- **target** — scalar target value of the controlled variable (e.g., 70 for MAP in mmHg)

The function accepts the data as provided and does not apply any additional processing to the controlled variable signal. Data acquisition and preprocessing can be considered by the user prior to calling the function.

NOTE: *ind_CL_start* and *ind_CL_end* use 1-based indexing in both MATLAB and Python, where index 1 corresponds to the first element of the vector. In Python, the function converts these to 0-based indices internally; the user does not need to subtract 1.

Step-by-Step Instructions — MATLAB

1. **Step 1:** Place `evaluate_CL_metrics.m`, `test_data.mat`, and `run_example.m` in the same MATLAB working directory.
2. **Step 2:** Open MATLAB® R2023b or a compatible version.
3. **Step 3:** Run the example script `run_example.m` to verify correct installation and review the expected outputs using the provided example dataset.
4. **Step 4:** Review the output struct `PCLC_char` for the computed metric values. Access fields using dot notation, for example `PCLC_char.MDPE`. NaN values indicate metrics that could not be computed for the given subject.
5. **Step 5:** When reporting results across a cohort of multiple subjects, call `evaluate_CL_metrics.m` separately for each subject.

Step-by-Step Instructions — Python

Installation

Python 3.8 or later is required. Install the required packages by running the following command in a command line window:

```
pip3 install numpy pandas scipy matplotlib
```

1. **Step 1:** Place `evaluate_CL_metrics.py`, `test_data.mat`, and `run_example.py` in the same folder.
2. **Step 2:** Open a command line window and navigate to the folder containing the files. For example:

```
cd "/path/to/your/folder"
```
3. **Step 3:** Run the example script to verify correct installation and review the expected outputs:

```
python3 run_example.py
```
4. **Step 4:** Review the output dictionary `PCLC_char` for the computed metric values. Access fields using bracket notation, for example `PCLC_char['MDPE']`. NaN values (`float('nan')`) indicate metrics that could not be computed for the given subject. To check for NaN in Python use `import math; math.isnan(PCLC_char['risetime'])`.
5. **Step 5:** When reporting results across a cohort of multiple subjects, call `evaluate_CL_metrics` separately for each subject.

NOTE: If your .mat file was saved in MATLAB version 7.3 format (HDF5), `scipy.io.loadmat` will raise a `NotImplementedError`. In that case, use `h5py`: `import h5py; f = h5py.File('my_data.mat', 'r')`.

Output

MATLAB

The function returns a MATLAB struct named `PCLC_char`. Fields are accessed using dot notation: `PCLC_char.MDPE`, `PCLC_char.risetime`, etc.

Python

The function returns a Python dictionary named `PCLC_char`. Fields are accessed using bracket notation: `PCLC_char['MDPE']`, `PCLC_char['risetime']`, etc. All values are Python floats. NaN is represented as `float('nan')`.

Output Metrics Table

The following metrics are returned in both implementations [1,2]. Units shown reflect the provided example dataset (MAP-based automated fluid resuscitation).

Metric	Symbol	Units	MATLAB field	Python key
Median Performance Error	MDPE	%	<code>PCLC_char.MDPE</code>	<code>PCLC_char['MDPE']</code>
Median Absolute Performance Error	MDAPE	%	<code>PCLC_char.MDAPE</code>	<code>PCLC_char['MDAPE']</code>
Wobble	Wob	%	<code>PCLC_char.Wobble</code>	<code>PCLC_char['Wobble']</code>
Percentage of Time Within 10% of Target	POT(10)	%	<code>PCLC_char.per10</code>	<code>PCLC_char['per10']</code>
Percentage of Time Within 20% of Target	POT(20)	%	<code>PCLC_char.per20</code>	<code>PCLC_char['per20']</code>
Percentage of Time Within 30% of Target	POT(30)	%	<code>PCLC_char.per30</code>	<code>PCLC_char['per30']</code>
Modified Rise Time	Tr	min	<code>PCLC_char.risetime</code>	<code>PCLC_char['risetime']</code>
Percentage Overshoot	%O.S.	%	<code>PCLC_char.overshoot</code>	<code>PCLC_char['overshoot']</code>
Settling Time	Ts	min	<code>PCLC_char.settlingtime</code>	<code>PCLC_char['settlingtime']</code>
Divergence	D	%/min	<code>PCLC_char.divergence</code>	<code>PCLC_char['divergence']</code>
Steady State Error	SSE	mmHg	<code>PCLC_char.SSE</code>	<code>PCLC_char['SSE']</code>

NaN is returned for `risetime`, `overshoot`, and `settlingtime` when the conditions for their computation are not met. When applying statistical analyses and reporting summary statistics across a cohort, methods for handling NaN values can be considered.

These metrics are designed for PCLC evaluation scenarios in which the controller may be subject to disturbances during the engagement period. In the representative use case of automated fluid resuscitation, these disturbances take the form of hemorrhagic events that perturb MAP during closed-loop controller engagement. Several metrics, including settling time and steady state error, include specific provisions to account for the presence of such disturbances. Users applying this function to other PCLC applications can supply the appropriate disturbance signal for their specific context, or a zero vector if no disturbance is applied.

Example Results

The following example results are generated by running `run_example.m` (MATLAB) or `run_example.py` (Python) with the provided example dataset (`test_data.mat`). The example dataset represents a MAP-based automated fluid resuscitation scenario in which a closed-loop controller regulates MAP toward a target of 70 mmHg in an anesthetized swine subject undergoing hemorrhage and fluid resuscitation, consistent with the experimental protocol described in [1]. This example is provided for demonstration and test purposes only. It illustrates the expected format and range of outputs and does not constitute a performance benchmark or acceptance criterion.

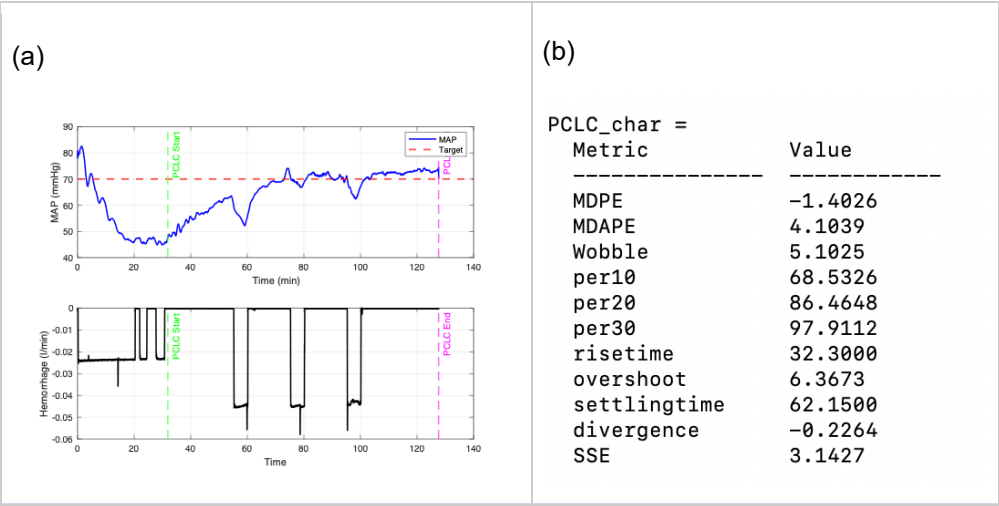


Figure 1: (a) MAP response and hemorrhage profile generated by running `run_example.m` or `run_example.py` with the provided example dataset. The top panel shows MAP over time with the target value, PCLC start, and PCLC end indicated. The bottom panel shows the hemorrhage disturbance profile with the same engagement markers; (b) PCLC performance metrics for the example.

For applications other than MAP-based fluid resuscitation, the example results will differ based on the controlled variable, target value, disturbance profile, and controller behavior specific to the application.

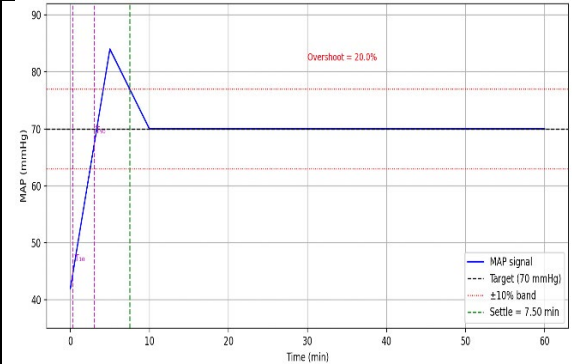
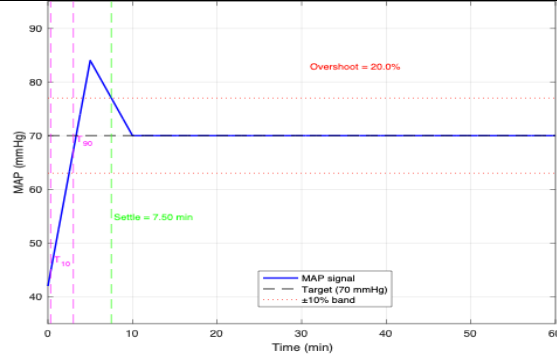
Software Code Testing Examples

Manufactured data with known PCLC performance metrics are used for code testing. The tables below show results from both MATLAB and Python implementations across three manufactured signal examples, confirming equivalence between implementations.

MATLAB

Python

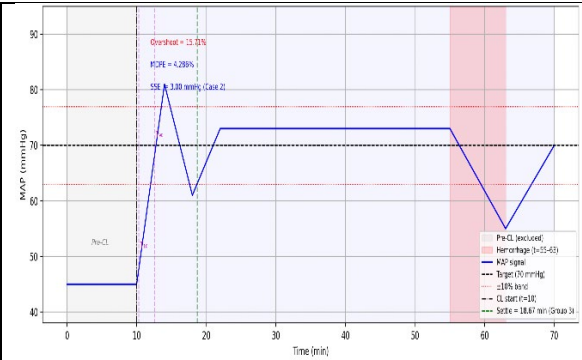
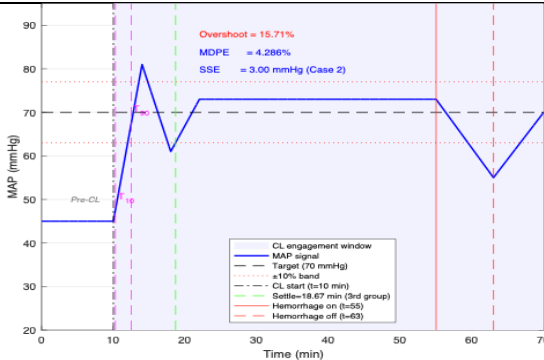
Example 1



Metric	Expected	Computed	Status
MDPE (%)	0.0000	0.0000	PASS
MDAPE (%)	0.0000	0.0000	PASS
Wobble (%)	0.0000	0.0000	PASS
per10 (%)	90.2516	90.2516	PASS
per20 (%)	97.2005	97.2005	PASS
per30 (%)	98.6002	98.6002	PASS
risetime (min)	2.6700	2.6700	PASS
overshoot (%)	20.0000	20.0000	PASS
settlingtime (min)	7.5000	7.5000	PASS
divergence (%/min)	< 0	-0.1957	PASS (sign < 0)
SSE (mmHg)	0.0000	0.0000	PASS

Metric	Expected	Computed	Status
MDPE (%)	0.0000	0.0000	PASS
MDAPE (%)	0.0000	0.0000	PASS
Wobble (%)	0.0000	0.0000	PASS
per10 (%)	90.2516	90.2516	PASS
per20 (%)	97.2005	97.2005	PASS
per30 (%)	98.6002	98.6002	PASS
risetime (min)	2.6700	2.6700	PASS
overshoot (%)	20.0000	20.0000	PASS
settlingtime (min)	7.5000	7.5000	PASS
divergence (%/min)	< 0	-0.1957	PASS (sign < 0)
SSE (mmHg)	0.0000	0.0000	PASS

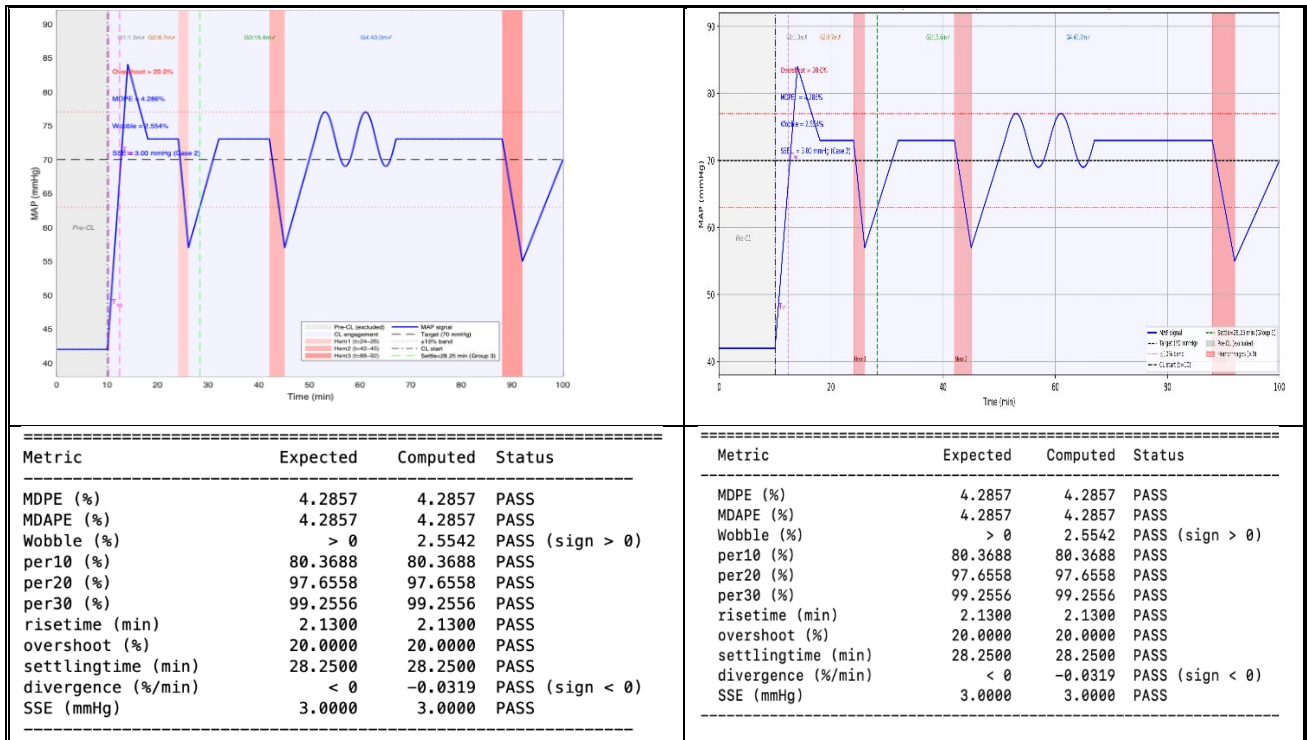
Example 2



Metric	Expected	Computed	Status
MDPE (%)	4.2857	4.2857	PASS
MDAPE (%)	4.2857	4.2857	PASS
Wobble (%)	0.0000	0.0000	PASS
per10 (%)	80.6366	80.6366	PASS
per20 (%)	96.4339	96.4339	PASS
per30 (%)	99.2501	99.2501	PASS
risetime (min)	2.2200	2.2200	PASS
overshoot (%)	15.7143	15.7143	PASS
settlingtime (min)	18.6700	18.6700	PASS
divergence (%/min)	> 0	0.0207	PASS (sign > 0)
SSE (mmHg)	3.0000	3.0000	PASS

Metric	Expected	Computed	Status
MDPE (%)	4.2857	4.2857	PASS
MDAPE (%)	4.2857	4.2857	PASS
Wobble (%)	0.0000	0.0000	PASS
per10 (%)	80.6366	80.6366	PASS
per20 (%)	96.4339	96.4339	PASS
per30 (%)	99.2501	99.2501	PASS
risetime (min)	2.2200	2.2200	PASS
overshoot (%)	15.7143	15.7143	PASS
settlingtime (min)	18.6700	18.6700	PASS
divergence (%/min)	> 0	0.0207	PASS (sign > 0)
SSE (mmHg)	3.0000	3.0000	PASS

Example 3



References

- [1] Chalumuri et al. *Annals of Biomedical Engineering*, 54, pp. 857–868 (2026).
<https://doi.org/10.1007/s10439-025-03929-2>
- [2] Varvel et al. *Journal of Pharmacokinetics and Biopharmaceutics*, 20(1), 63–94 (1992).
<https://doi.org/10.1007/bf01143186>