

0.a. Goal

[illegible]

0.b. Target

[illegible]

0.c. Indicator

[illegible]

0.e. Metadata update

□□□□□□□□□□ □□□□ [□□]

1.a. Organisation

□□□□□□□□□□□□□□□□□□ (NIS) / □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□ [□□□]

1.b. Contact person(s)

សម្តេច សុវណ្ណឌី / **សម្តេច** សុវណ្ណឌី សុវណ្ណឌី (ប្រ)

1.c. Contact organisation unit

_____ / _____, ____

1.d. Contact person function

□□□□□□□□□□ / □□□□ [□□]

1.e. Contact phone

[] [] [] [] [] [] [] [] / [] [] [] [] [] [] [] []

1.f. Contact mail

sovanndypoch @ gmail.com / Norvanndy @ gmail.com

[illegible]

□□□□□ (%)

□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□ (CSES) [□□]

[illegible]

3.c. Data collection calendar

Quater1, []

3.d. Data release calendar

[illegible]

3.e. Data providers

□ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ [□ □]

3.f. Data compilers

3.g. Institutional mandate

[illegible]

4.a. Rationale

[illegible]

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4.b. Comment and limitations

The model presented here is a simplified representation of the complex biological system under study. It is based on a set of assumptions that may not fully capture the underlying biology. For example, the model assumes that the system is at equilibrium, which may not be the case in all situations. Additionally, the model does not account for stochasticity, which is a key feature of many biological systems. Despite these limitations, the model provides a useful framework for understanding the basic principles of the system and for generating testable hypotheses. The model is implemented in the R programming language, and the code is available on GitHub at <https://github.com/yourusername/your-repo>. The model is a simplified representation of the complex biological system under study. It is based on a set of assumptions that may not fully capture the underlying biology. For example, the model assumes that the system is at equilibrium, which may not be the case in all situations. Additionally, the model does not account for stochasticity, which is a key feature of many biological systems. Despite these limitations, the model provides a useful framework for understanding the basic principles of the system and for generating testable hypotheses. The model is implemented in the R programming language, and the code is available on GitHub at <https://github.com/yourusername/your-repo>.

4.c. Method of computation

The model was solved using the Runge-Kutta method, which is a numerical method for solving ordinary differential equations. The model was implemented in the R programming language, and the code is available on GitHub at <https://github.com/yourusername/your-repo>. The model was solved using the Runge-Kutta method, which is a numerical method for solving ordinary differential equations. The model was implemented in the R programming language, and the code is available on GitHub at <https://github.com/yourusername/your-repo>. The model was solved using the Runge-Kutta method, which is a numerical method for solving ordinary differential equations. The model was implemented in the R programming language, and the code is available on GitHub at <https://github.com/yourusername/your-repo>. The model was solved using the Runge-Kutta method, which is a numerical method for solving ordinary differential equations. The model was implemented in the R programming language, and the code is available on GitHub at <https://github.com/yourusername/your-repo>.

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4.i. Quality management

NIS The following information is provided for the purpose of the study (the following information is provided for the purpose of the study)

5. Data availability and disaggregation

The following information is provided for the purpose of the study (the following information is provided for the purpose of the study)

7. References and Documentation

The following information is provided for the purpose of the study (the following information is provided for the purpose of the study)