

# **Branching stochastic processes as models of Covid-19 epidemic development**

**AntiguaAndBarbuda - week 53**

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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 AntiguaAndBarbuda. The data comes from European Centre for Disease Prevention and Control available at <https://opendata.ecdc.europa.eu/covid19/casedistribution/csv>.

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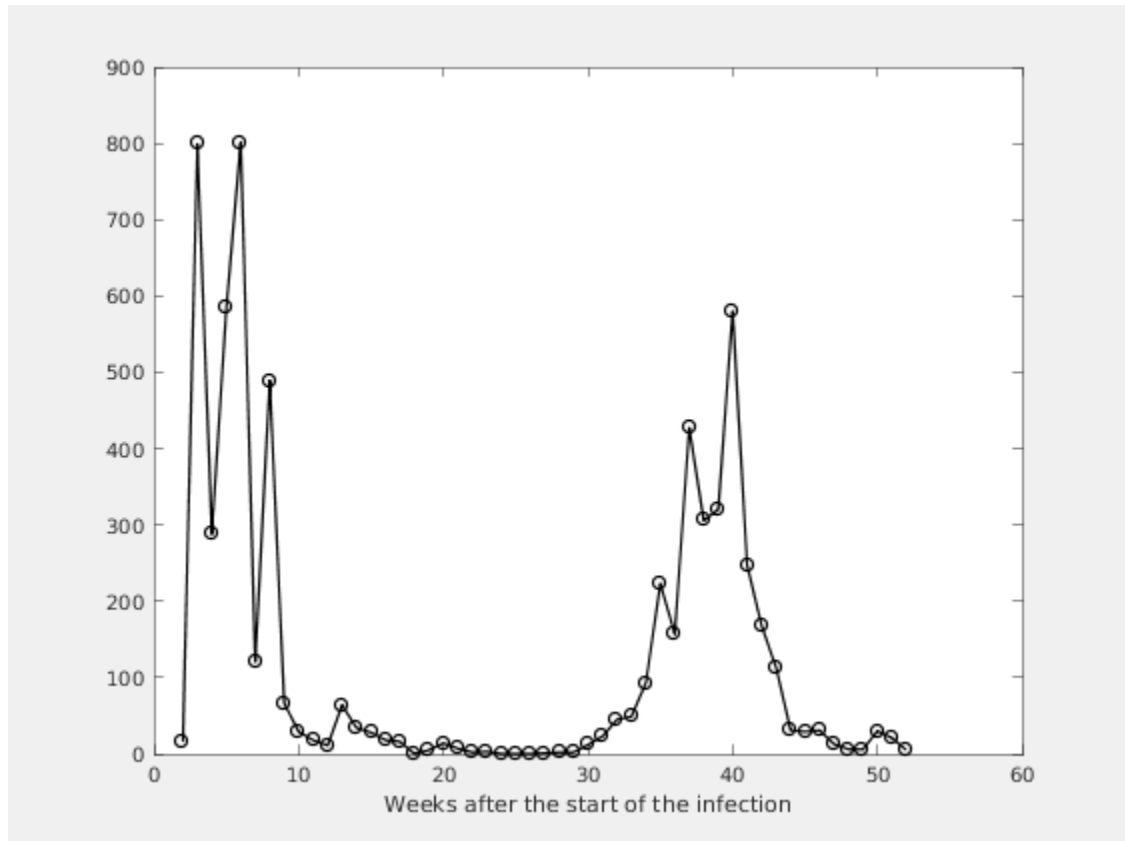
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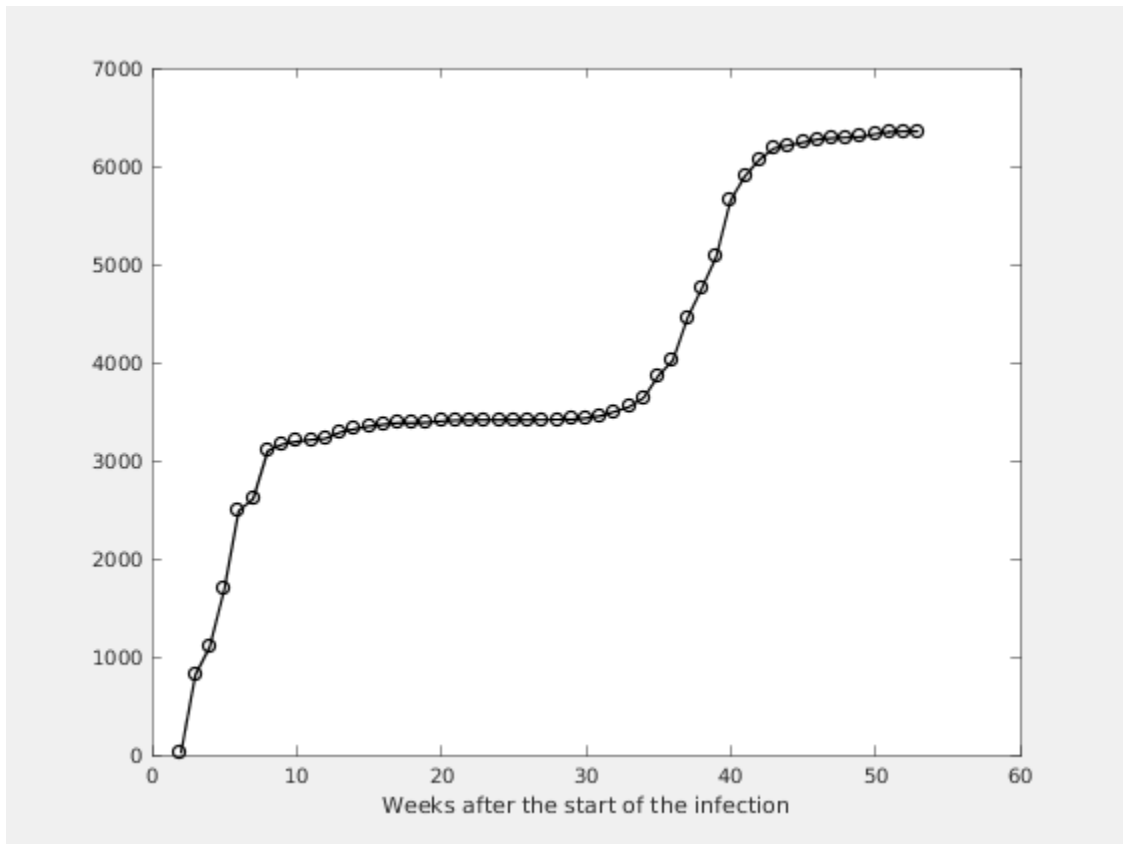
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# Chapter 1. Observed Infection data

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



**Figure 1.2. Number of the total registered cases**



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# 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

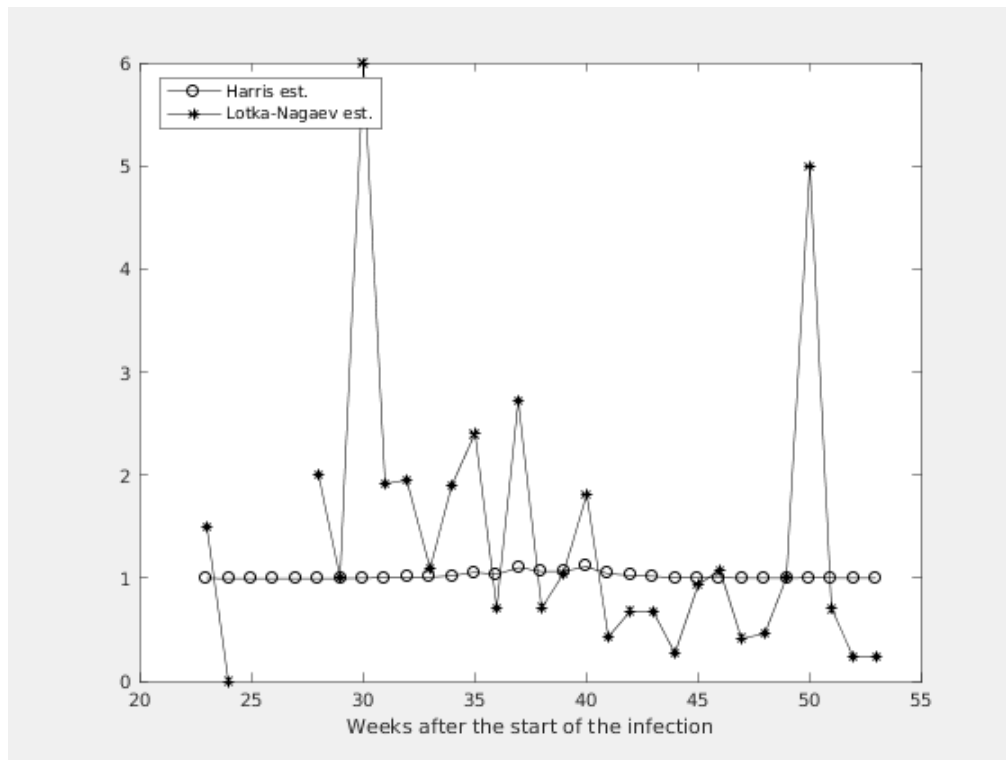
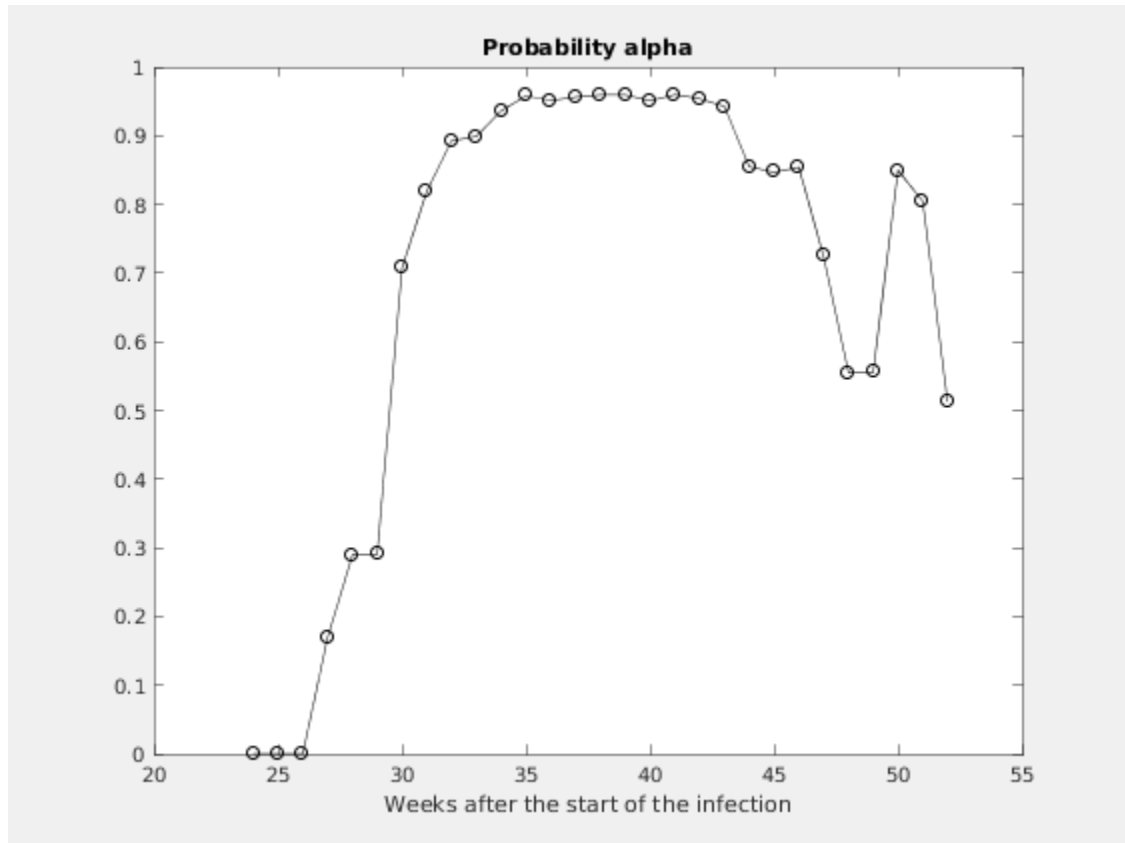
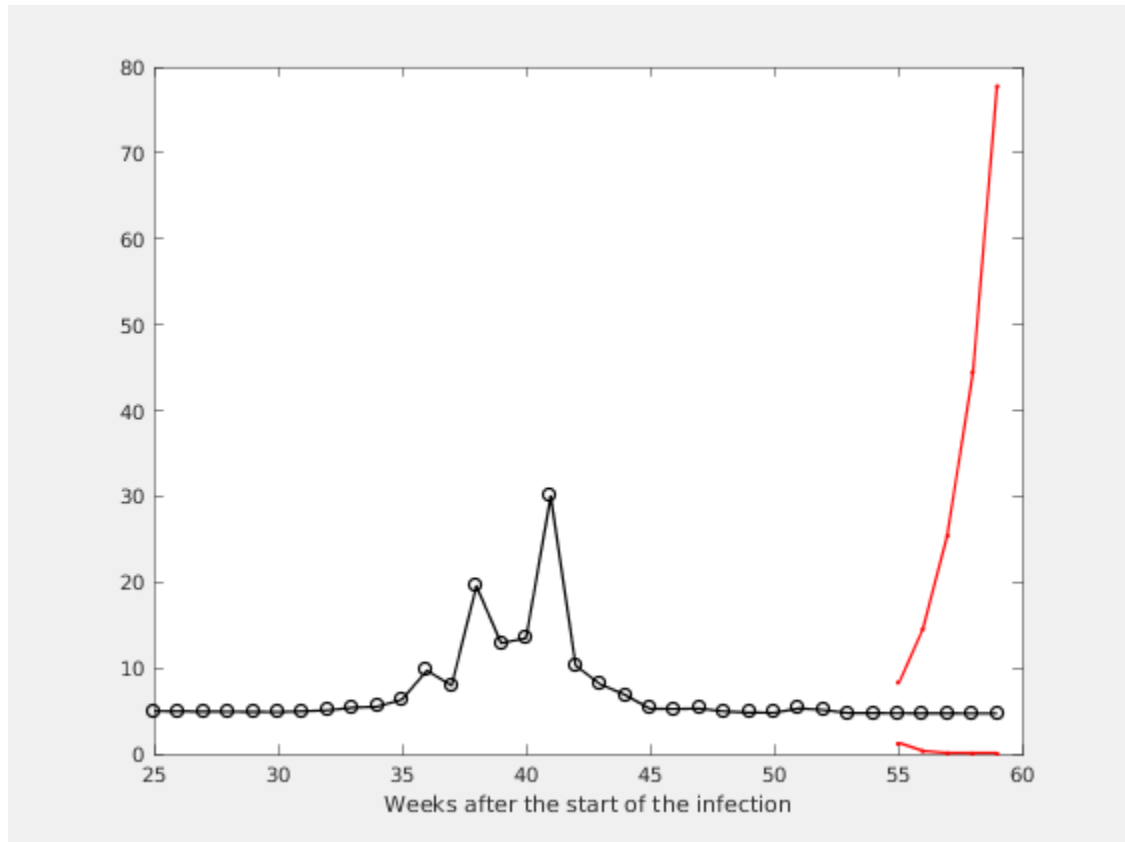


Figure 2.2. Figure

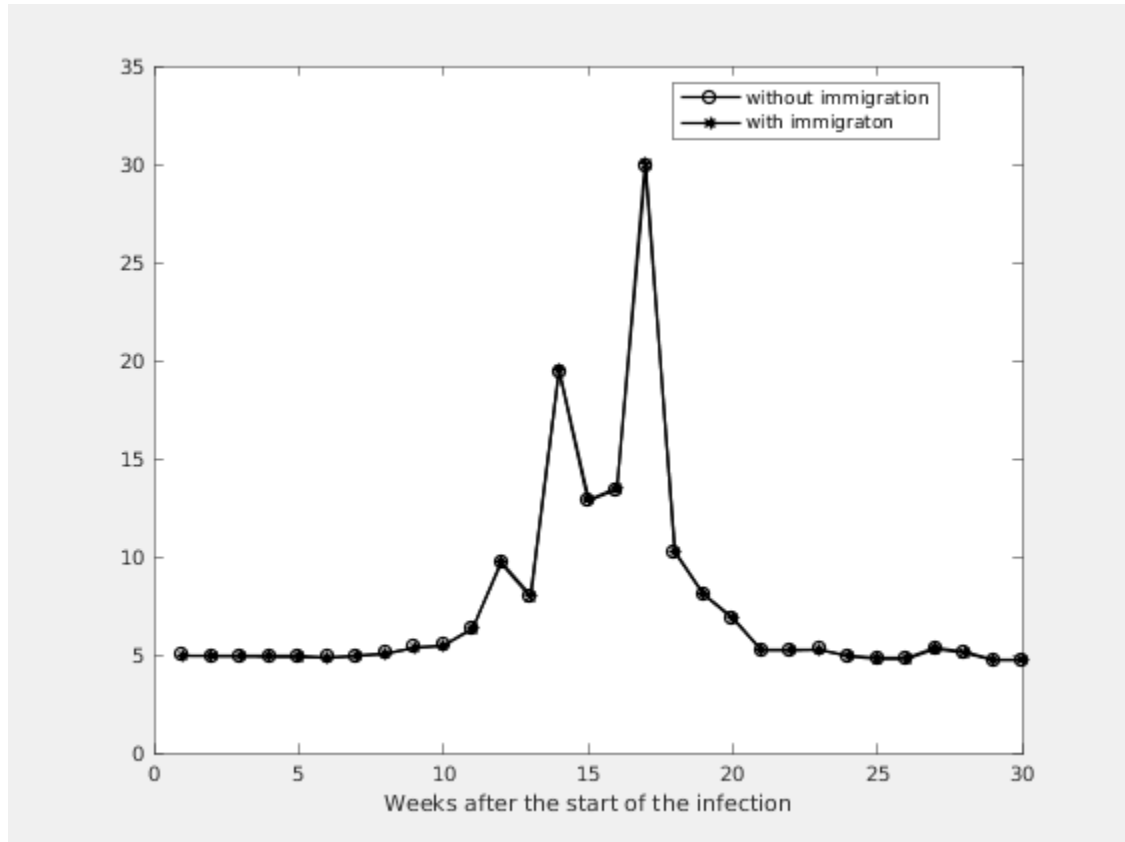




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

k	m	ci	alpha	A1	M1
4	0.9984	0.1992 - 1.7976	0.7245	5	5
3	1.0022	0.2134 - 1.7911	0.5553	5	5
2	1.0008	0.2215 - 1.7801	0.5557	5	5
1	0.9983	0.2274 - 1.7691	0.8496	5	5
0	0.9983	0.2378 - 1.7587	0.8042	5	5