

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

**Var174 - week 53**

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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 Var174. 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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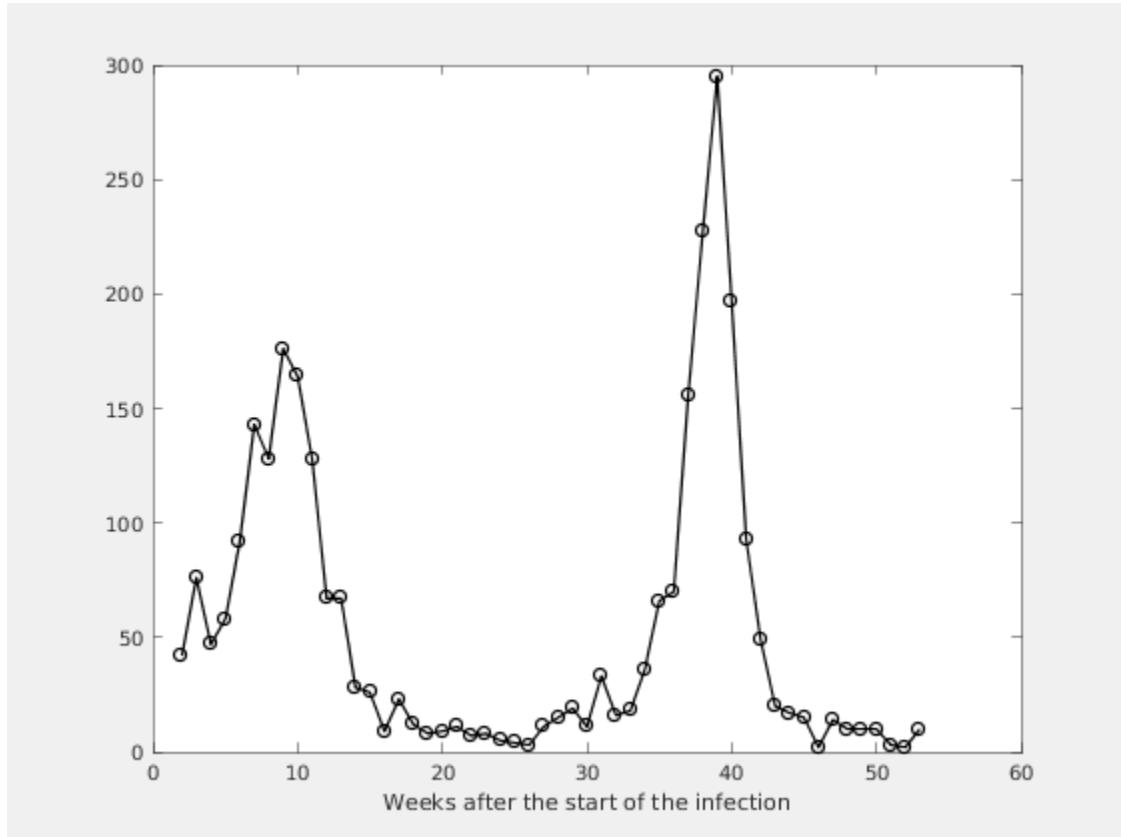
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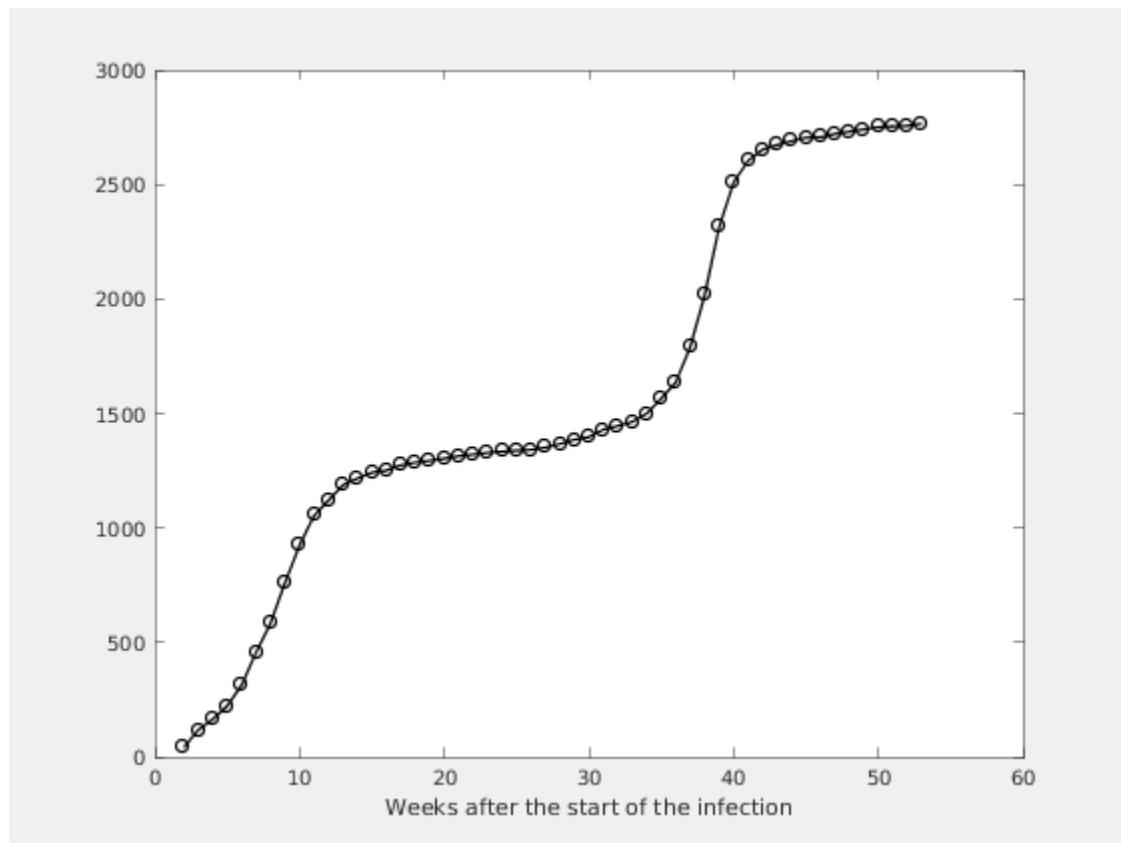
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# Chapter 1. Observed Infection data

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



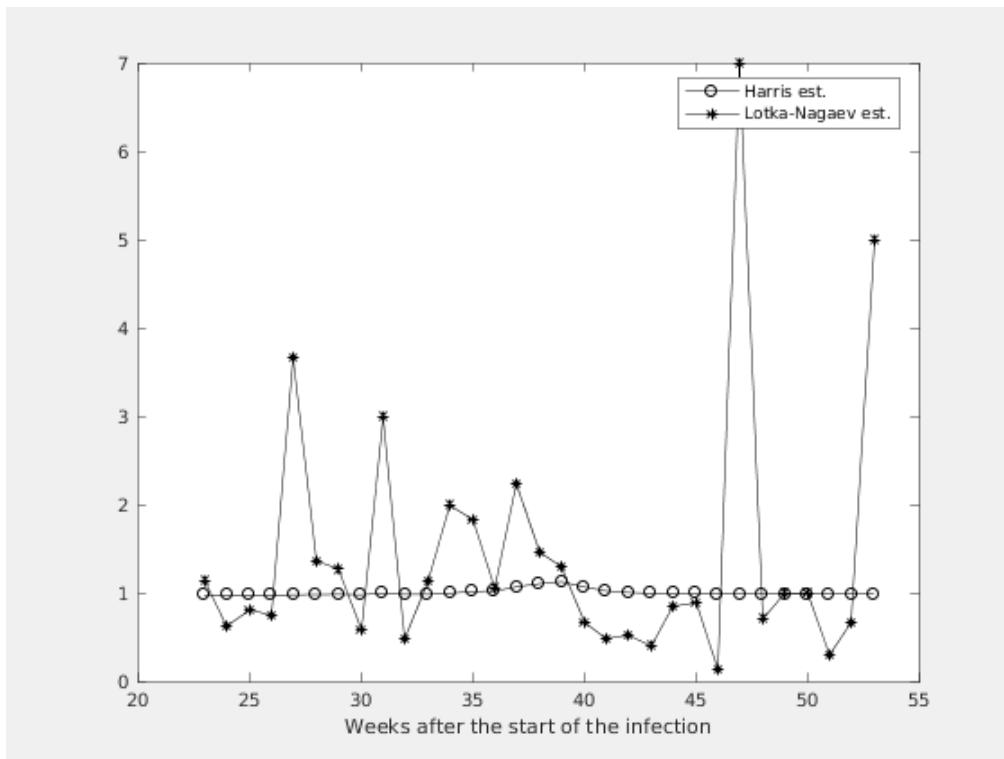
**Figure 1.2. Number of the total registered cases**



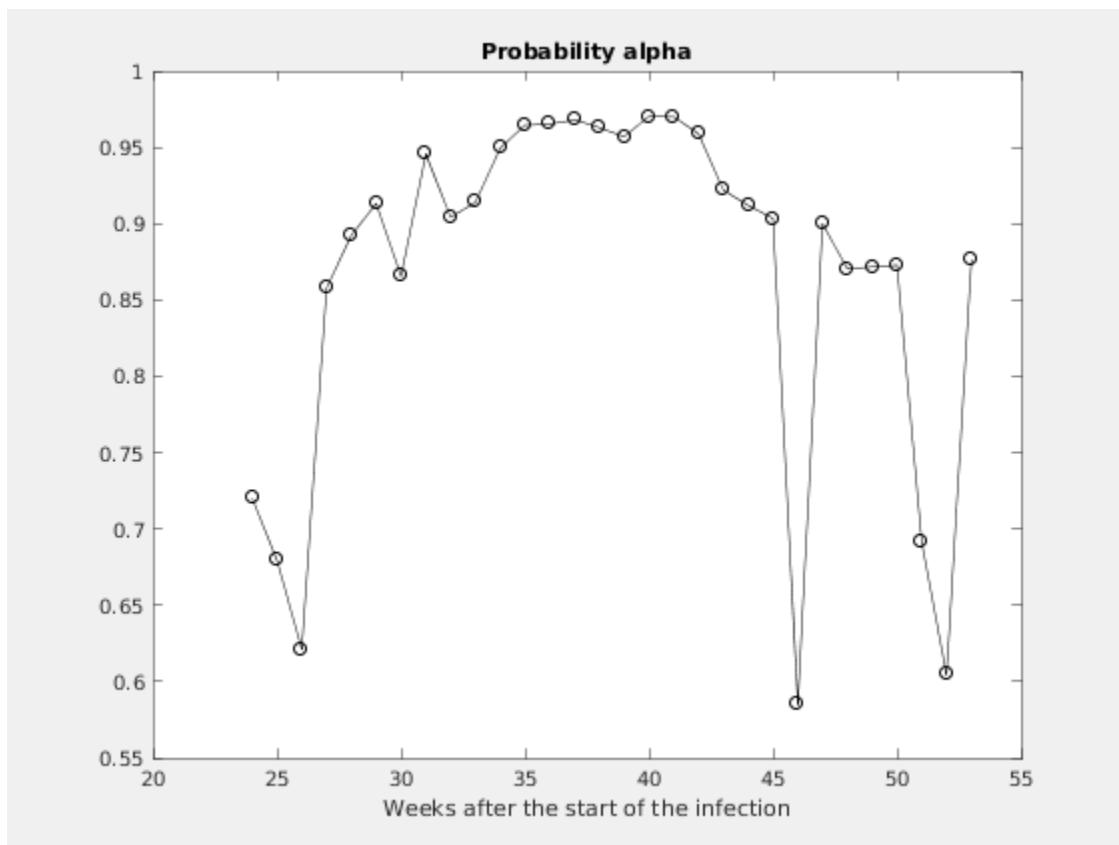
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# Chapter 2. Estimating of the main parameter and some predictions

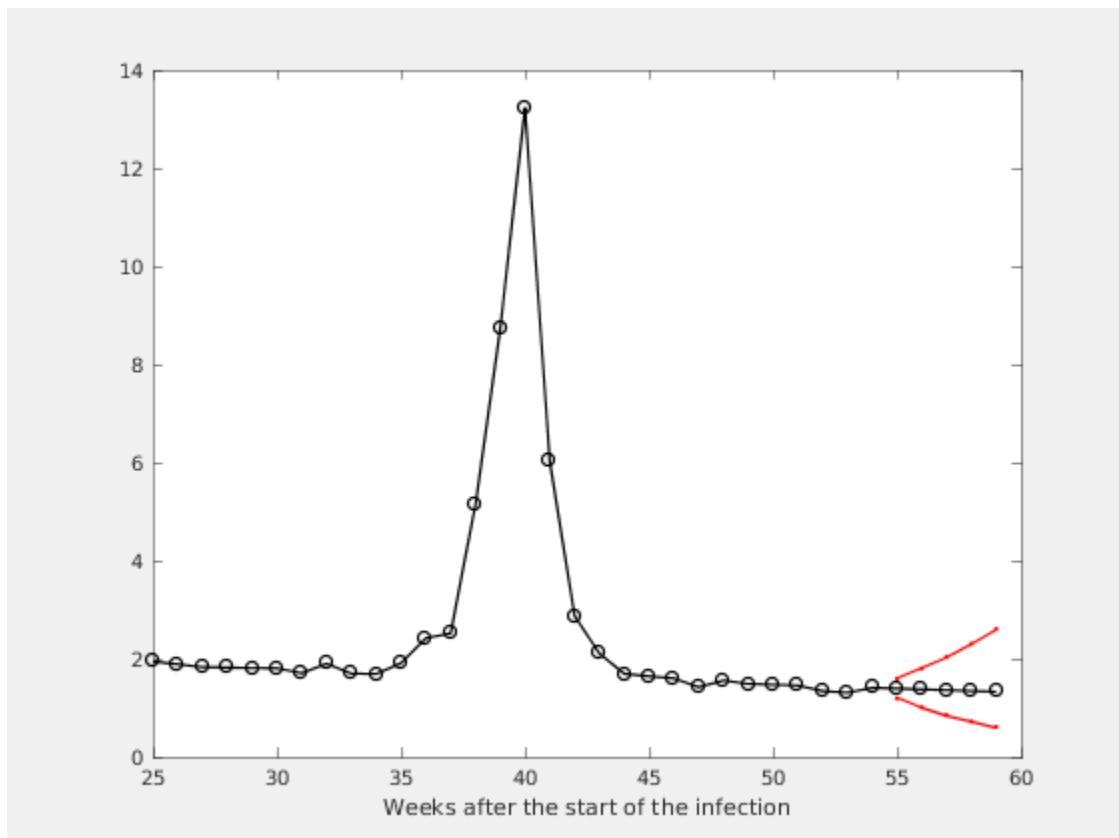
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



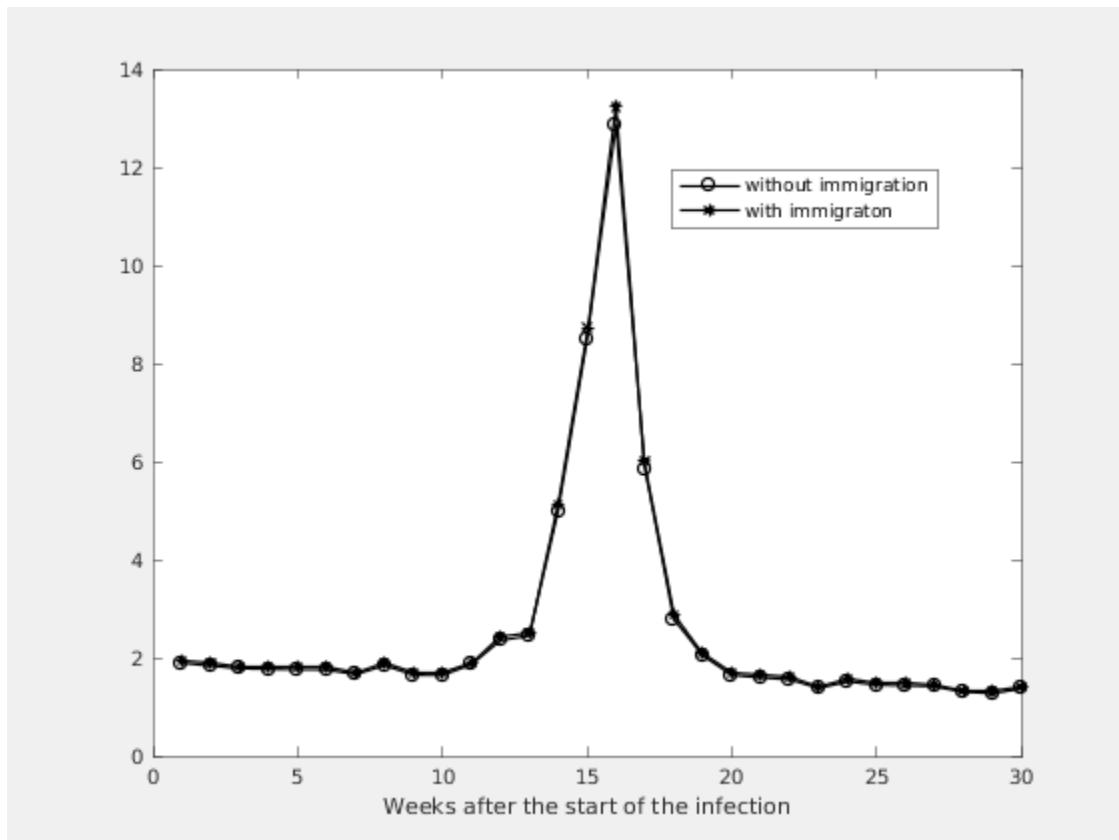
**Figure 2.2. Figure**



**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	0.9883	0.8393 - 1.1373	0.8998	2	2	
3	0.9883	0.8413 - 1.1354	0.8705	1	1	
2	0.9858	0.8406 - 1.1311	0.8717	1	1	
1	0.9855	0.8420 - 1.1289	0.8729	1	1	
0	0.9884	0.8462 - 1.1306	0.6911	1	1	