

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

**Barbados - 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 Barbados. 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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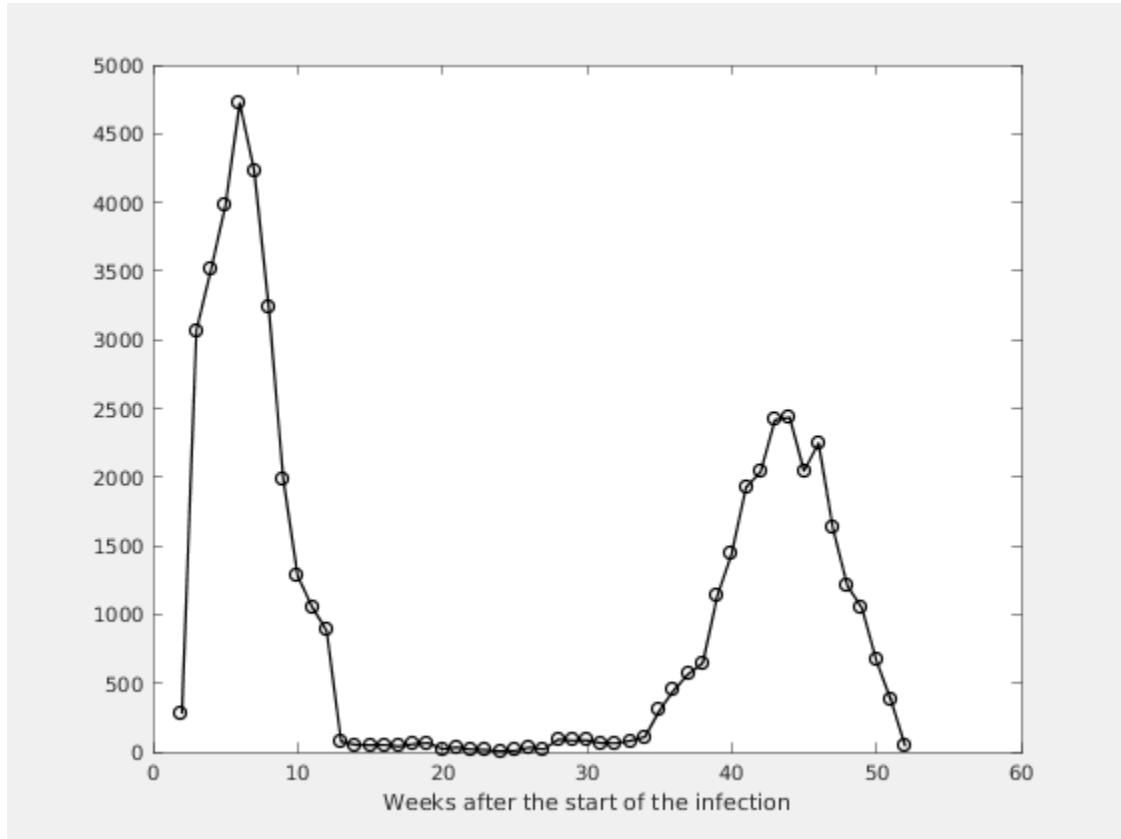
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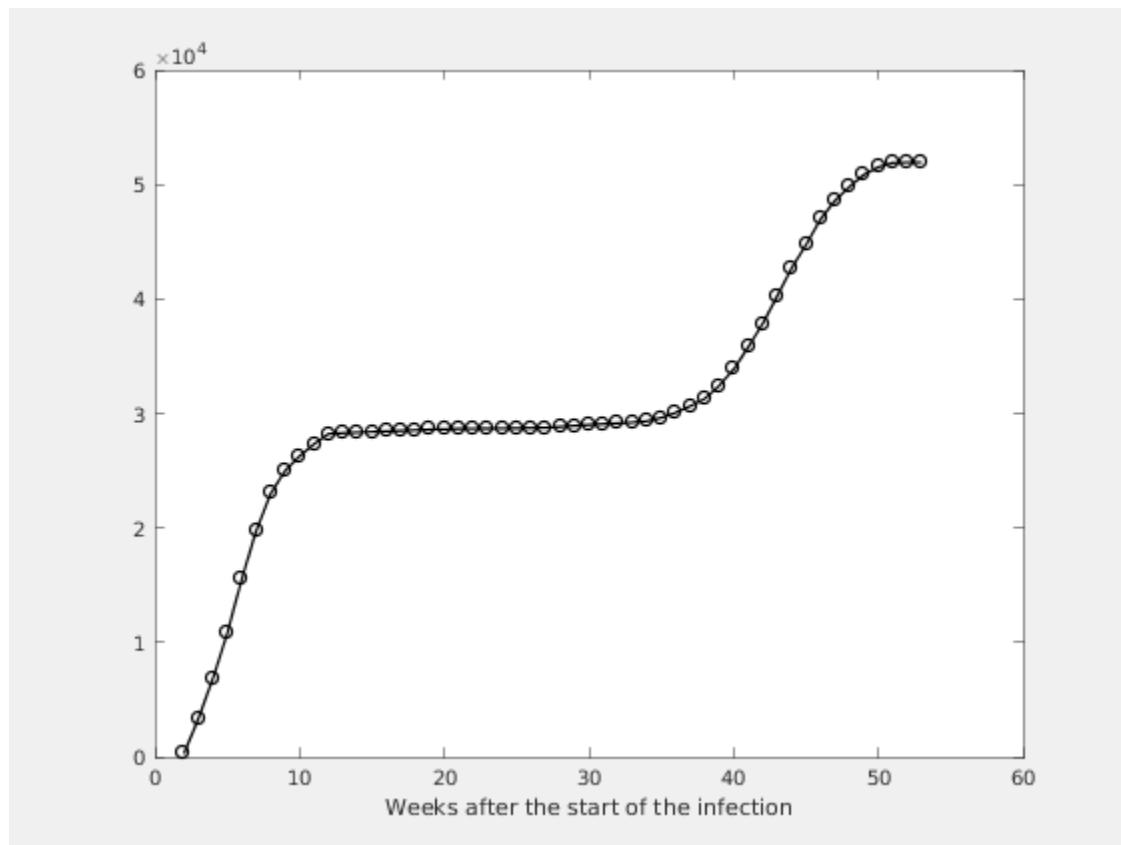
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# Chapter 1. Observed Infection data

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



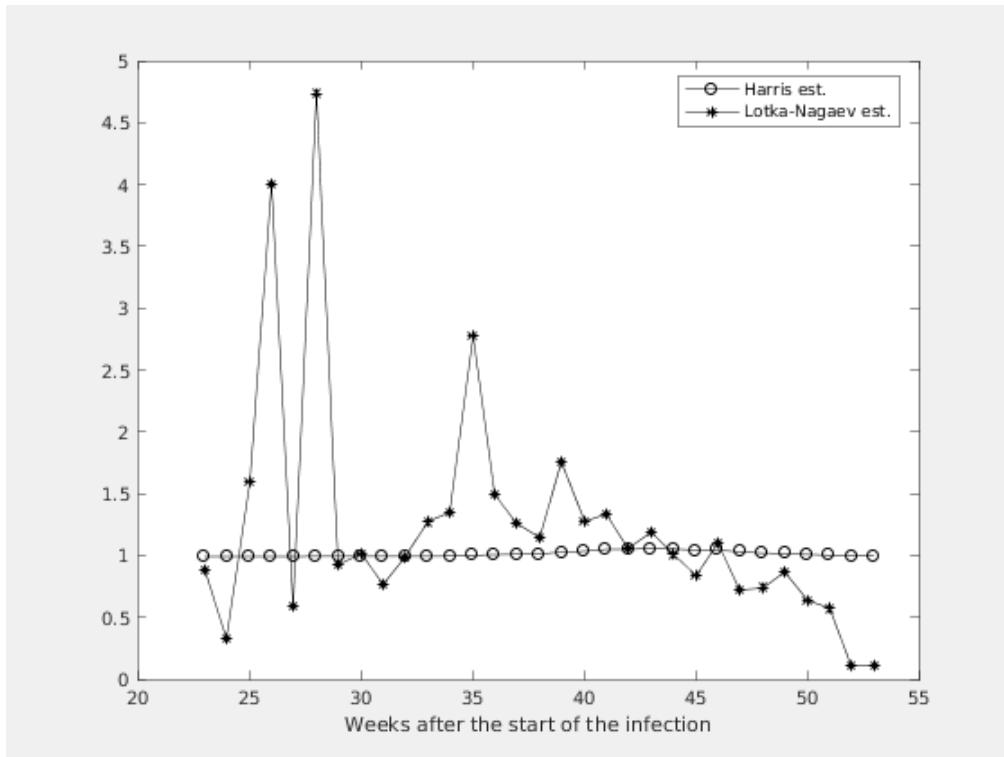
**Figure 1.2. Number of the total registered cases**



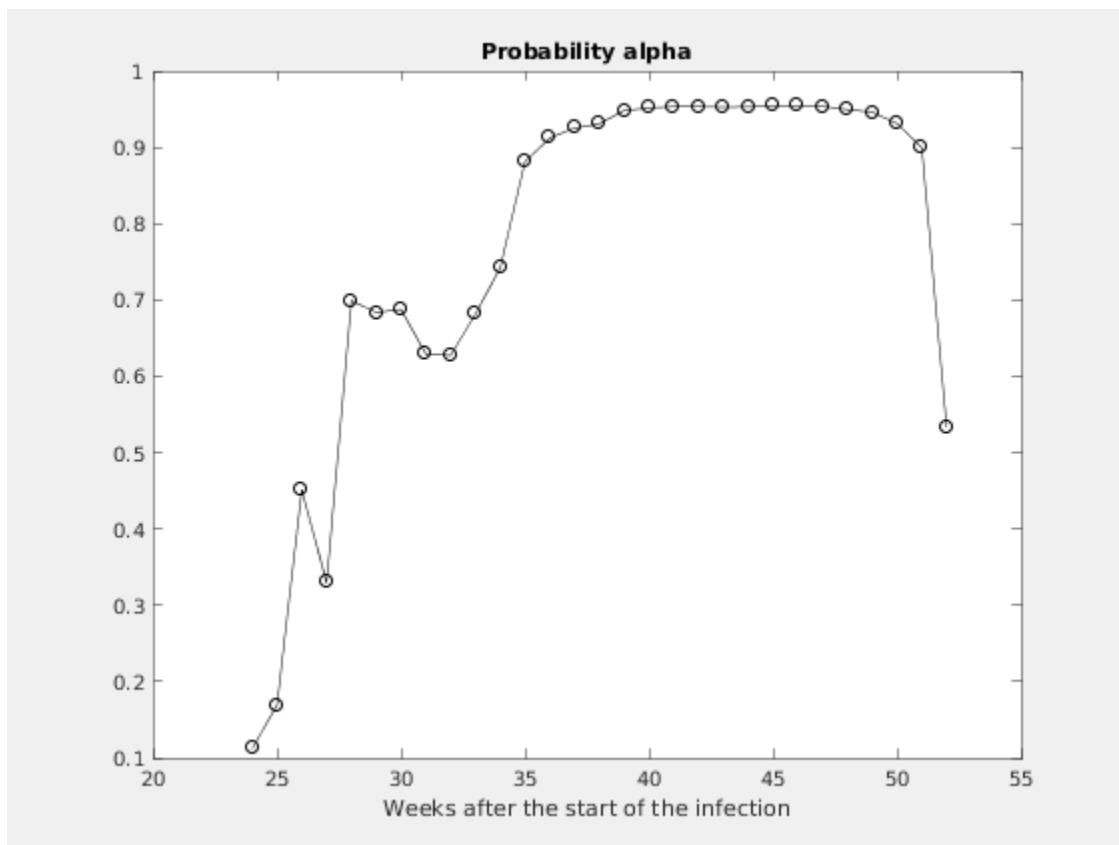
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# Chapter 2. Estimating of the main parameter and some predictions

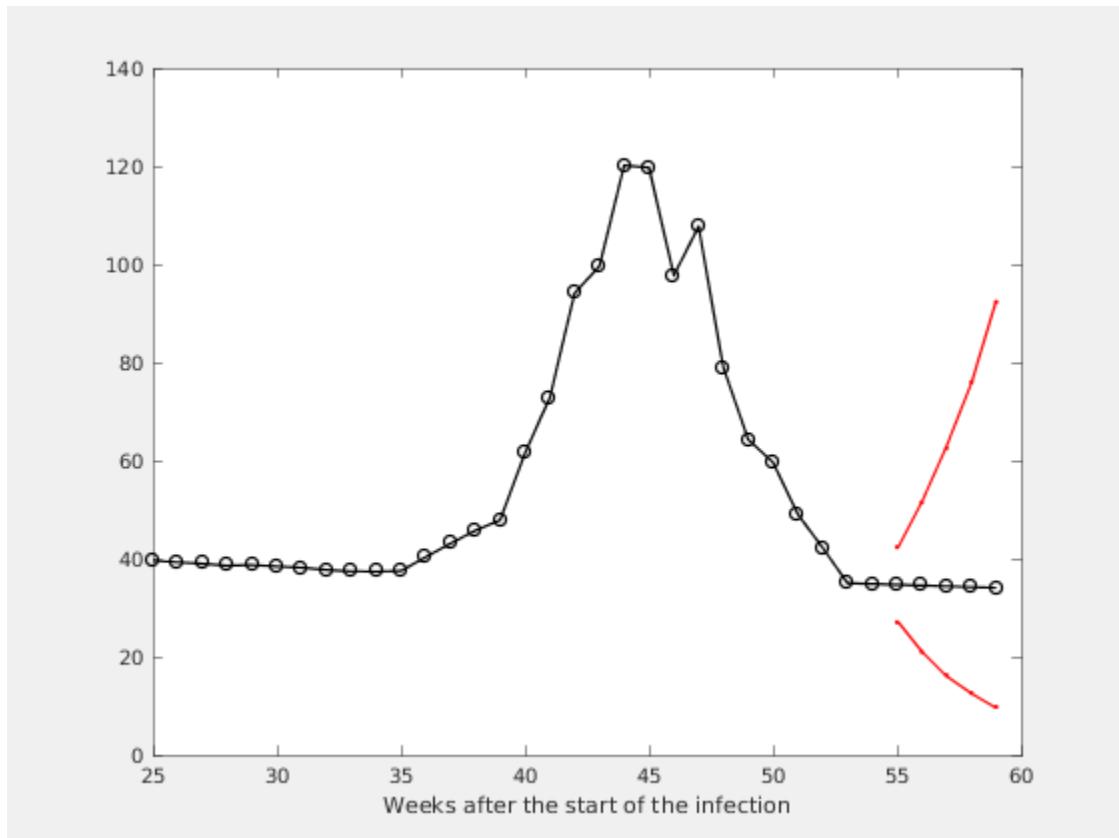
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



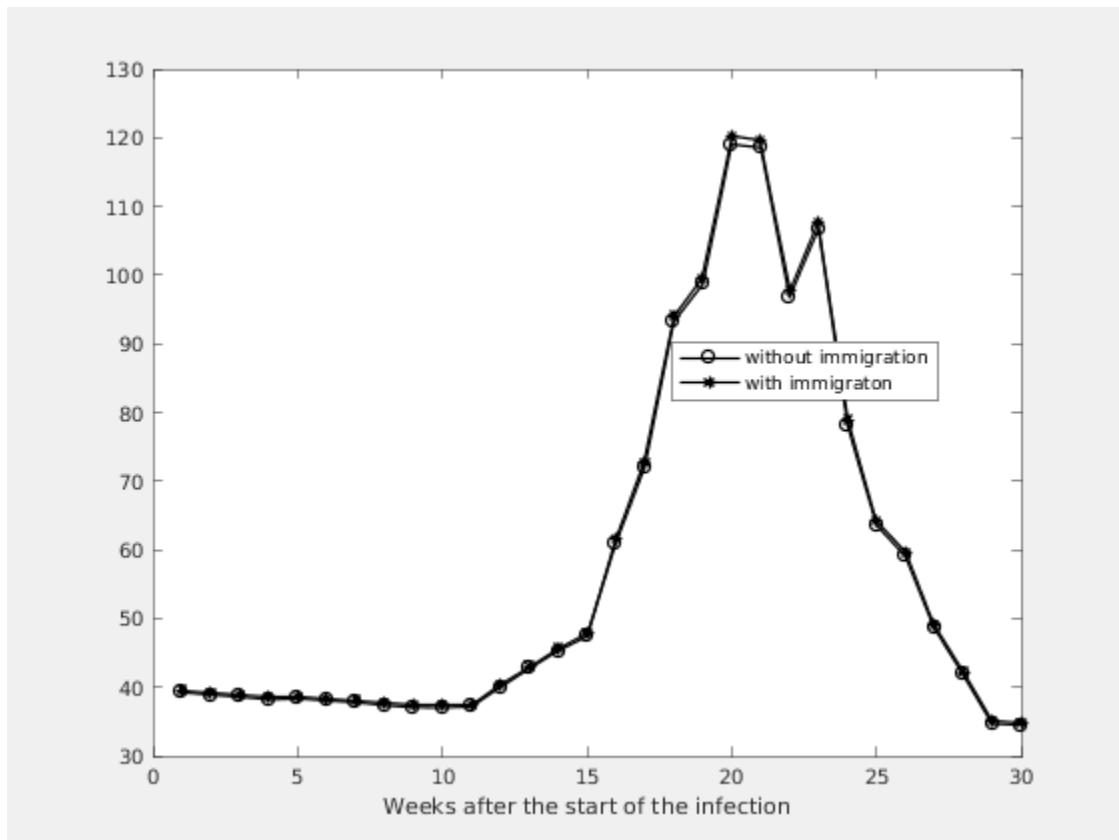
**Figure 2.2. Figure**



**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	1.0155	0.7747 - 1.2562	0.9537	79	78	
3	1.0076	0.7730 - 1.2421	0.9495	64	64	
2	1.0019	0.7726 - 1.2313	0.9462	60	59	
1	0.9954	0.7703 - 1.2205	0.9312	49	49	
0	0.9954	0.7737 - 1.2172	0.8995	42	42	