

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

**CaboVerde - week 53**

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## **Branching stochastic processes as models of Covid-19 epidemic development : CaboVerde - 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 CaboVerde. 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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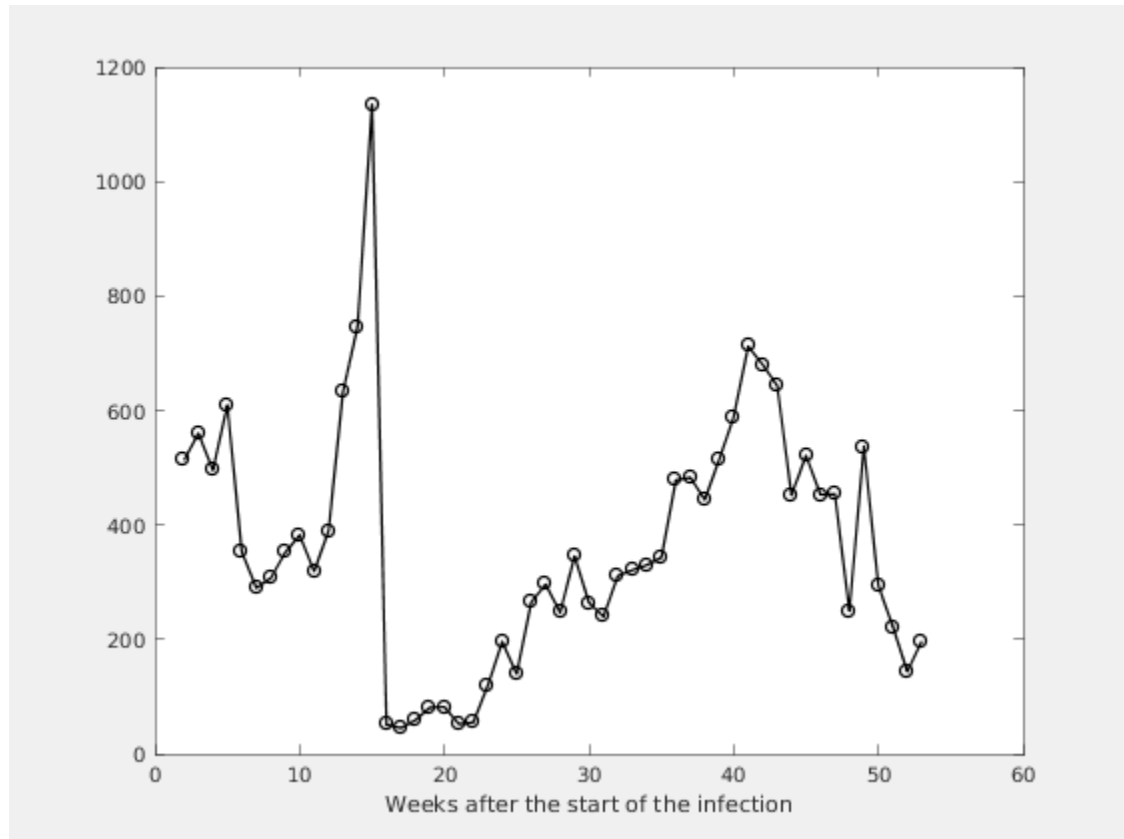
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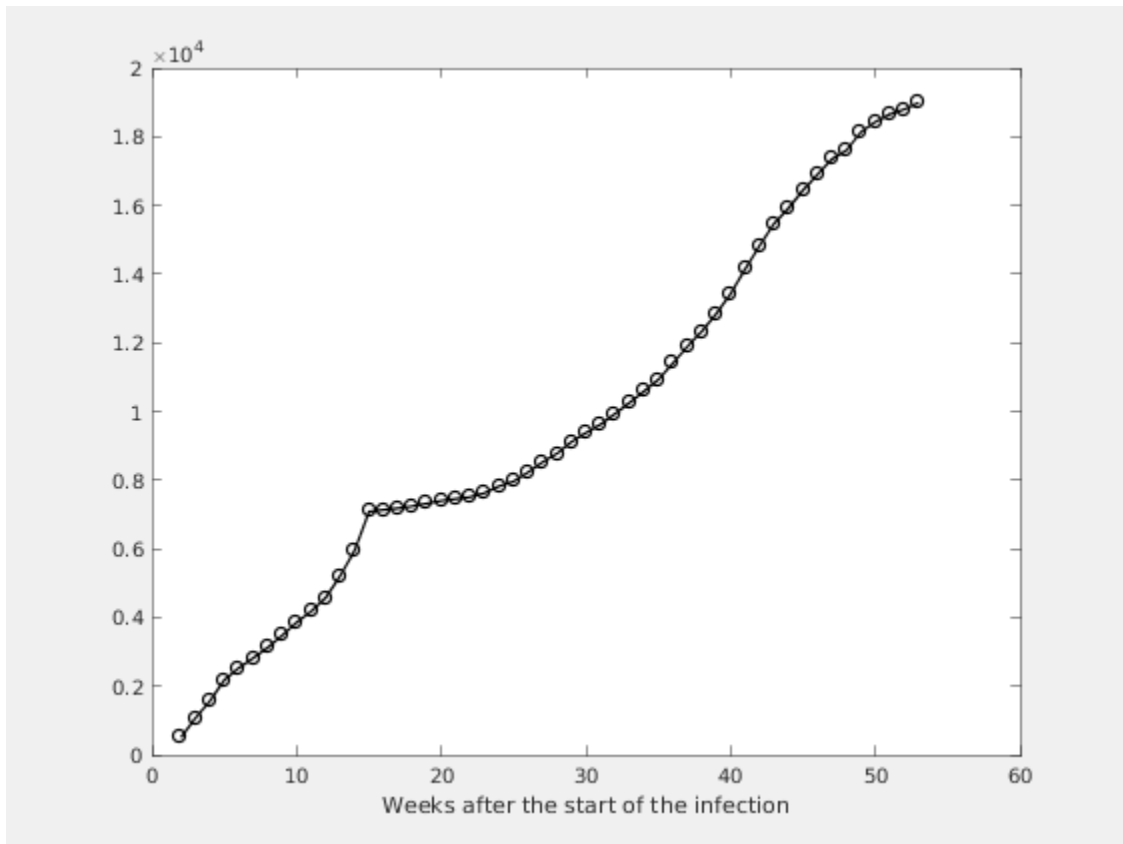
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# Chapter 1. Observed Infection data

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

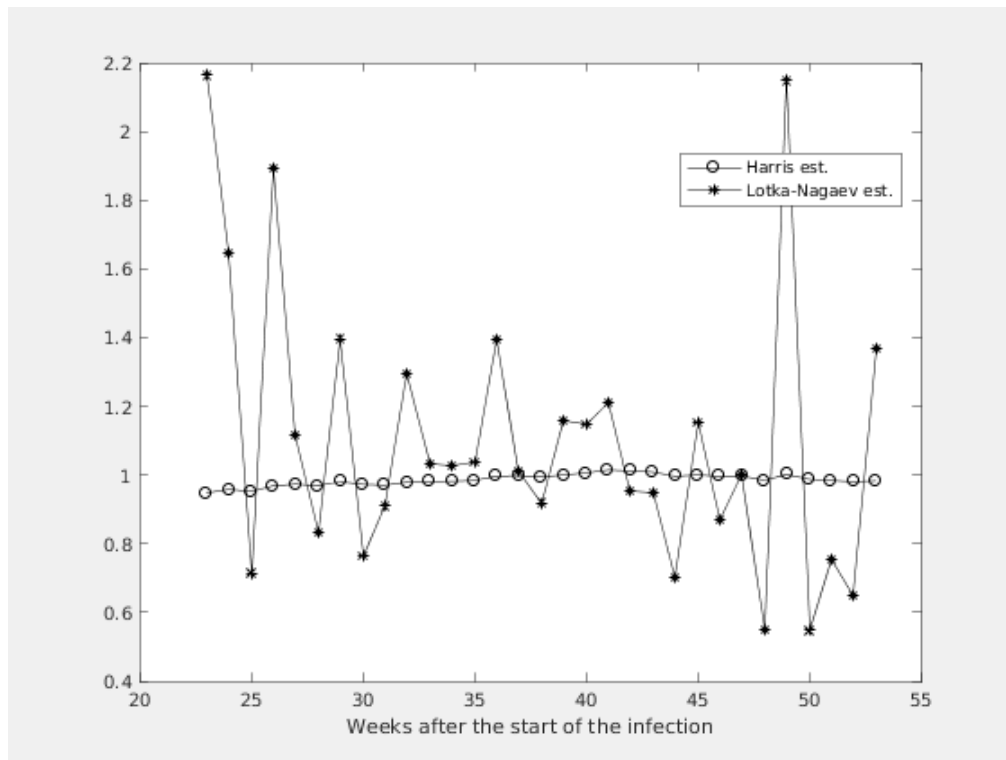
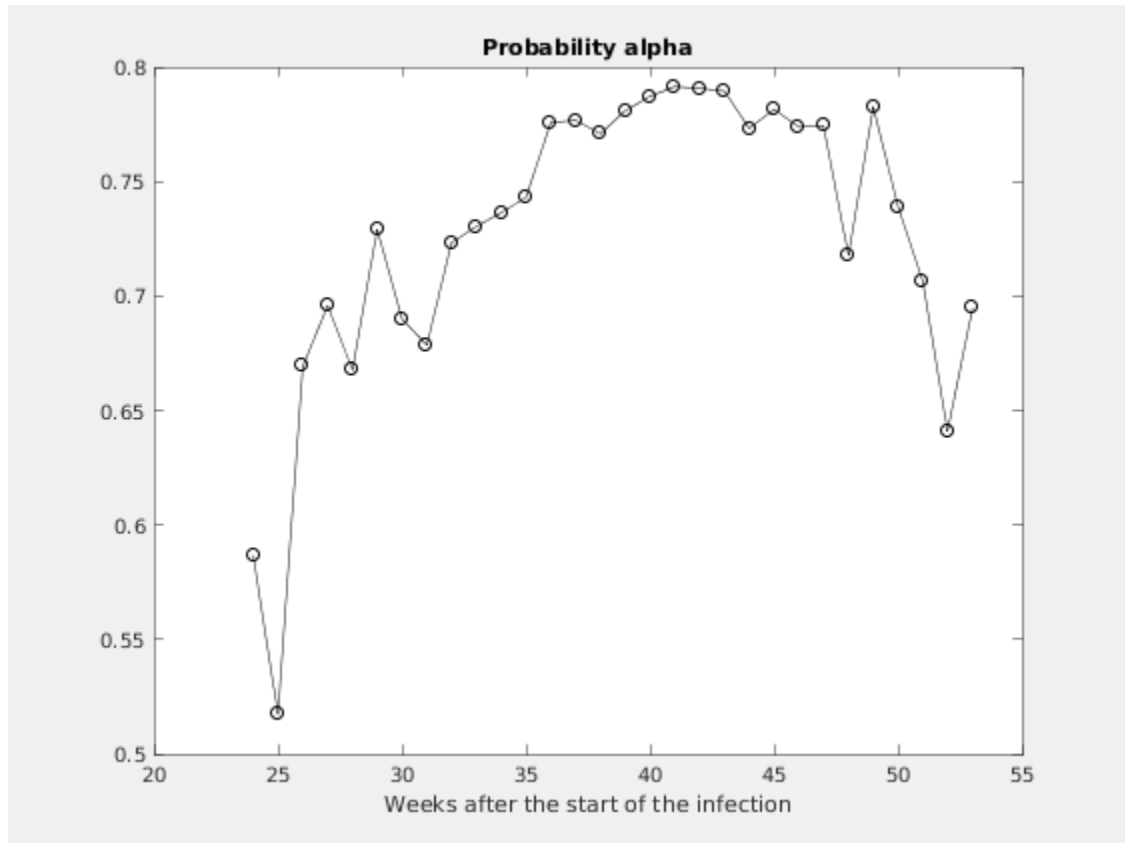
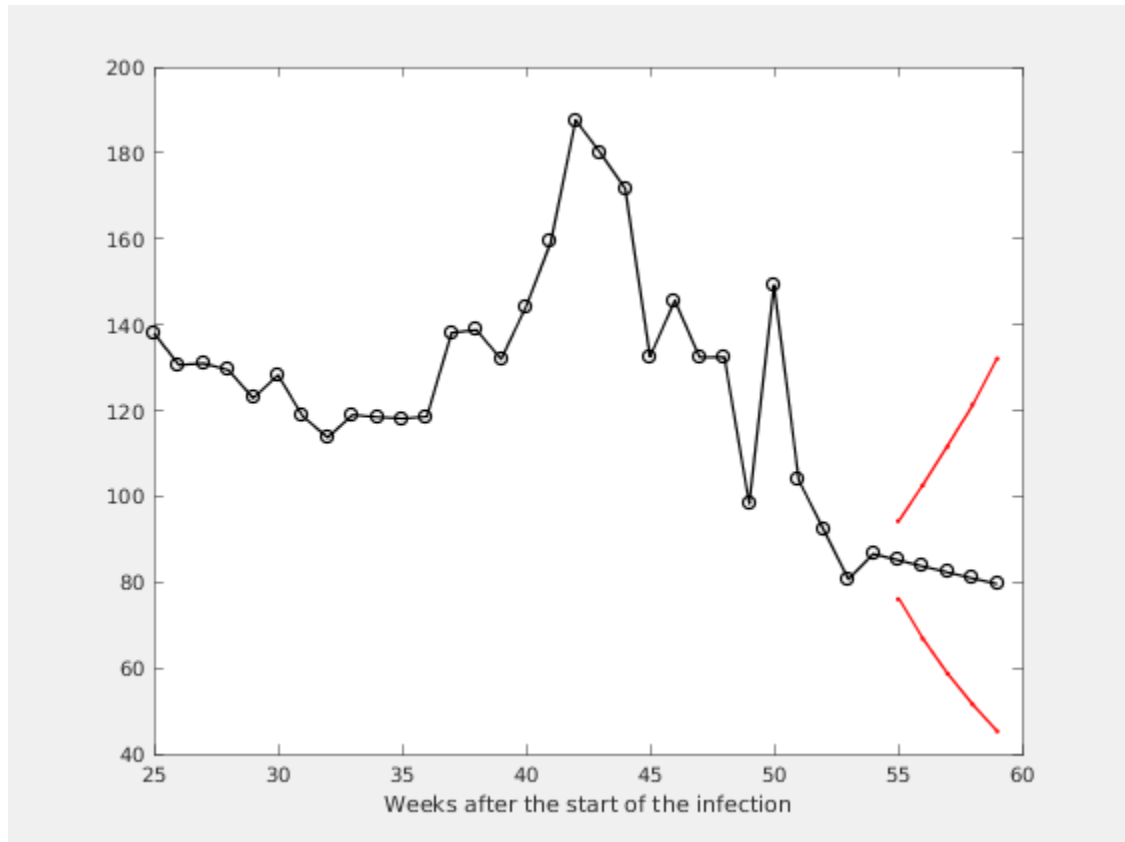


Figure 2.2. Figure

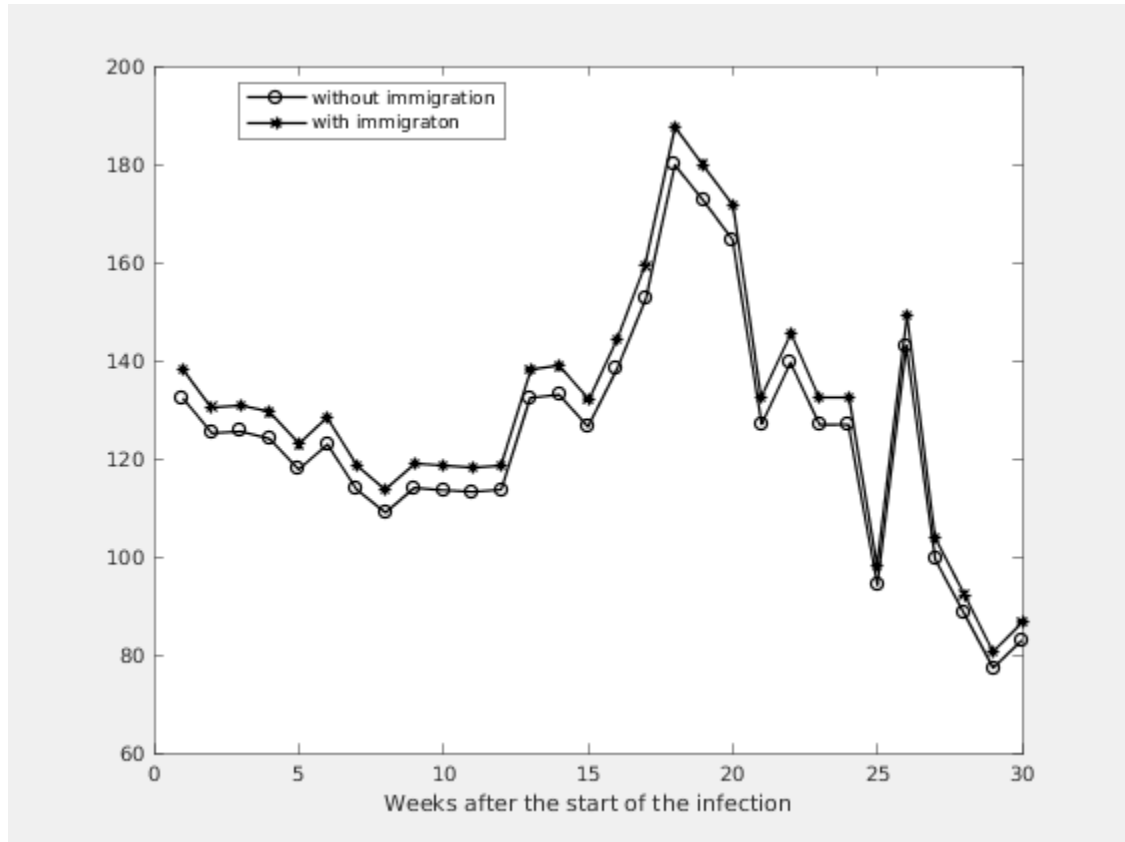




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

k	m	ci	alpha	A1	M1
4	1.0014	0.8970 - 1.1057	0.7742	132	127
3	0.9879	0.8840 - 1.0918	0.7178	98	94
2	0.9842	0.8747 - 1.0937	0.7826	149	143
1	0.9802	0.8714 - 1.0891	0.7392	104	99
0	0.9832	0.8760 - 1.0903	0.7065	92	88