

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

**Var213 - 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 Var213. 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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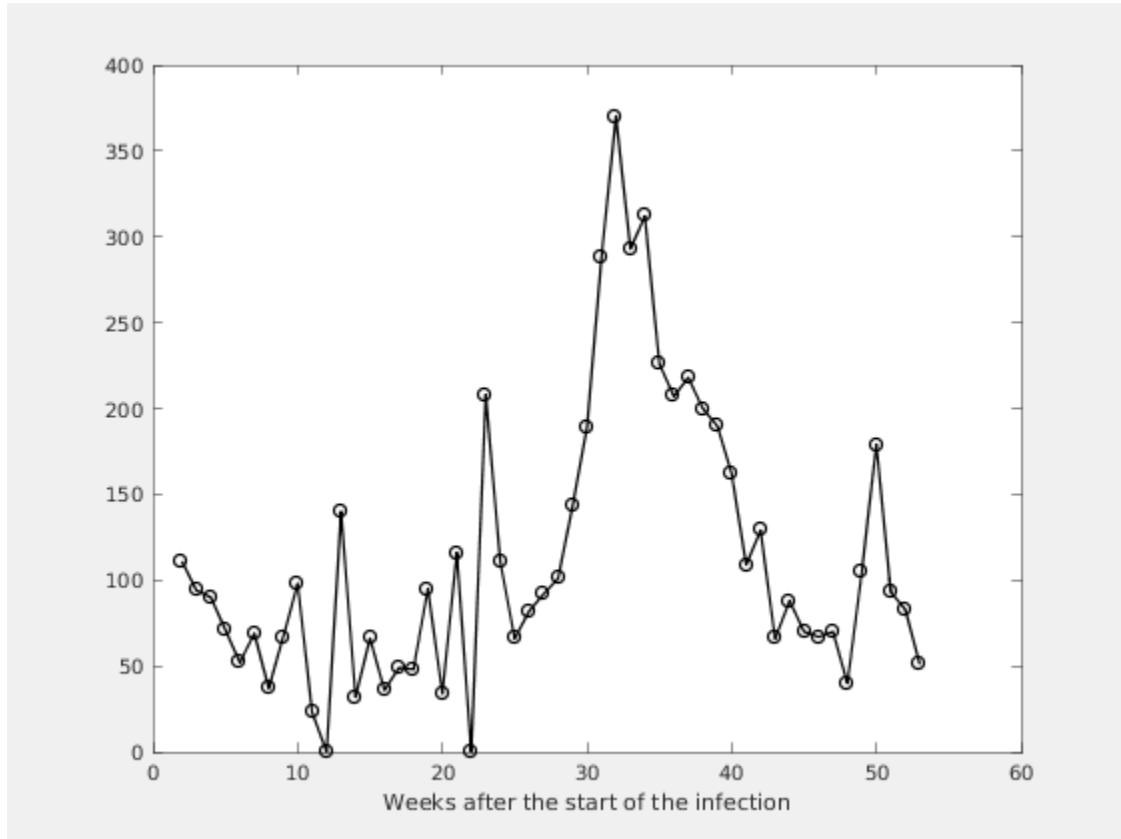
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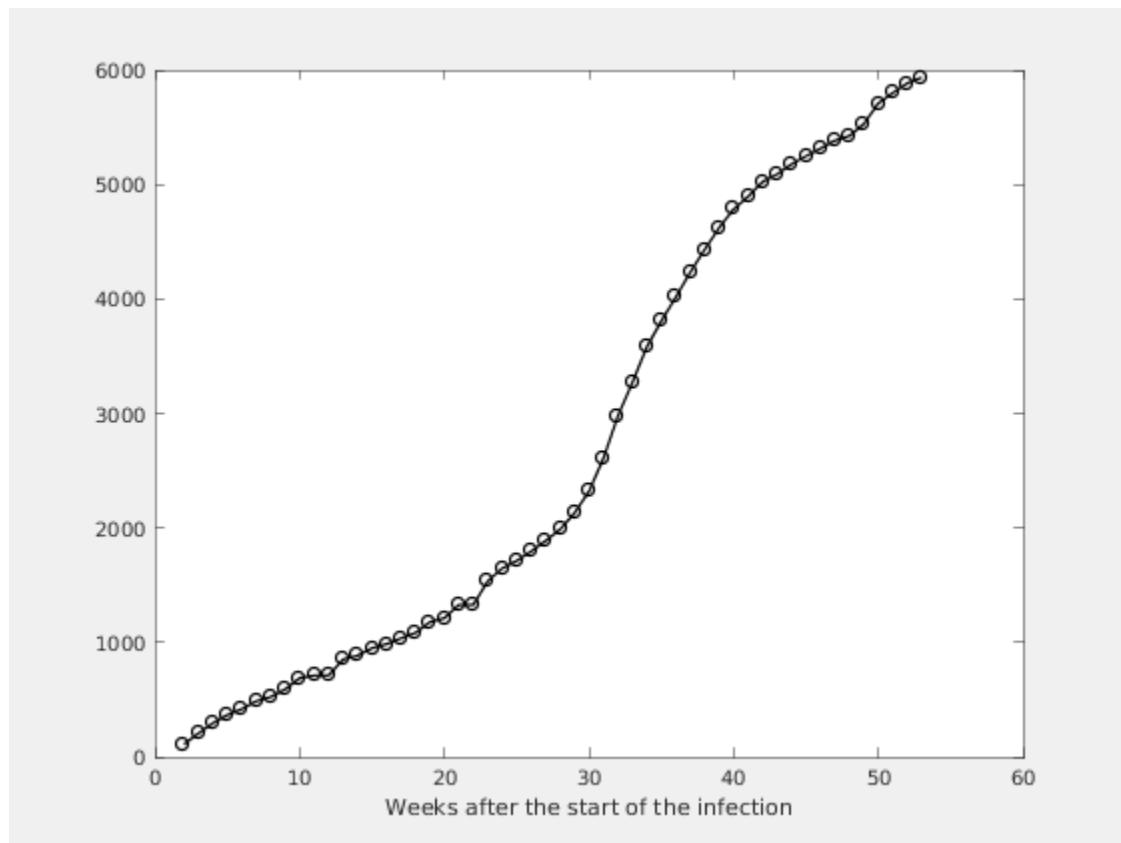
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# Chapter 1. Observed Infection data

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



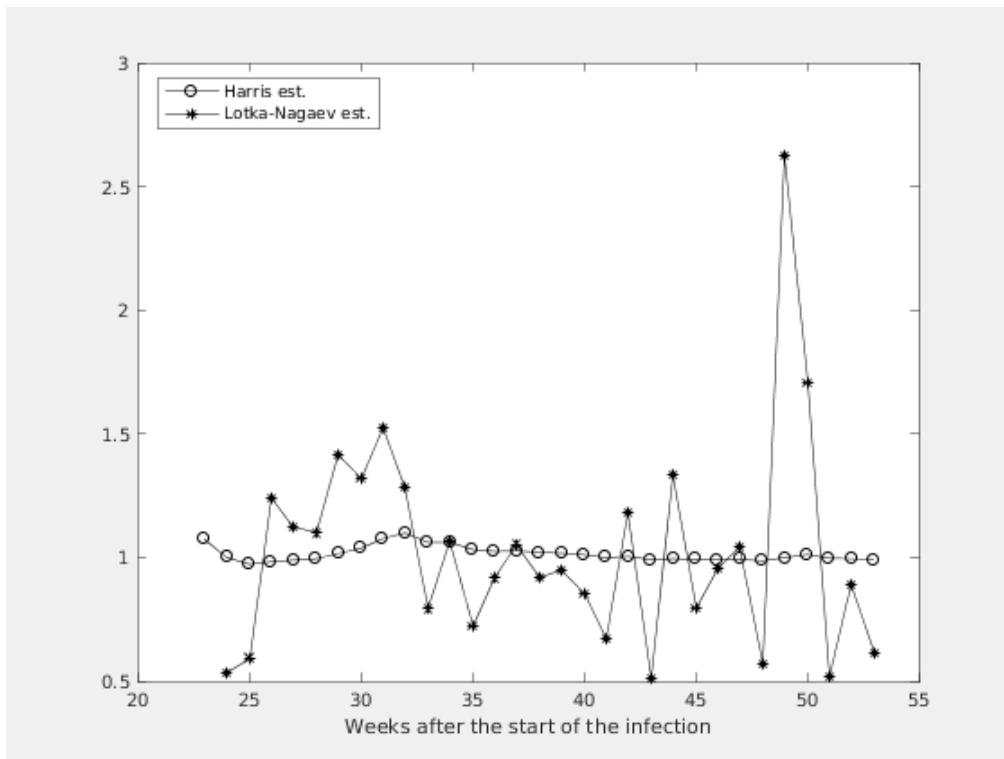
**Figure 1.2. Number of the total registered cases**



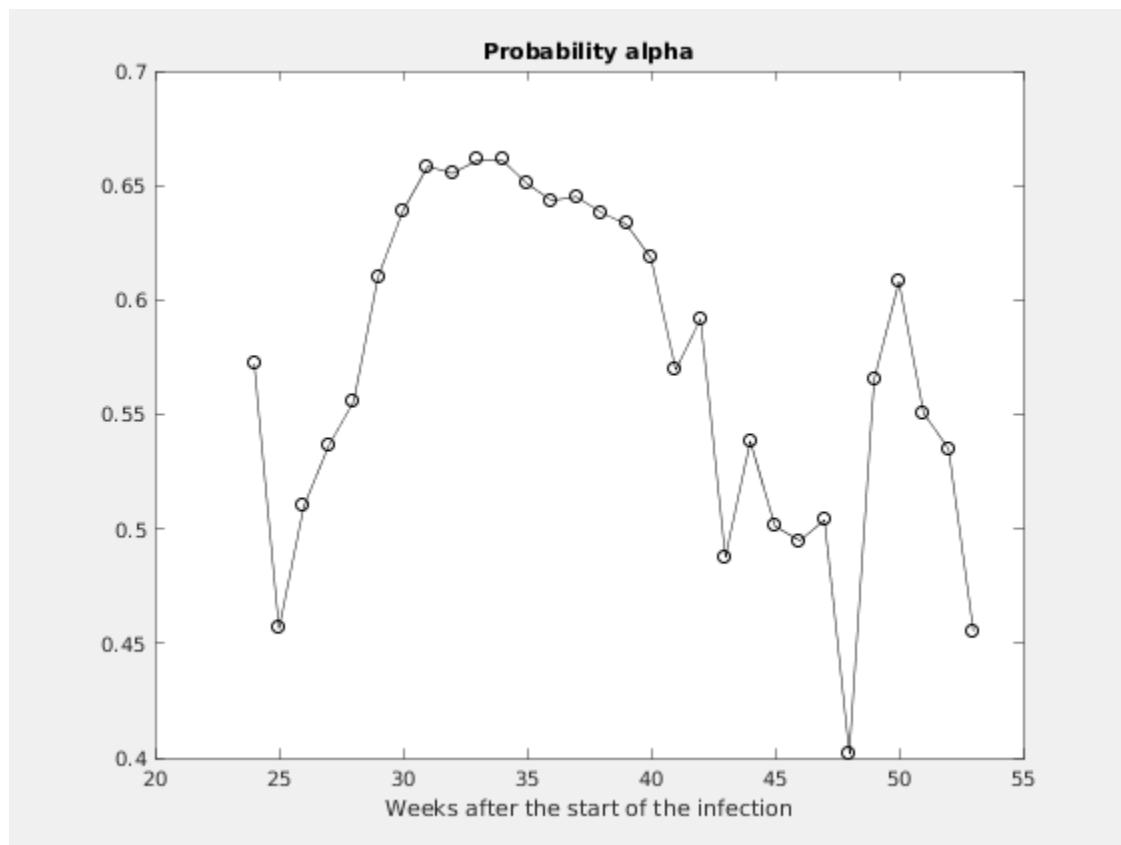
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# Chapter 2. Estimating of the main parameter and some predictions

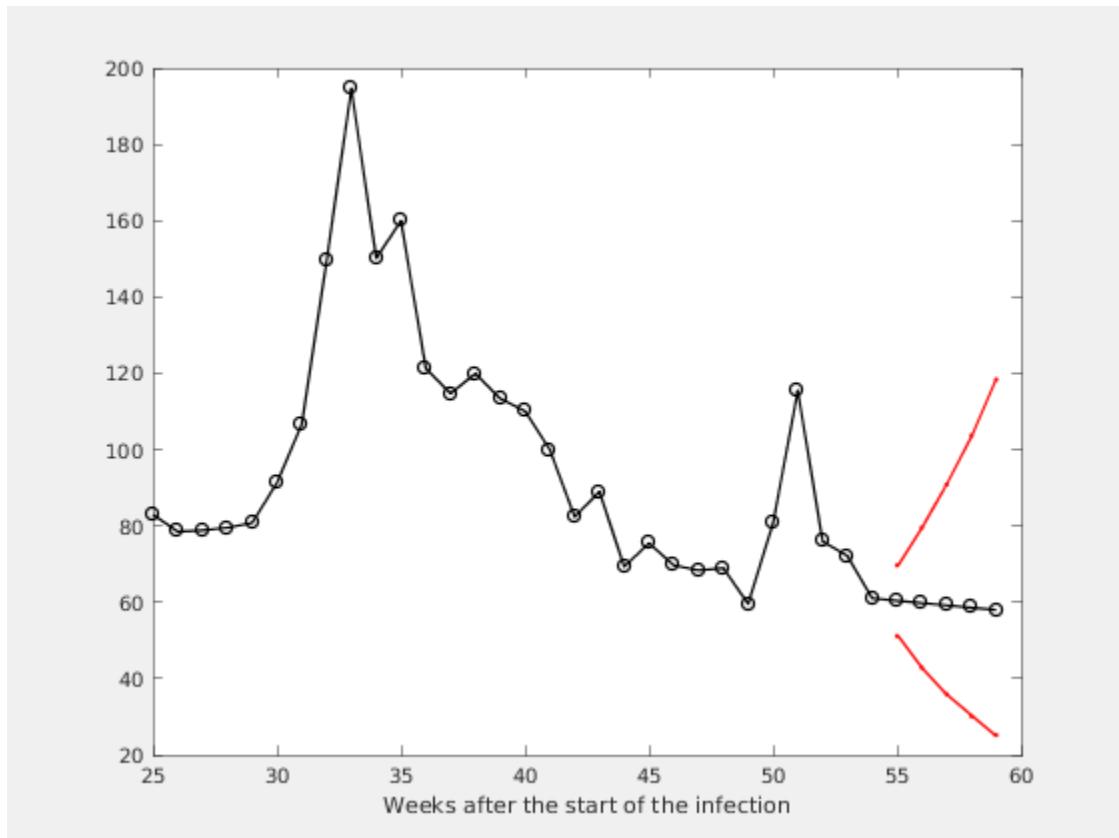
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



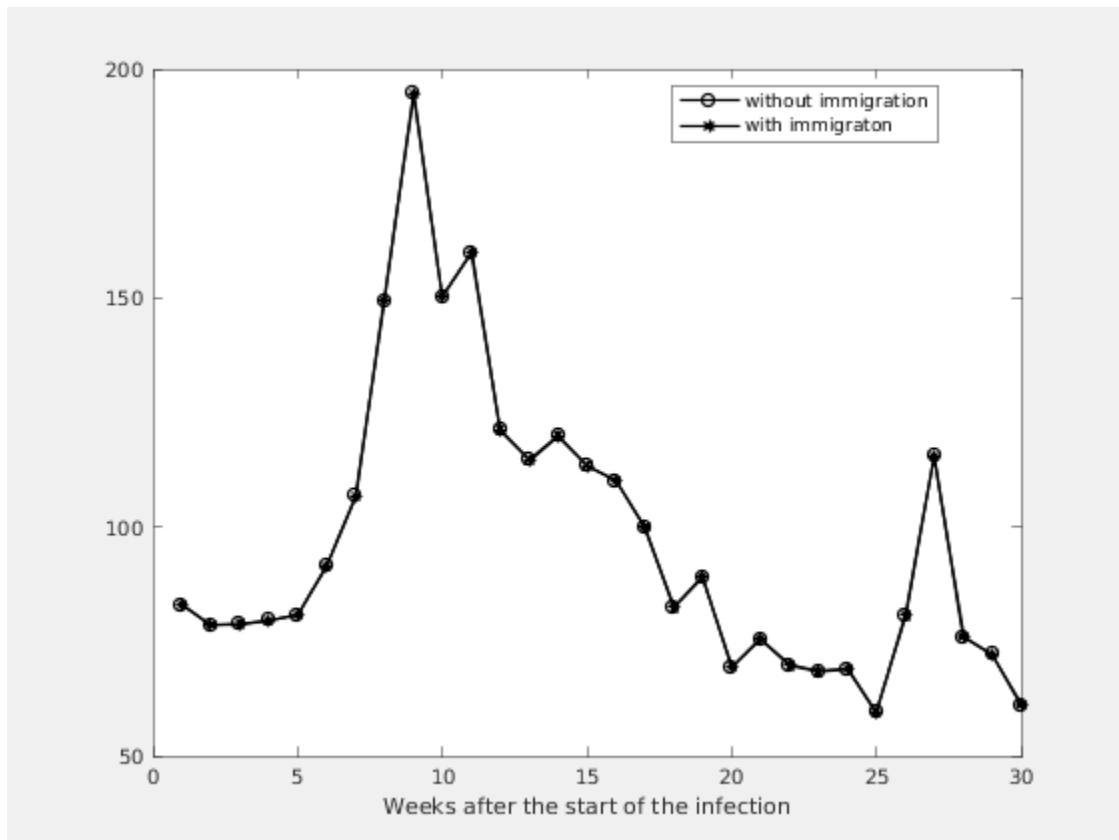
**Figure 2.2. Figure**



**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	0.9989	0.8398	- 1.1580	0.5039	69	69
3	1.0123	0.8552	- 1.1694	0.4018	60	60
2	0.9968	0.8369	- 1.1568	0.5656	81	81
1	0.9952	0.8360	- 1.1544	0.6079	115	115
0	0.9898	0.8329	- 1.1467	0.5504	76	76