

Branching stochastic processes as models of Covid-19 epidemic development

Myanmar - 20201214

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Abstract

The results presented here are obtained using the methodology proposed in the paper <https://arxiv.org/abs/2004.14838> for the country Myanmar. The data comes from European Centre for Disease Prevention and Control available at <https://opendata.ecdc.europa.eu/covid19/casedistribution/csv>.

Table of Contents

1. Observed Infection data	1
2. Estimating of the main parameter and some predictions	3

List of Figures

1.1. Number of the daily reported laboratory-confirmed cases	1
1.2. Number of the total registered cases	2
2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate	3
2.2. Figure	4
2.3. Expected number of the nonregistered infected individuals without immigra- tion	5
2.4. Expected number of the nonregistered infected individuals with immigration	6

Chapter 1. Observed Infection data

Figure 1.1. Number of the daily reported laboratory-confirmed cases

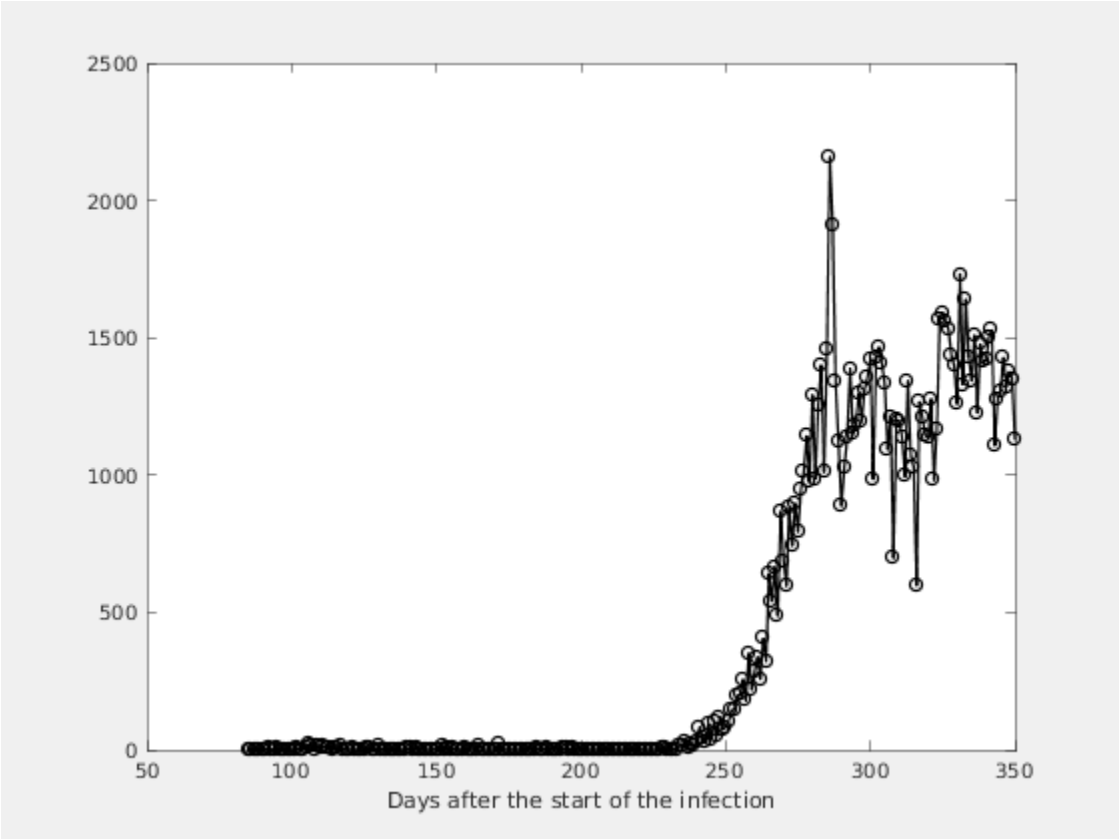
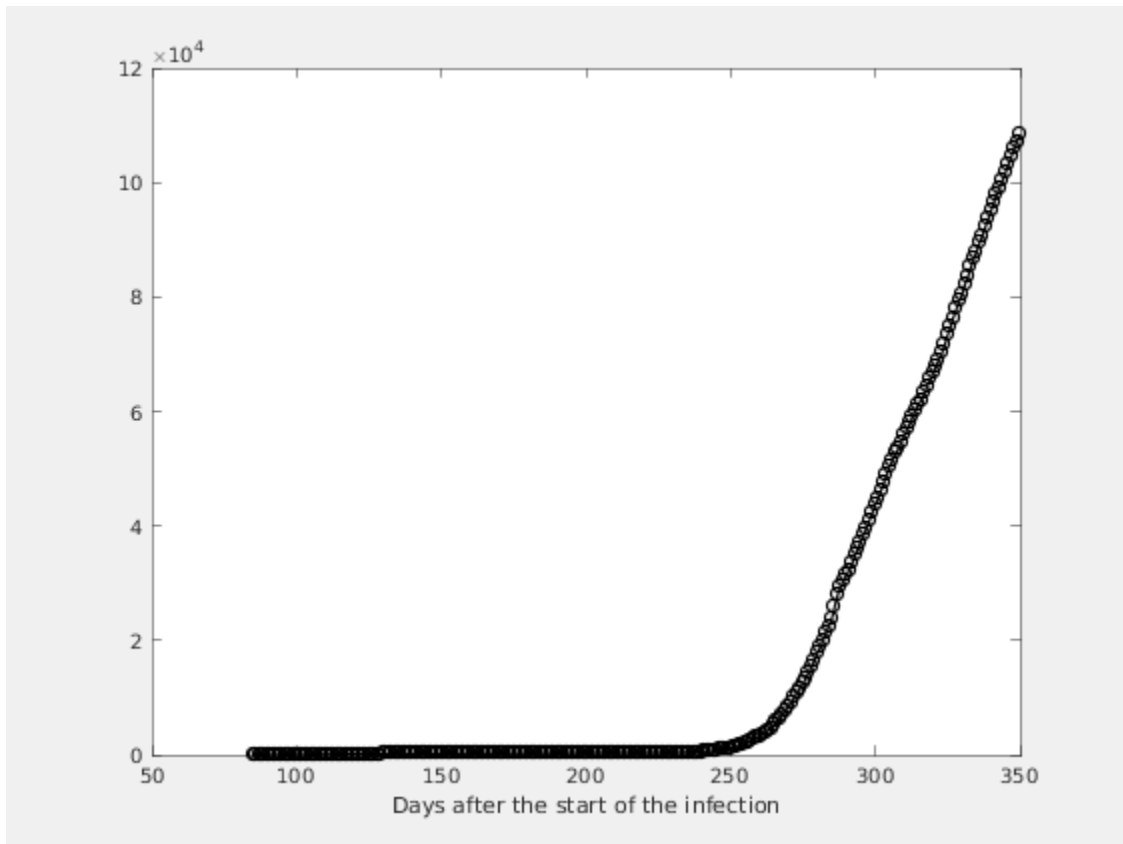


Figure 1.2. Number of the total registered cases



Chapter 2. Estimating of the main parameter and some predictions

Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate

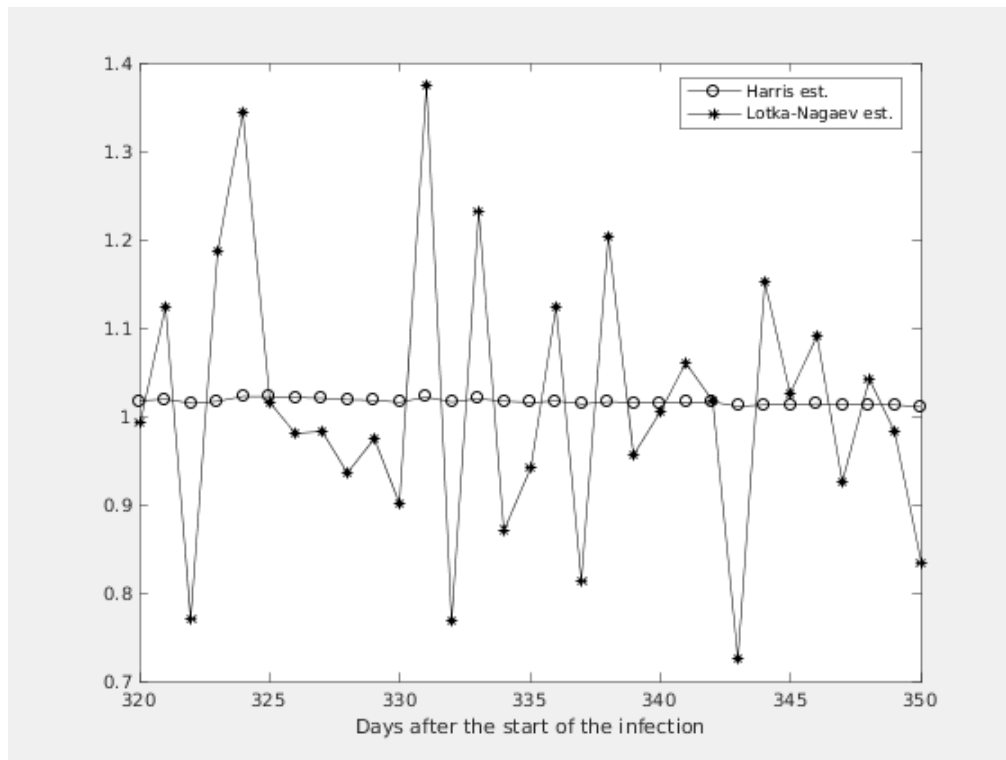


Figure 2.2. Figure

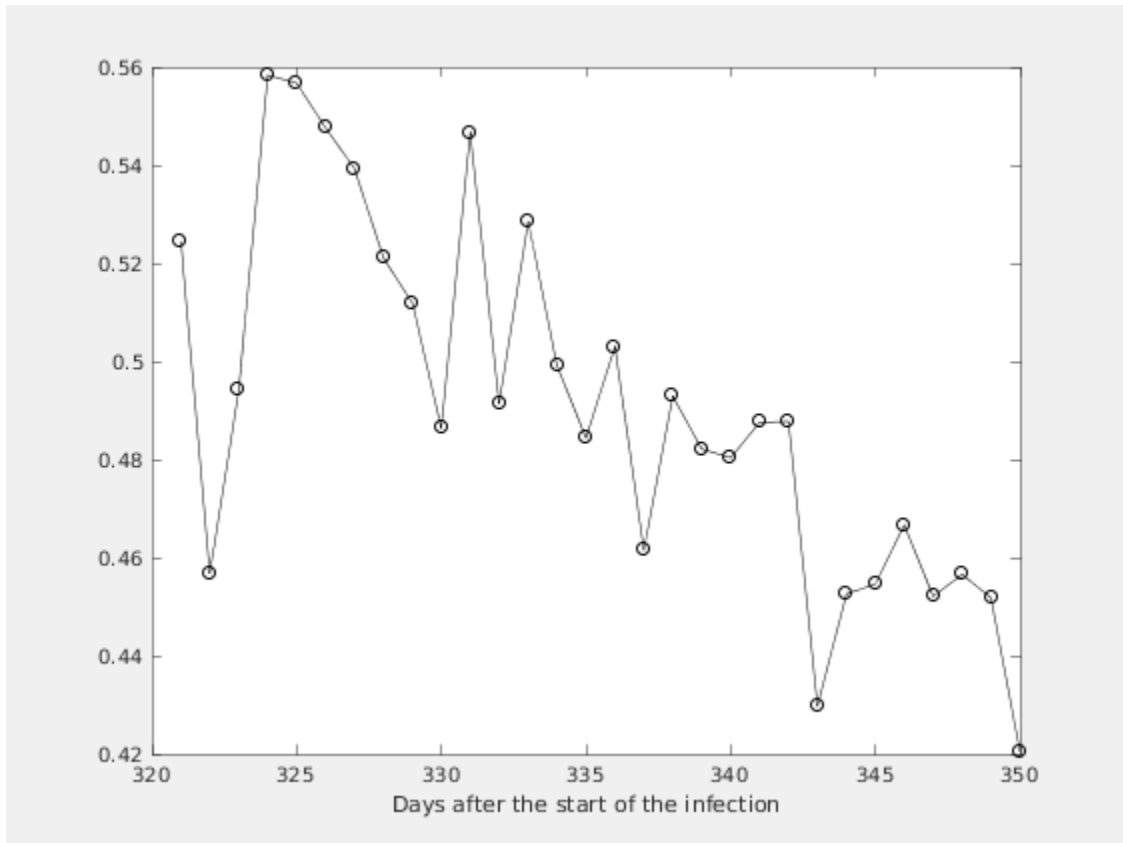


Figure 2.3. Expected number of the nonregistered infected individuals without immigration

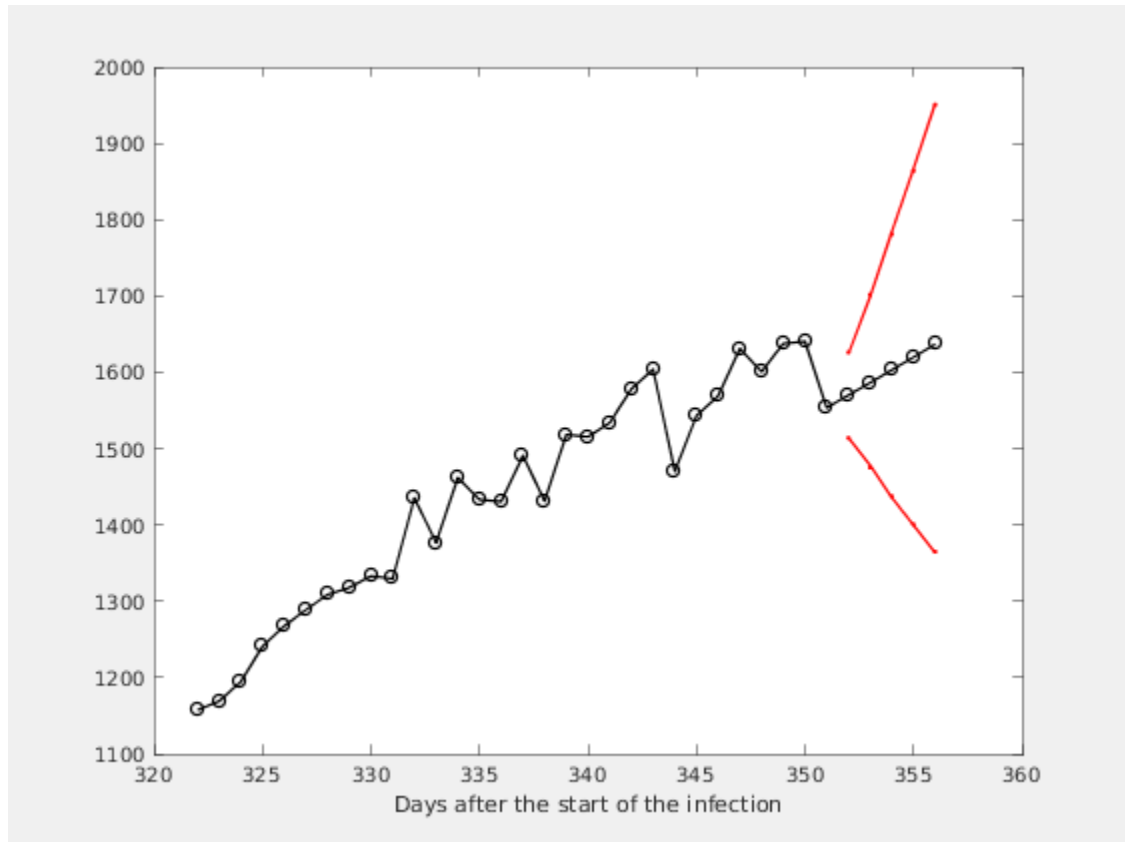
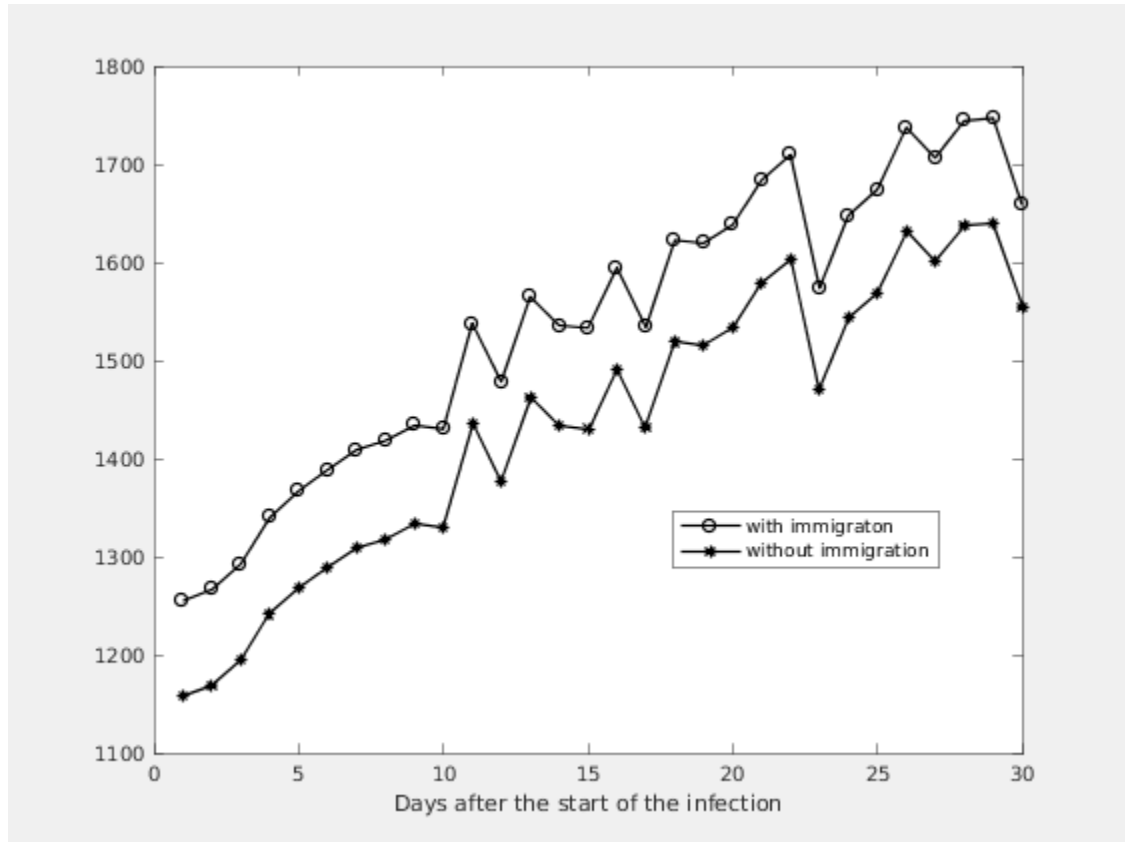


Figure 2.4. Expected number of the nonregistered infected individuals with immigration



Estimation of the model parameters.

k	m	ci	alpha	M1	A1
4	1.0140	0.9761 - 1.0519	0.4526	1543	1648
3	1.0128	0.9752 - 1.0503	0.4546	1569	1674
2	1.0131	0.9759 - 1.0504	0.4667	1631	1737
1	1.0128	0.9758 - 1.0497	0.4521	1601	1707
0	1.0105	0.9739 - 1.0471	0.4566	1638	1744