

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

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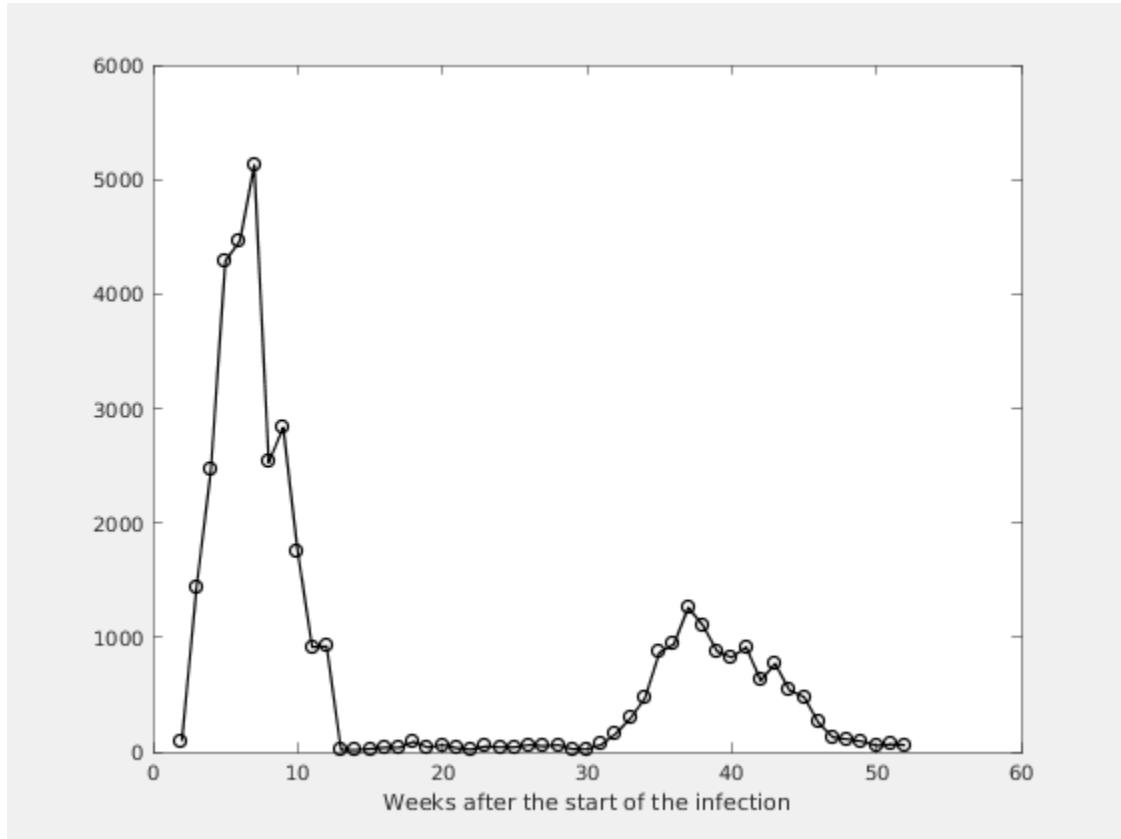
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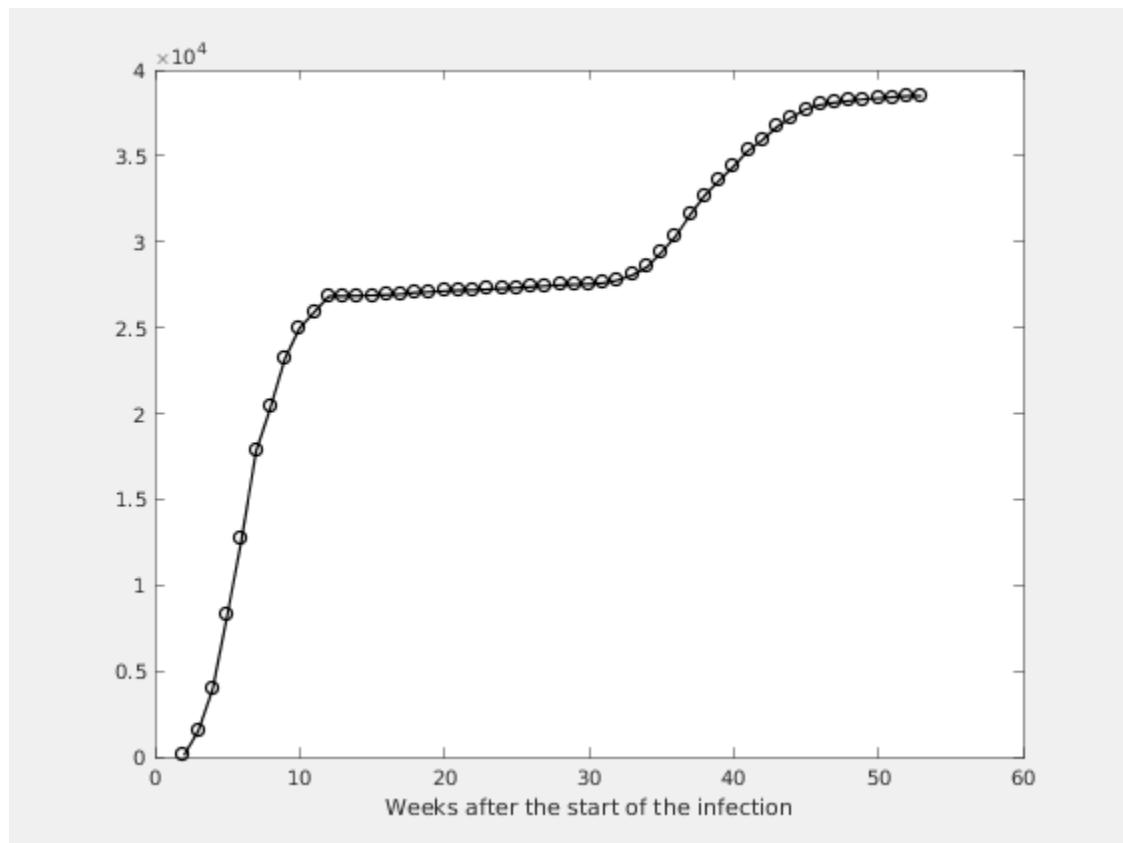
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# Chapter 1. Observed Infection data

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



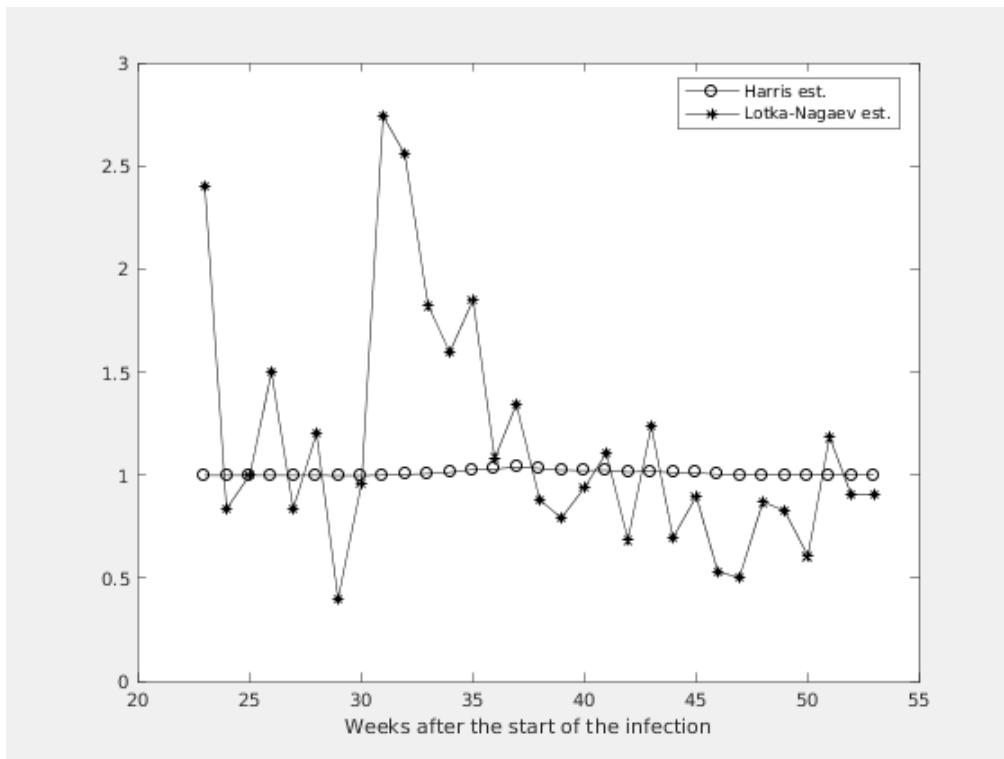
**Figure 1.2. Number of the total registered cases**



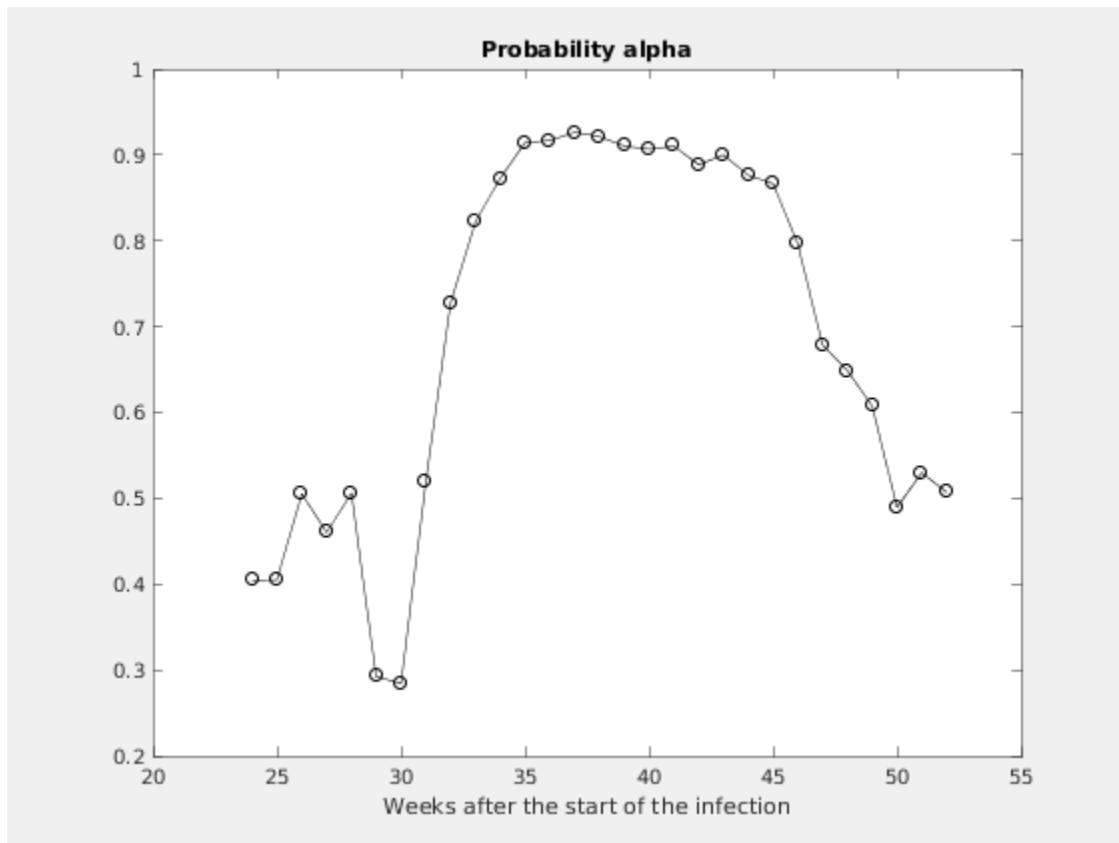
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# Chapter 2. Estimating of the main parameter and some predictions

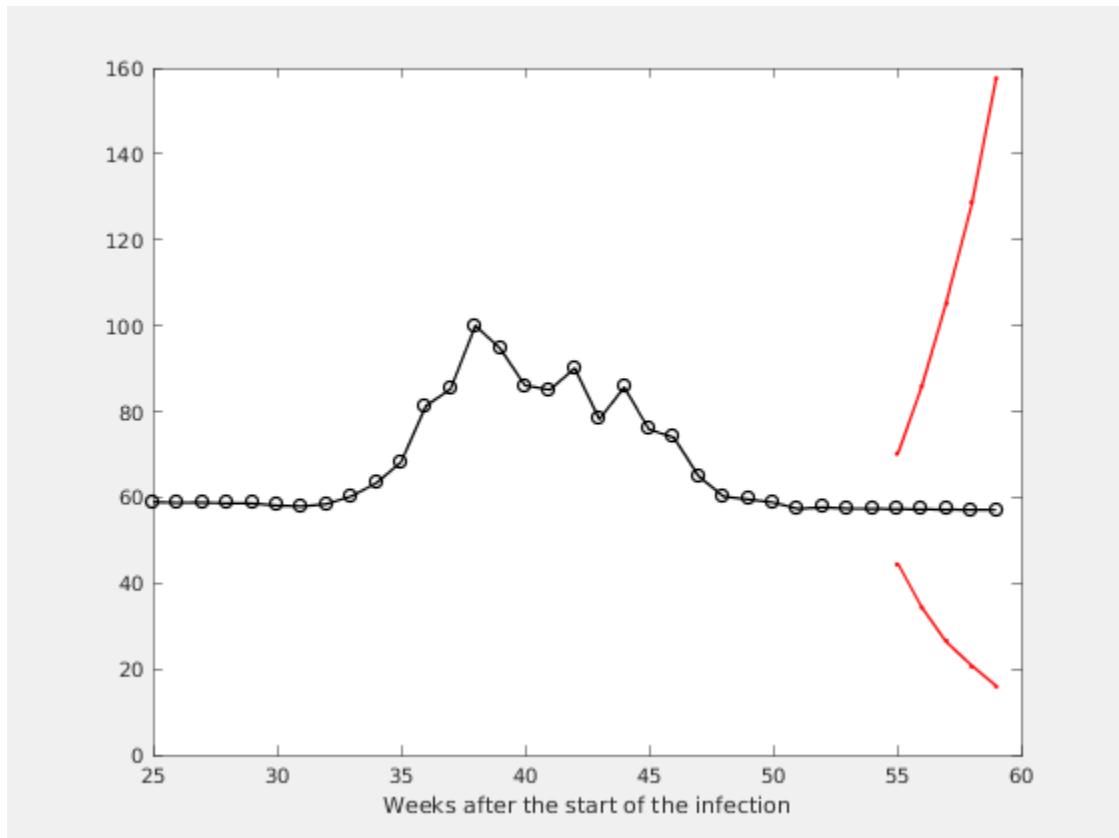
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



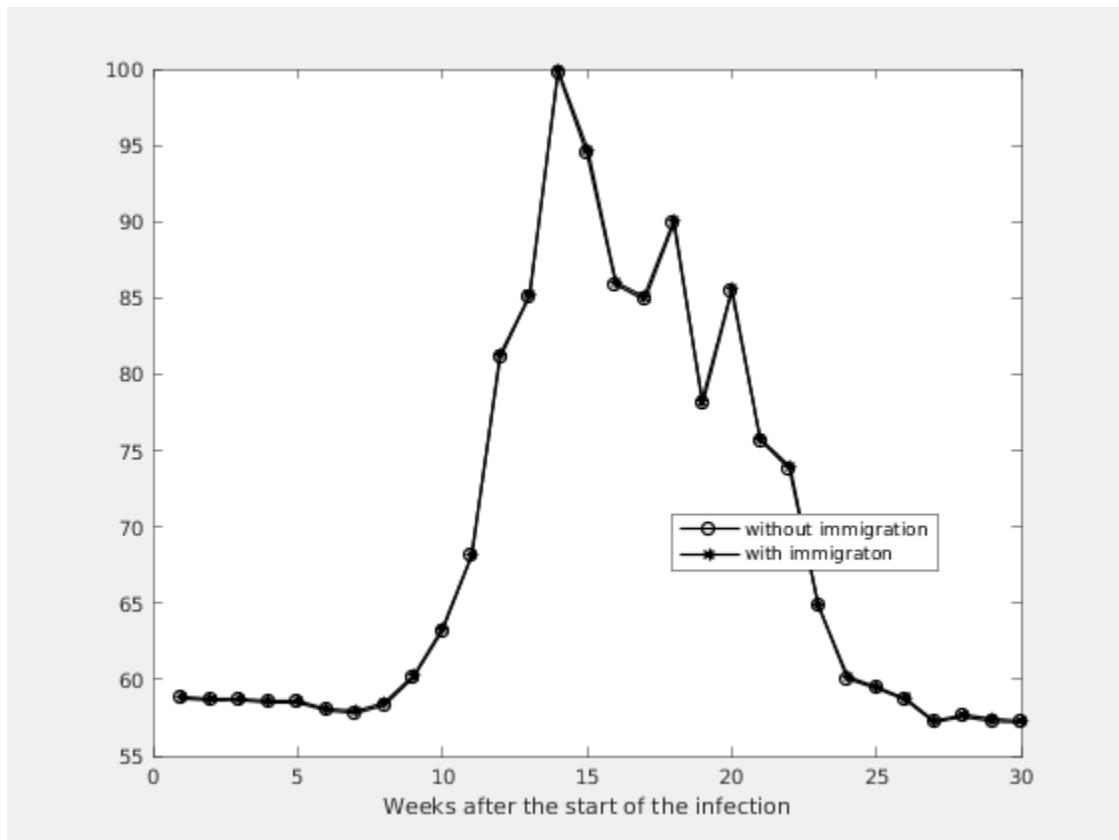
**Figure 2.2. Figure**



**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	0.9999	0.7612 - 1.2385	0.6785	60	60	
3	0.9989	0.7633 - 1.2346	0.6488	60	59	
2	0.9992	0.7664 - 1.2320	0.6075	59	59	
1	0.9990	0.7689 - 1.2292	0.4897	57	57	
0	0.9990	0.7714 - 1.2267	0.5298	58	58	