Assessing Covid-19 Situation in Vietnam Using a Data-Driven Epidemiological Compartmental Model

Vo Le Tung Assessor: Assoc. Prof. Huynh Trung Hieu Co-Assessor: Assoc. Prof. Nguyen Tuan Duc

> Vietnamese - German University Frankfurt University of Applied Sciences

> > January 19, 2022



Outline

- Introduction
- 2 Literature review
- Methodologies
- 4 Results
- Discussion
- 6 Conclusion

Outline for Introduction

Introduction

The Covid-19 disease

- 230 million infections
- 4.7 deaths
- Economic impacts
- Mental impacts

Why do we need a model?

- Lack of research on the impacts of Non-Pharmaceutical Intervention (NPI) in Vietnam
- Need of a simple tool for assessing the disease current and future progression

Goals

- A model that works with the data availability in Vietnam
- An explainable model that can be used by experts
- Assessment of the effectiveness of mobility data in predicting future cases

Outline for Literature review

- 2 Literature review
 - Related works
 - Artificial Neural Network

Compartmental models [1]

Susceptible-Exposed-Infective-Removed (SEIR) model

$$S' = -\beta(N)SI$$

$$E' = \beta(N)SI - \kappa E$$

$$I' = \kappa E - \alpha I$$

$$R' = f\alpha I$$

$$N' = -(1 - f)\alpha I$$

Mathematical models II

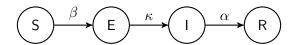


Figure: Graph of transitions between each compartment in the SEIR model

Mathematical models III

Changes made to classical compartmental model for Covid-19 [2]–[6]:

- Inclusion of compartments that represent government interventions
- Inclusion of compartments that specifically represent the behavior of SARS-NCOV-2
- Separation of the infectious compartment into multiple compartments representing the severity of the patients

Mathematical models IV

Agent-based models [7]-[9]

- Model of individualistic behaviors
- Based on population size, age structure, transmission networks

Mathematical models V

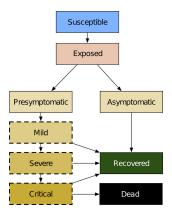


Figure: Agent states used in Covasim [7]

Mathematical models VI

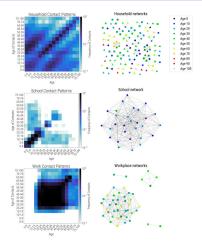


Figure: Transmission networks used in Covasim [7]

Pros

- Explainable
- Based on many years of research
- Easy to understand and implement

Cons

- Low representational capabilities
- The represented dynamics are stationary
- Unrealistic assumptions about real-world scenarios

Data-driven models I

ARIMA models [10]-[12]

- Suitable for modeling changing trends, periodic changes, and random noises
- Suitable for different types of data
- Can capture temporal dependency of time series

Data-driven models II

Deep-learning models [13]-[15]

- Long Short Term Memory (LSTM)
- Bidirectional Long Short Term Memory (Bi-LSTM)
- Gated Recurrent Unit (GRU)

Data-driven models III

Pros

- High prediction accuracy
- Allow for modeling without needing domain knowledge

Cons

- Unexplainable
- Relied on large amount of data
- Inability to capture the true disease dynamics

Novel compartmental models I

Mitigate the disadvantages of compartmental models by incorporating covariates

- using predefined functions based on domain knowledge
- using neural networks

Novel compartmental models II

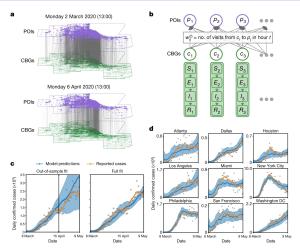


Figure: SEIR model weighted using complex mappings of point-to-point mobility data [16]



Novel compartmental models III

Related works

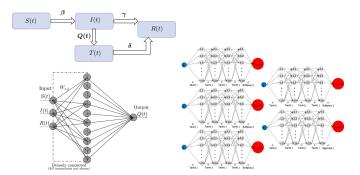


Figure: Using neural networks to predicts the parameters for compartmental models. On the left is the schematic for the QSIR model [17], on the right is the schematic for the time-dependent SIR model [18].

Artificial Neural Network

Multi-layer perceptron I

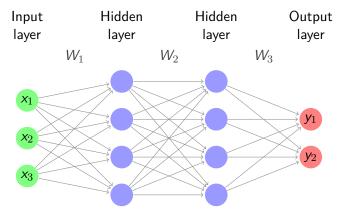


Figure: Graph representation of a multi-layer perceptron with four layers

Artificial Neural Network

Multi-layer perceptron II

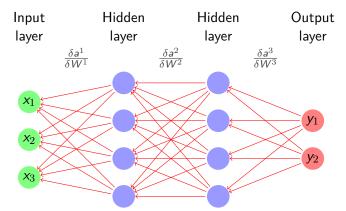


Figure: Graph representation of the back-propagation algorithm on a multi-layer perceptron with four layers

Outline for Methodologies

Methodologies

Methodologies

Outline for Results

4 Results

Results



Outline for Discussion

5 Discussion

Discussion

Outline for Conclusion

6 Conclusion

Conclusion

References I

- [1] F. Brauer, "Compartmental Models in Epidemiology," in *Mathematical Epidemiology*, ser. Lecture Notes in Mathematics, F. Brauer, P. van den Driessche, and J. Wu, Eds., red. by J. .-. Morel, F. Takens, and B. Teissier, vol. 1945, Berlin, Heidelberg: Springer Berlin Heidelberg, 2008, pp. 19–79.
- [2] S. Zhao and H. Chen, "Modeling the epidemic dynamics and control of COVID-19 outbreak in China," *Quantitative Biology*, vol. 8, no. 1, pp. 11–19, Mar. 2020.
- [3] S. He, Y. Peng, and K. Sun, "SEIR modeling of the COVID-19 and its dynamics," *Nonlinear Dynamics*, vol. 101, no. 3, pp. 1667–1680, Aug. 1, 2020.

References II

- [4] F. Ndaïrou, I. Area, J. J. Nieto, and D. F. Torres, "Mathematical modeling of COVID-19 transmission dynamics with a case study of Wuhan," *Chaos, Solitons, and Fractals*, vol. 135, p. 109 846, Jun. 2020.
- [5] S. B. Bastos and D. O. Cajueiro, "Modeling and forecasting the early evolution of the Covid-19 pandemic in Brazil," *Scientific Reports*, vol. 10, no. 1, p. 19457, Dec. 2020.
- [6] K. Sarkar, S. Khajanchi, and J. J. Nieto, "Modeling and forecasting the COVID-19 pandemic in India," *Chaos, Solitons, and Fractals*, vol. 139, p. 110 049, Oct. 2020.
- [7] C. C. Kerr, R. M. Stuart, D. Mistry, et al., "Covasim: An agent-based model of COVID-19 dynamics and interventions," *PLOS Computational Biology*, vol. 17, no. 7, e1009149, Jul. 26, 2021.

References III

- [8] P. C. Silva, P. V. Batista, H. S. Lima, M. A. Alves, F. G. Guimarães, and R. C. Silva, "COVID-ABS: An agent-based model of COVID-19 epidemic to simulate health and economic effects of social distancing interventions," *Chaos, Solitons, and Fractals*, vol. 139, p. 110 088, Oct. 2020.
- [9] N. Hoertel, M. Blachier, C. Blanco, et al., "A stochastic agent-based model of the SARS-CoV-2 epidemic in France," Nature Medicine, vol. 26, no. 9, pp. 1417–1421, 9 Sep. 2020.
- [10] Z. Ceylan, "Estimation of COVID-19 prevalence in Italy, Spain, and France," Science of The Total Environment, vol. 729, p. 138 817, Aug. 10, 2020.

References IV

- [11] R. K. Singh, M. Rani, A. S. Bhagavathula, et al., "Prediction of the COVID-19 Pandemic for the Top 15 Affected Countries: Advanced Autoregressive Integrated Moving Average (ARIMA) Model," JMIR Public Health and Surveillance, vol. 6, no. 2, e19115, May 13, 2020.
- [12] M. H. D. M. Ribeiro, R. G. da Silva, V. C. Mariani, and L. d. S. Coelho, "Short-term forecasting COVID-19 cumulative confirmed cases: Perspectives for Brazil," *Chaos, Solitons & Fractals*, vol. 135, p. 109 853, Jun. 1, 2020.
- [13] V. K. R. Chimmula and L. Zhang, "Time series forecasting of COVID-19 transmission in Canada using LSTM networks," *Chaos, Solitons, and Fractals*, vol. 135, p. 109864, Jun. 2020.

References V

- [14] F. Shahid, A. Zameer, and M. Muneeb, "Predictions for COVID-19 with deep learning models of LSTM, GRU and Bi-LSTM," *Chaos, Solitons, and Fractals*, vol. 140, p. 110 212, Nov. 2020.
- [15] A. Ramchandani, C. Fan, and A. Mostafavi, "DeepCOVIDNet: An Interpretable Deep Learning Model for Predictive Surveillance of COVID-19 Using Heterogeneous Features and Their Interactions," *IEEE Access*, vol. 8, pp. 159 915–159 930, 2020.
- [16] S. Chang, E. Pierson, P. W. Koh, et al., "Mobility network models of COVID-19 explain inequities and inform reopening," *Nature*, vol. 589, no. 7840, pp. 82–87, 7840 Jan. 2021.

References VI

- [17] R. Dandekar, C. Rackauckas, and G. Barbastathis, "A Machine Learning-Aided Global Diagnostic and Comparative Tool to Assess Effect of Quarantine Control in COVID-19 Spread," *Patterns*, vol. 1, no. 9, p. 100145, Dec. 11, 2020.
- [18] S. Y. Jung, H. Jo, H. Son, and H. J. Hwang, "Real-World Implications of a Rapidly Responsive COVID-19 Spread Model with Time-Dependent Parameters via Deep Learning: Model Development and Validation," *Journal of Medical Internet Research*, vol. 22, no. 9, e19907, Sep. 9, 2020.