

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

**Eswatini - week 53**

**N. Yanev, V. Stoimenova, D. Atanasov**

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### **Abstract**

The results presented here are obtained using the method proposed in the paper <https://arxiv.org/abs/2004.14838> for the country Eswatini. 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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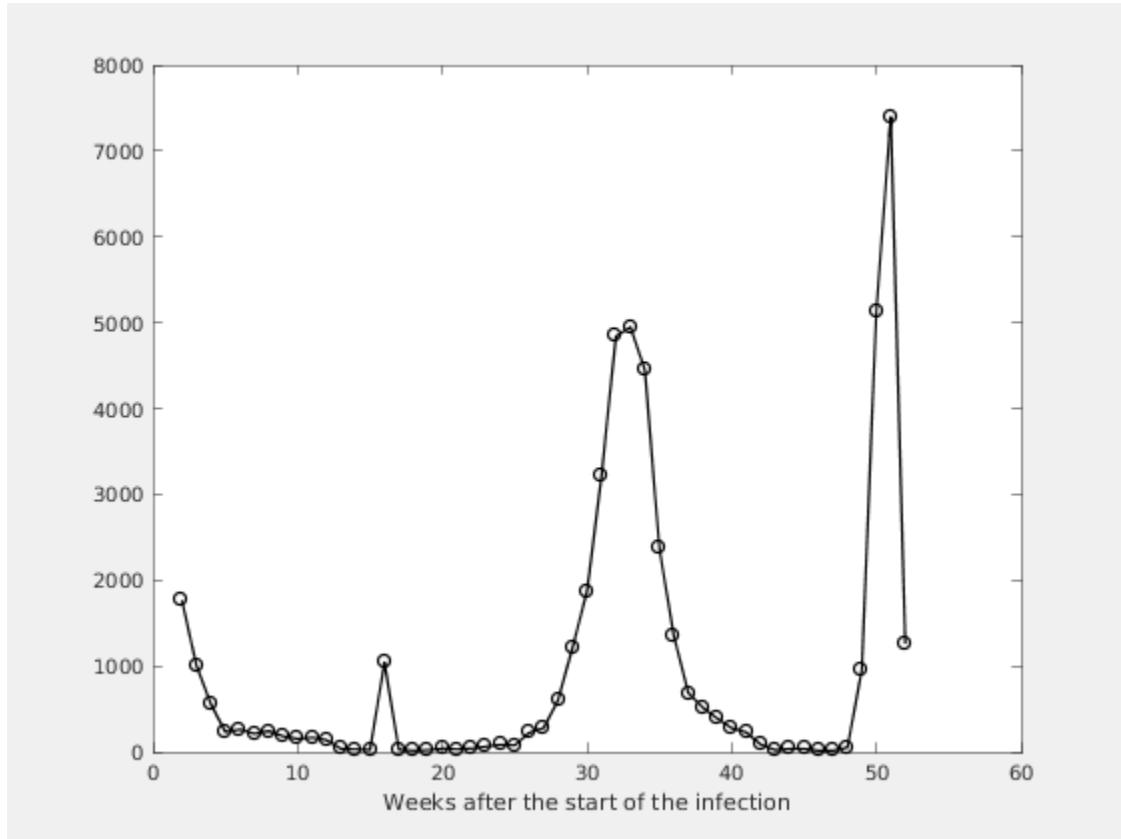
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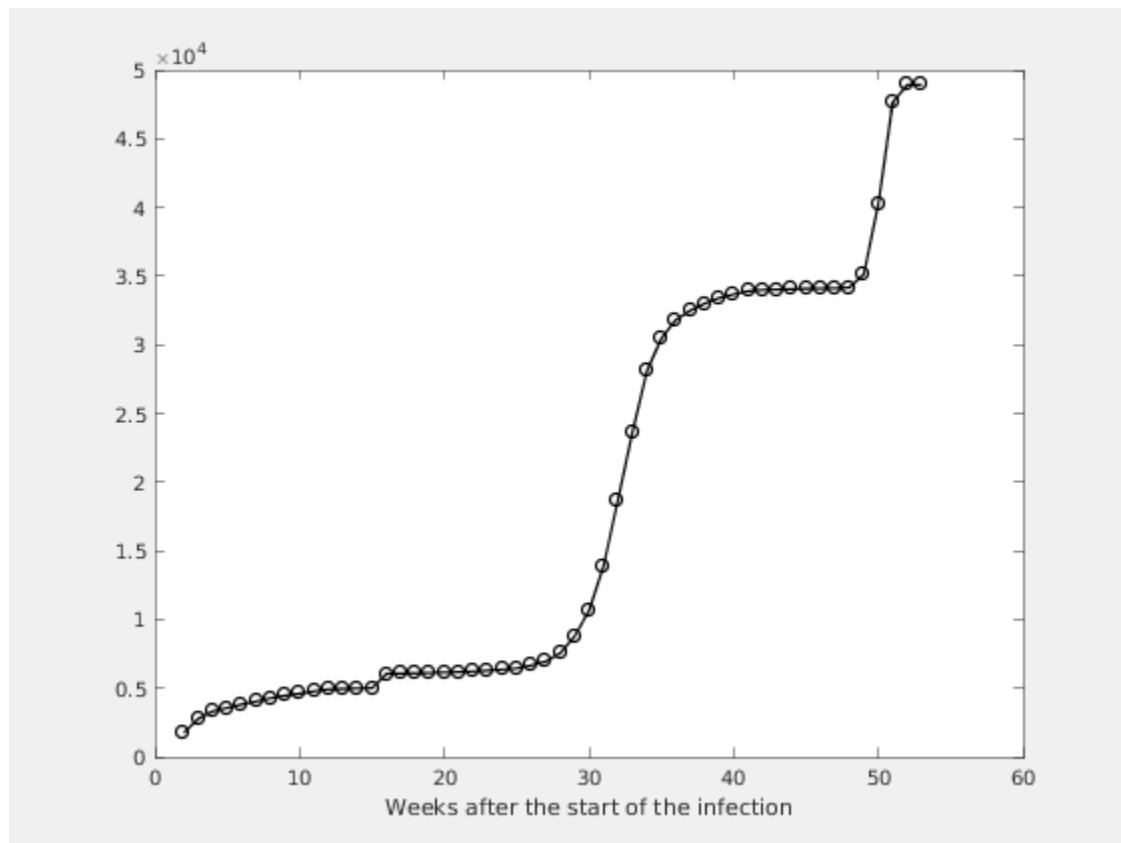
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# Chapter 1. Observed Infection data

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



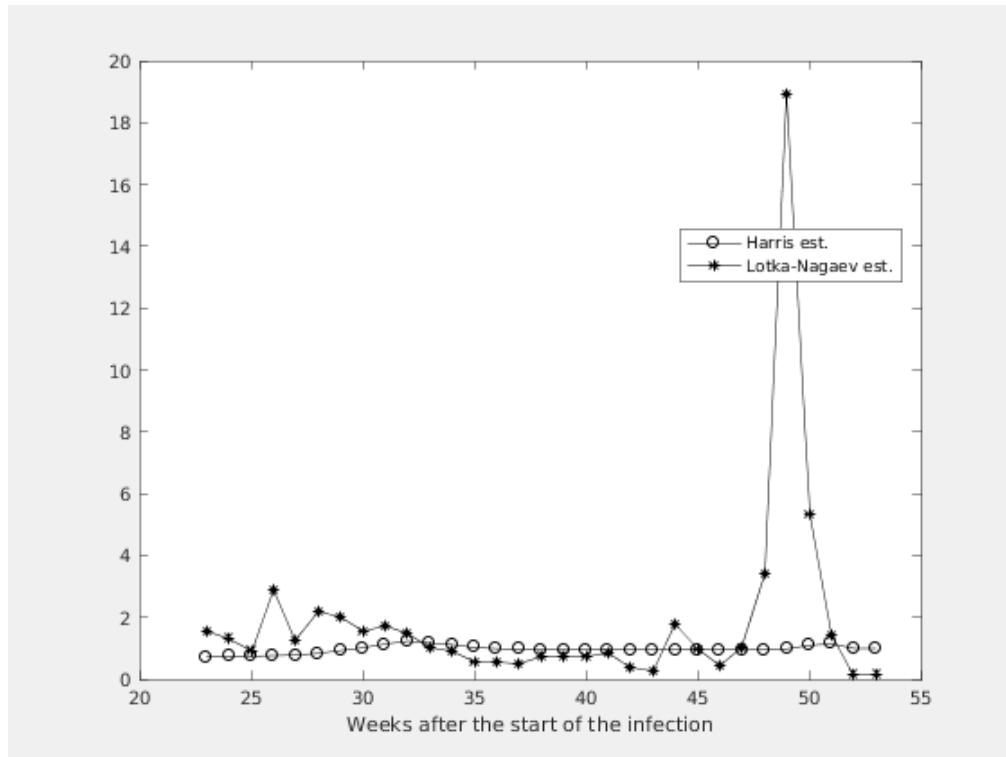
**Figure 1.2. Number of the total registered cases**



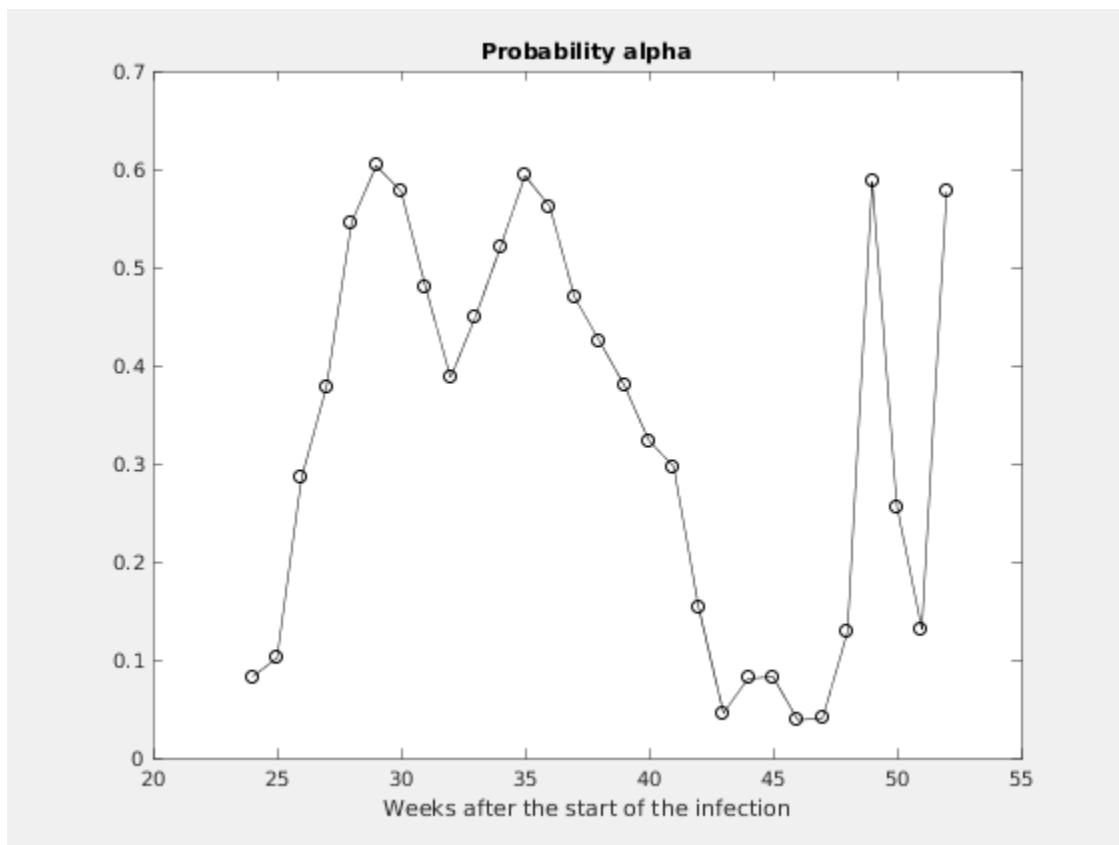
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# Chapter 2. Estimating of the main parameter and some predictions

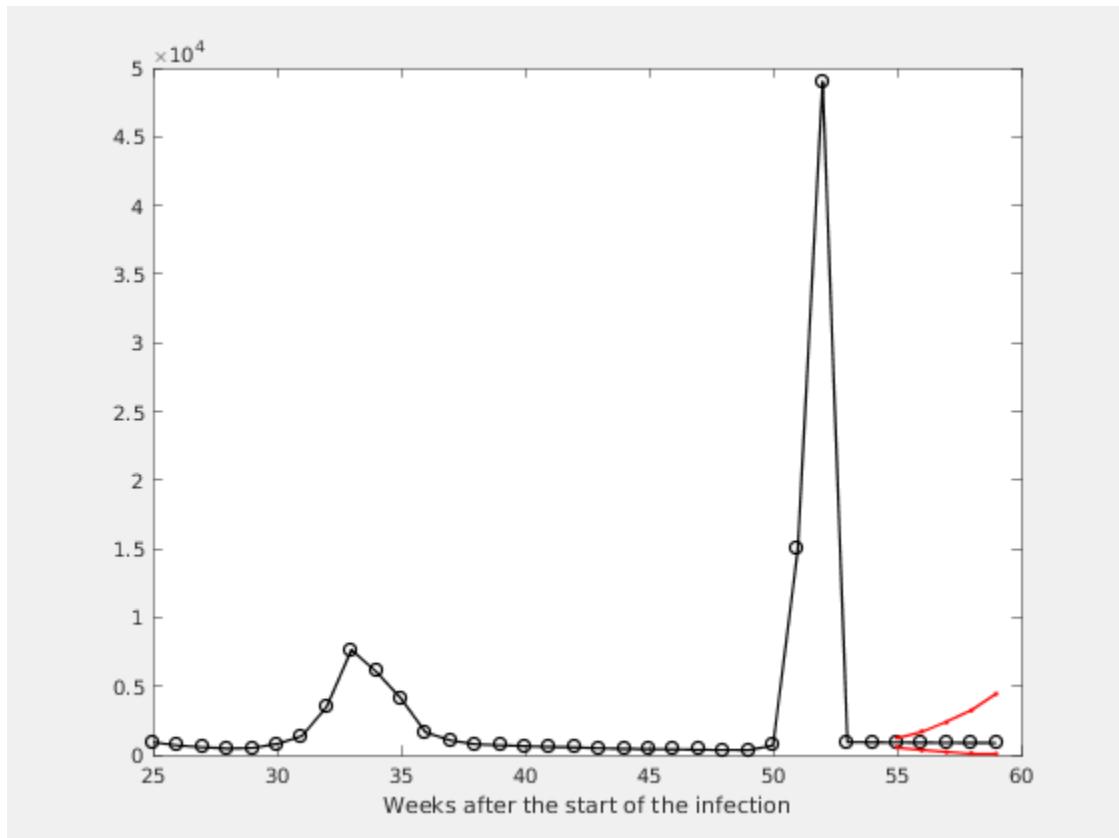
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



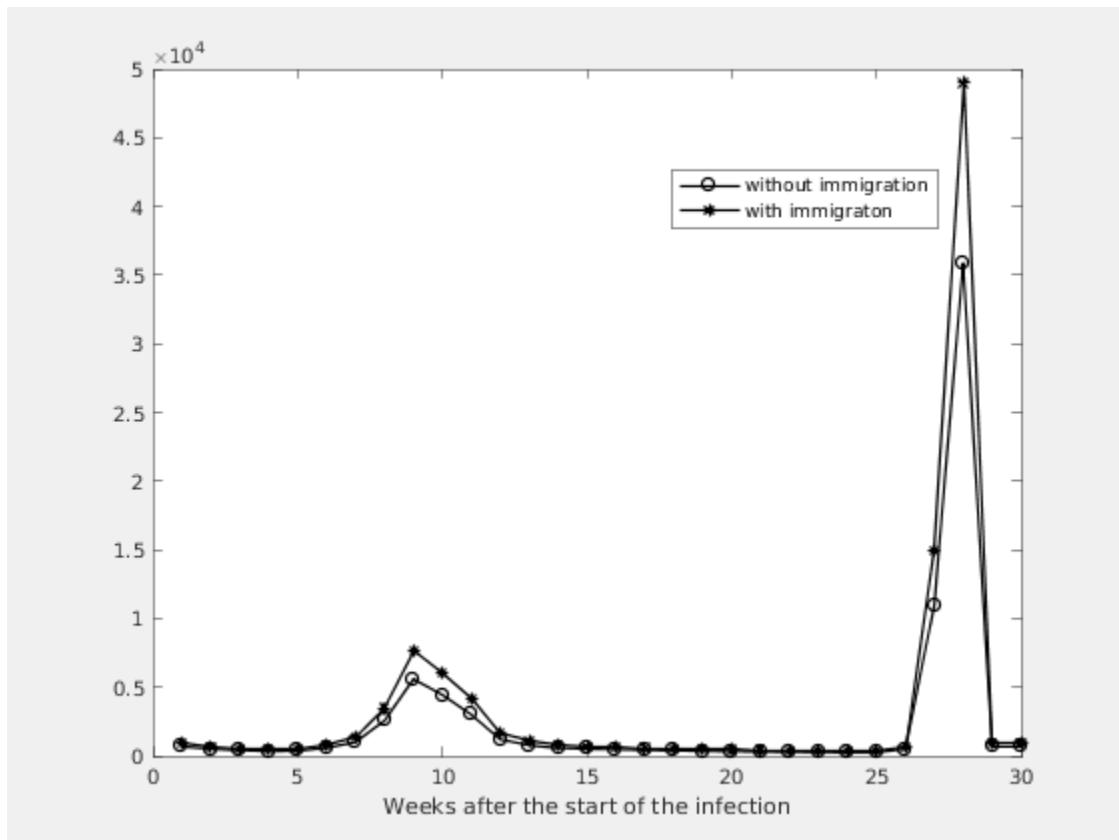
**Figure 2.2. Figure**



**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	
<hr/>						
4	0.9765	0.6214	- 1.3317	0.0405	355	260
3	1.0959	0.7420	- 1.4499	0.1283	347	254
2	1.1397	0.7368	- 1.5425	0.5871	679	497
1	0.9893	0.5492	- 1.4295	0.2558	14951	10942
0	0.9893	0.5799	- 1.3987	0.1311	48997	35859