

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

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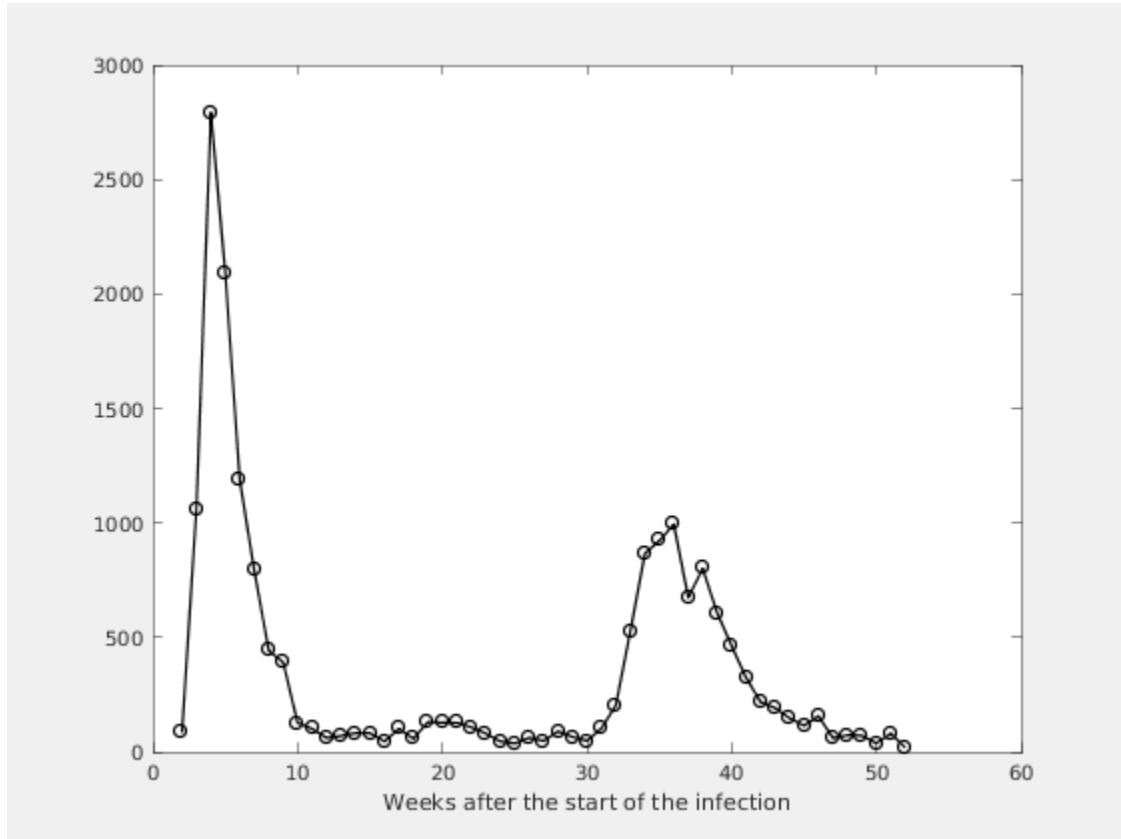
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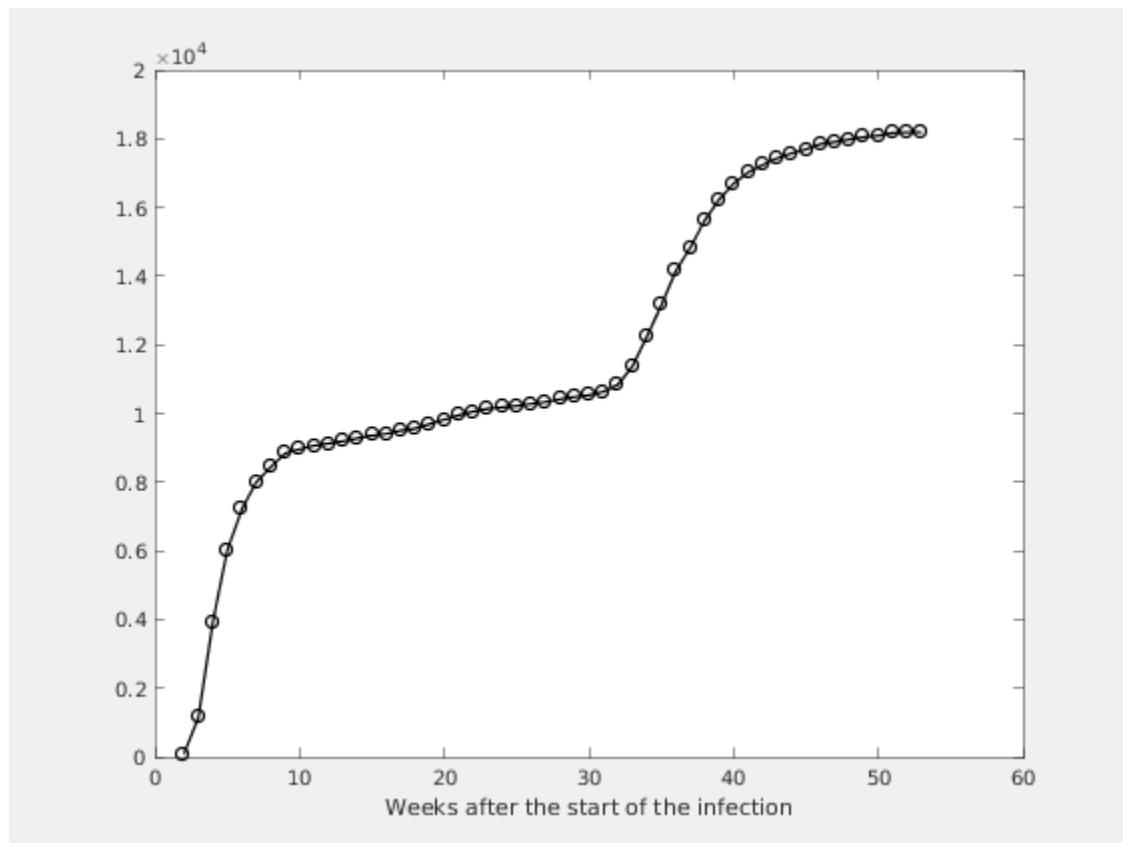
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# Chapter 1. Observed Infection data

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



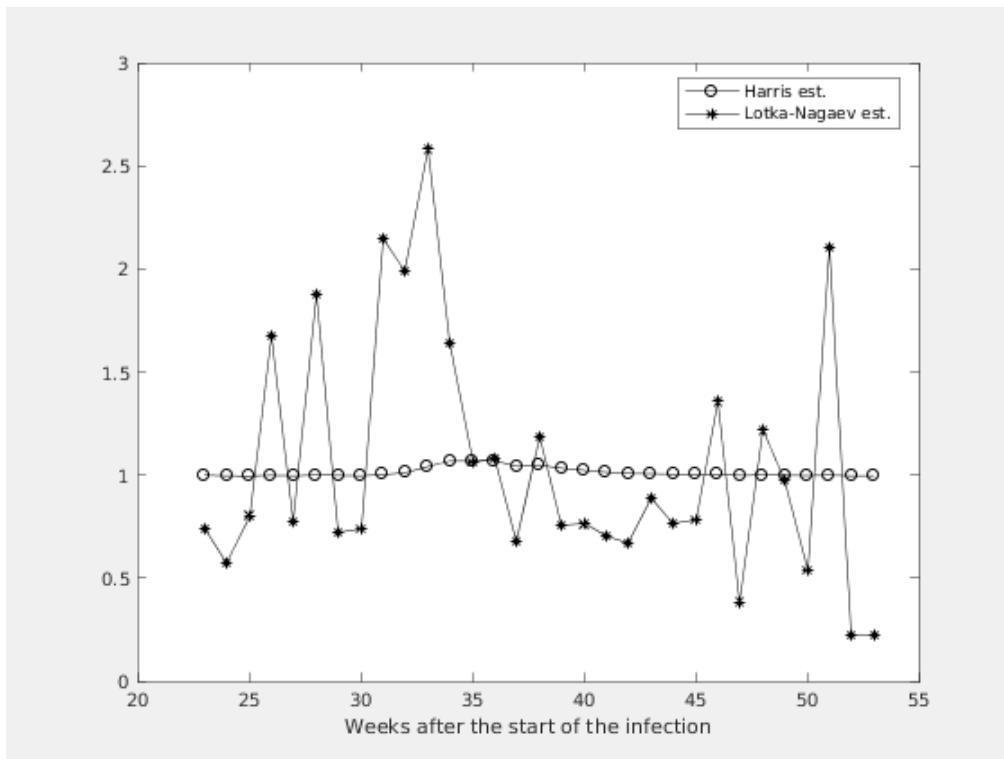
**Figure 1.2. Number of the total registered cases**



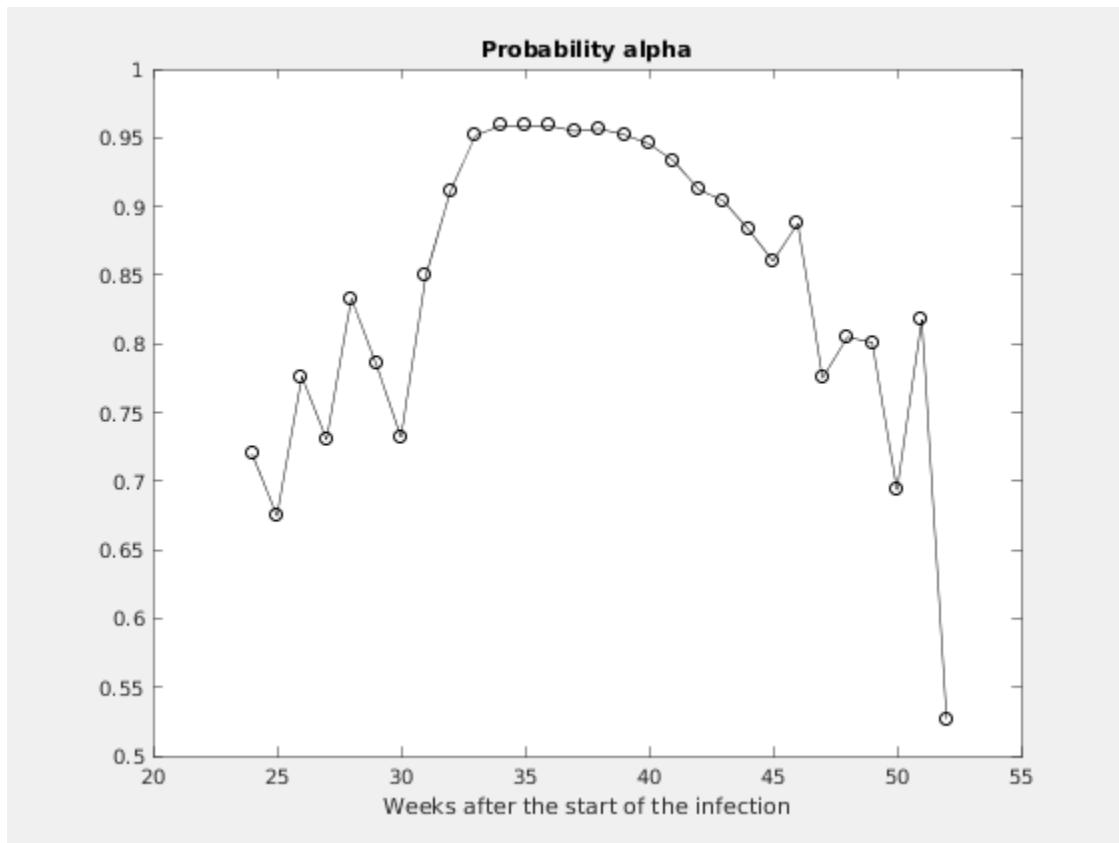
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# Chapter 2. Estimating of the main parameter and some predictions

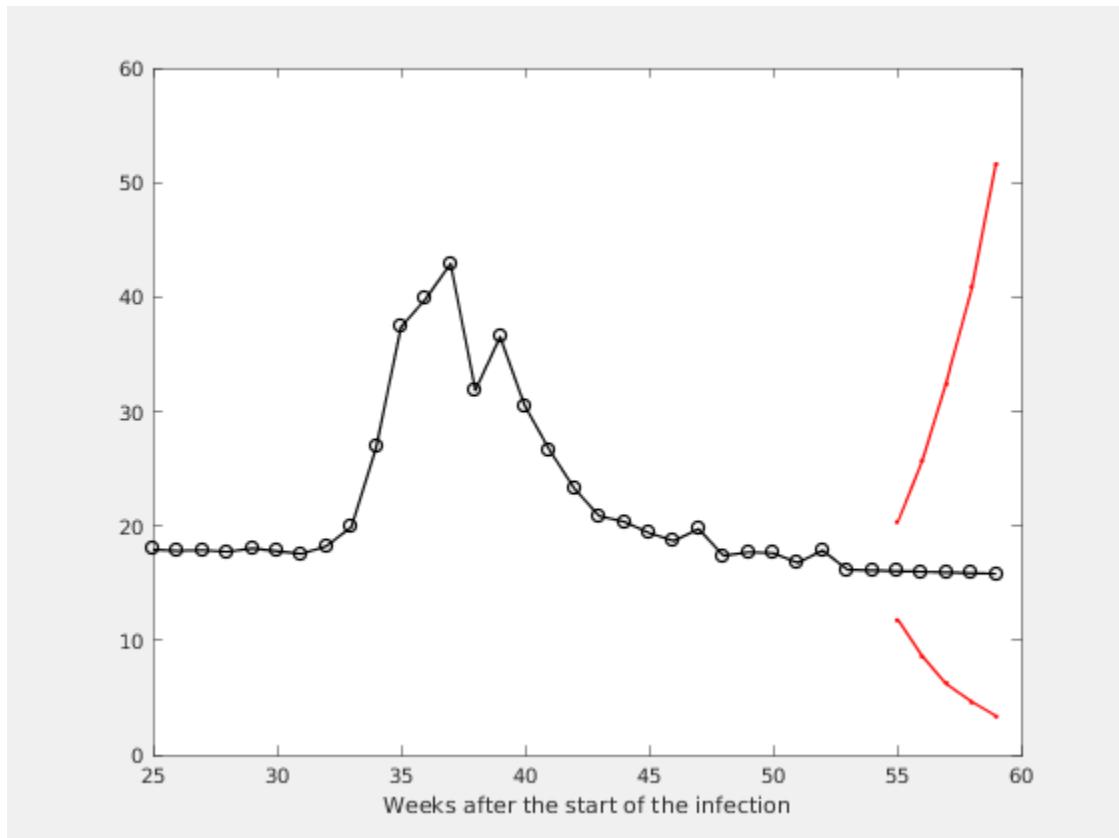
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



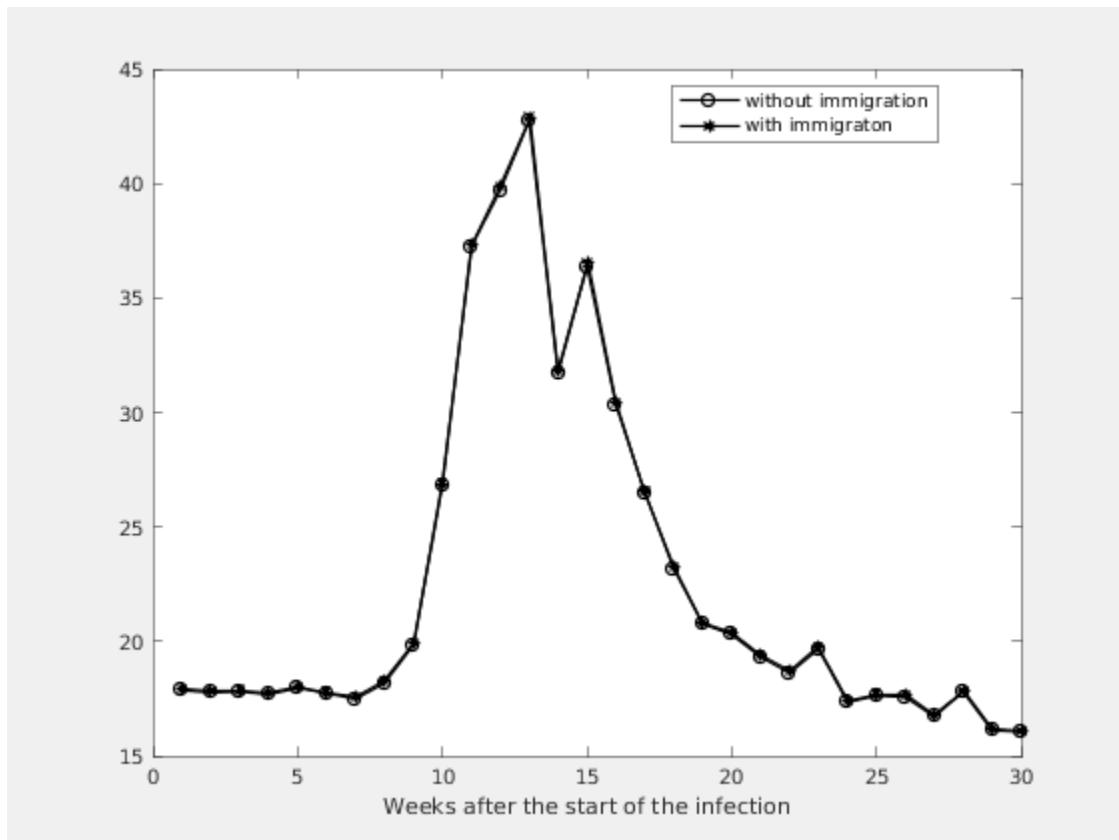
**Figure 2.2. Figure**



**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	
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4	0.9992	0.7174	- 1.2810	0.7751	17	17
3	0.9974	0.7191	- 1.2757	0.8048	18	18
2	0.9997	0.7250	- 1.2745	0.8010	18	18
1	0.9963	0.7248	- 1.2678	0.6937	17	17
0	0.9963	0.7275	- 1.2651	0.8175	18	18