

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

**Guinea - week 53**

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## **Branching stochastic processes as models of Covid-19 epidemic development : Guinea - week 53**

### **Abstract**

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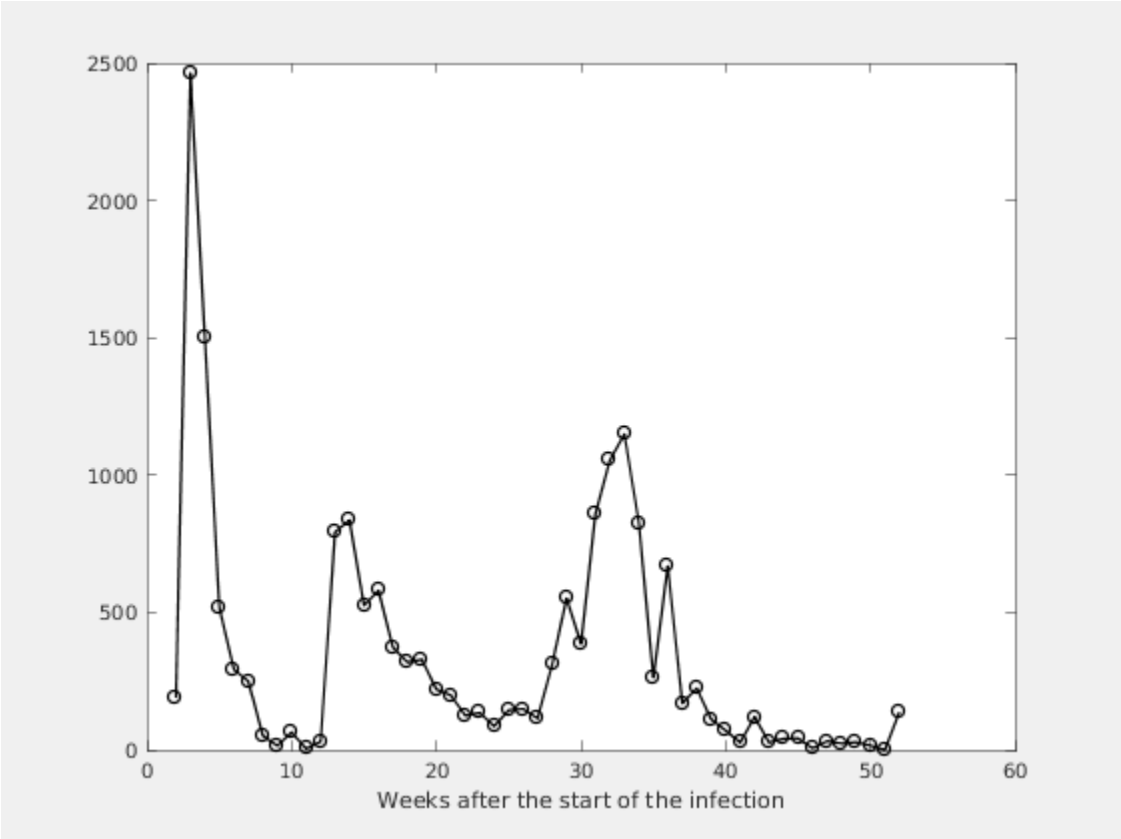
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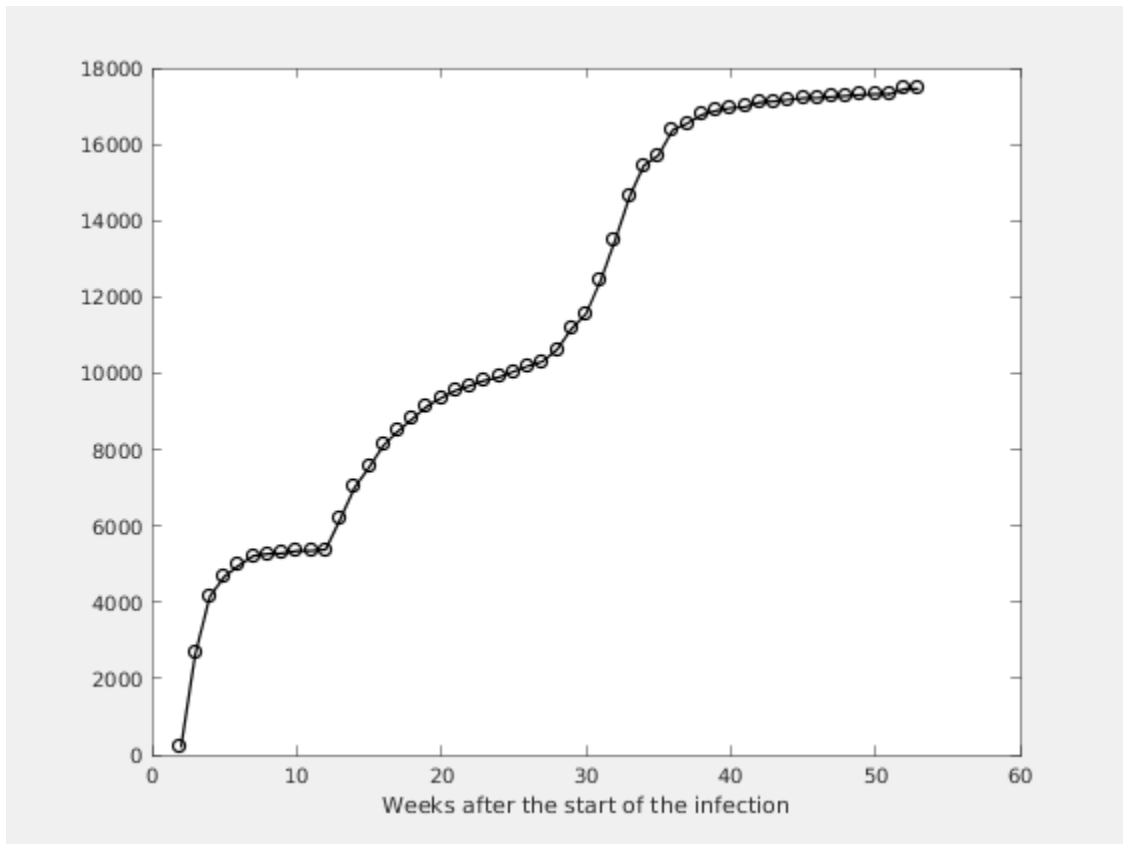
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# Chapter 1. Observed Infection data

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

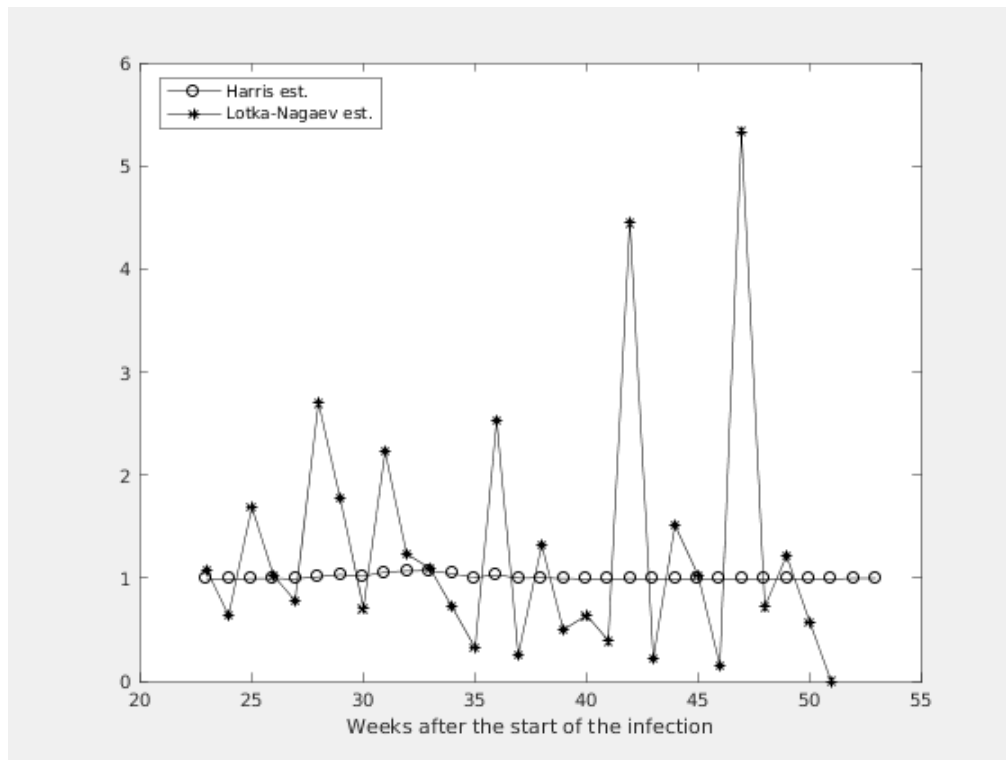
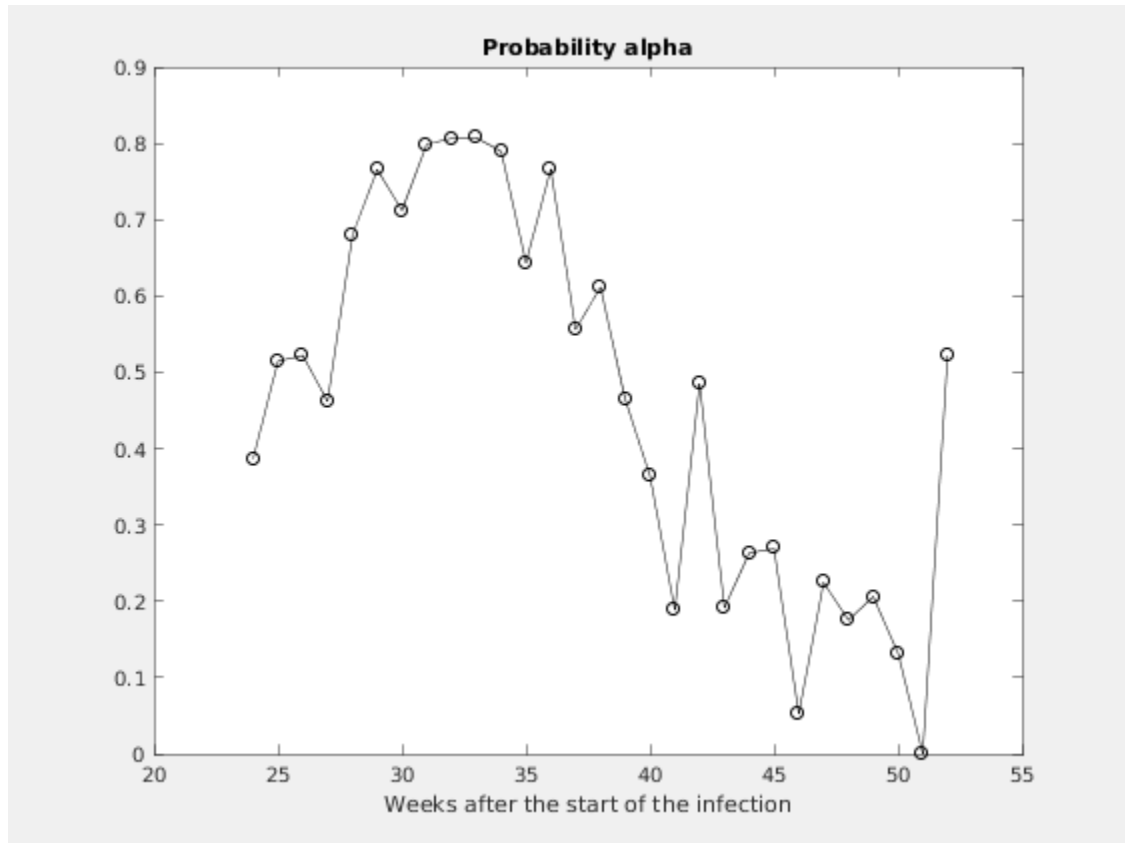
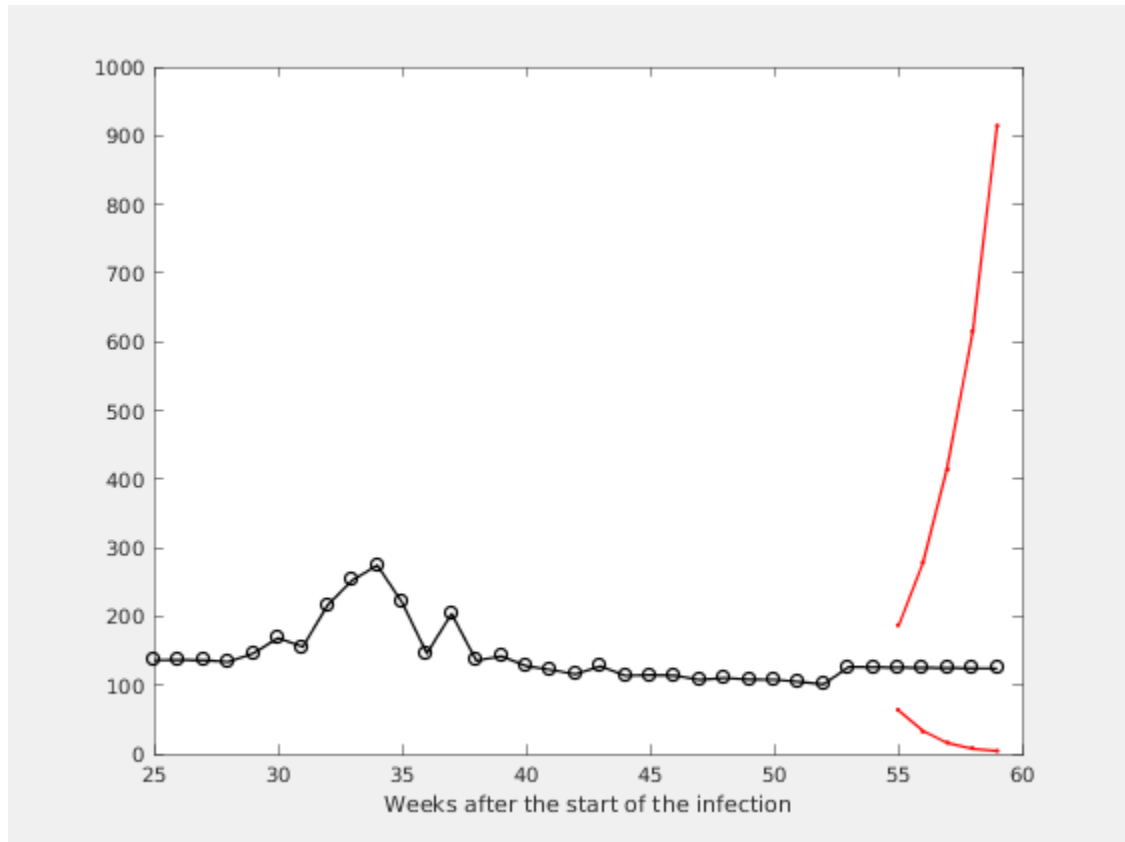


Figure 2.2. Figure

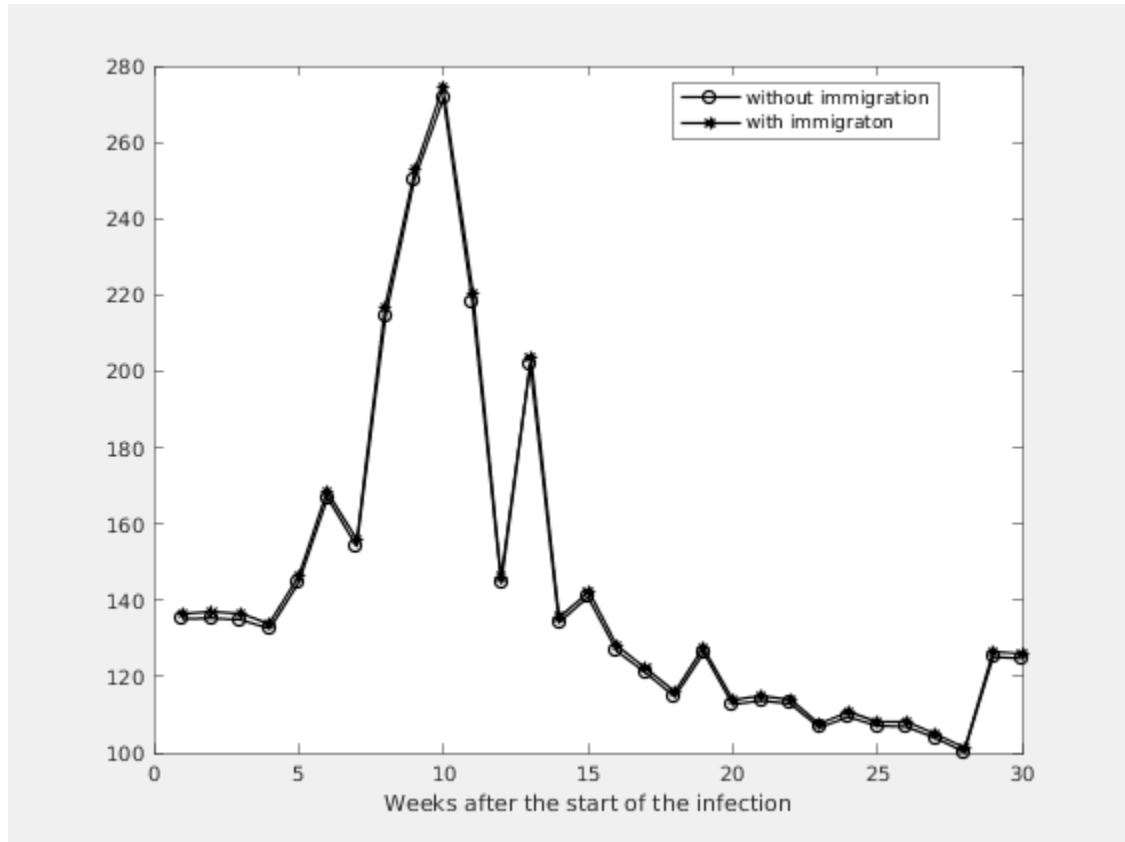




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

k	m	ci	alpha	A1	M1
4	0.9906	0.4831 - 1.4981	0.2244	111	109
3	0.9899	0.4884 - 1.4915	0.1753	108	107
2	0.9890	0.4932 - 1.4849	0.2059	108	107
1	0.9970	0.5067 - 1.4873	0.1322	105	104
0	0.9970	0.5070 - 1.4870	0.0000	101	100