

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

**Var206 - week 53**

**N. Yanev, V. Stoimenova, D. Atanasov**

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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 Var206. 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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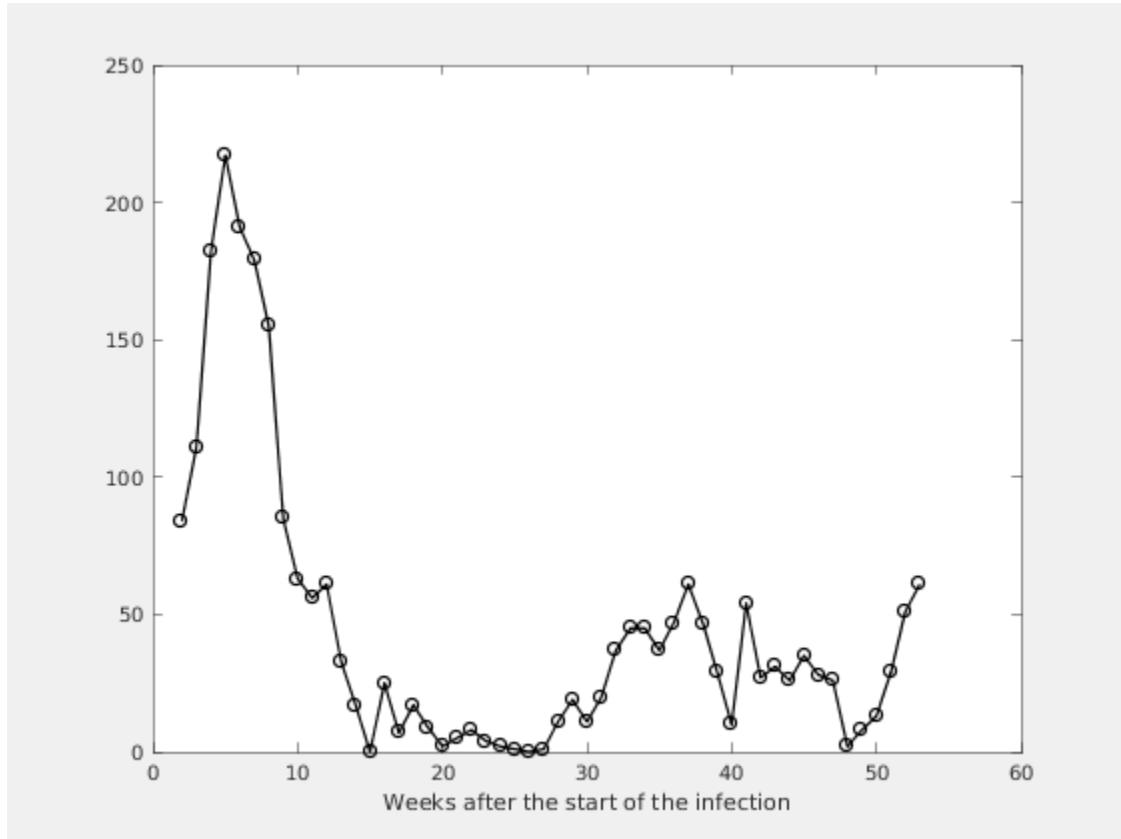
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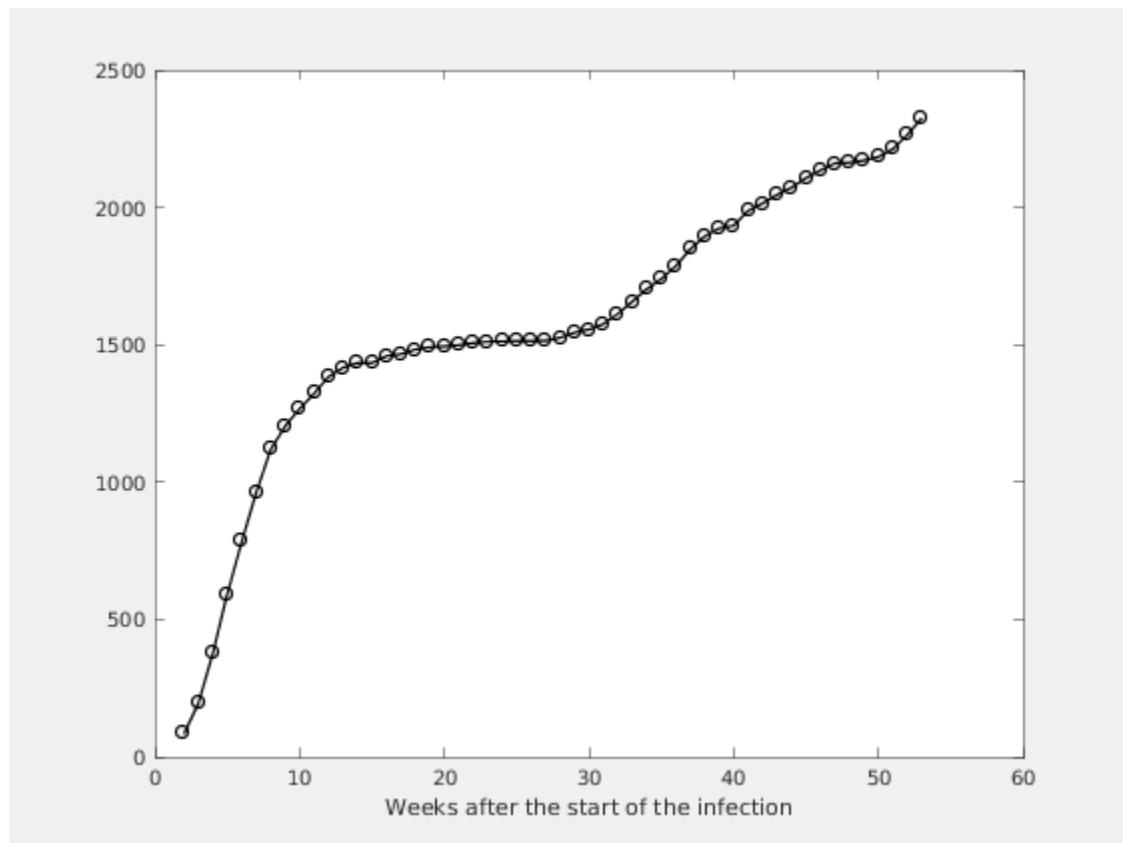
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# Chapter 1. Observed Infection data

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



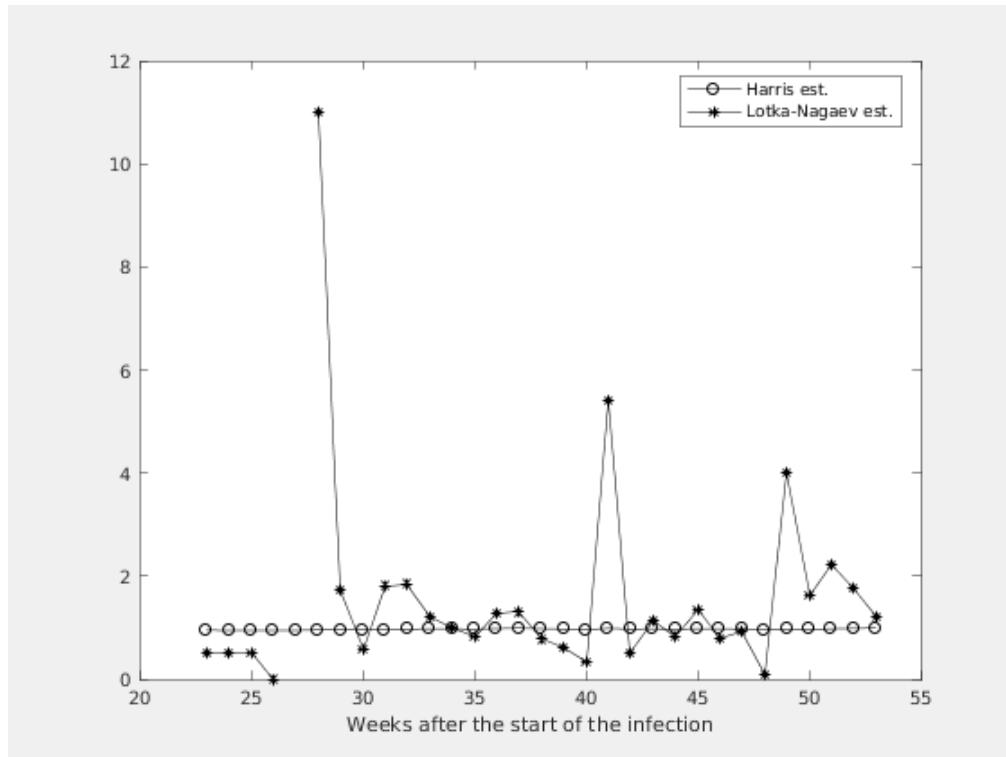
**Figure 1.2. Number of the total registered cases**



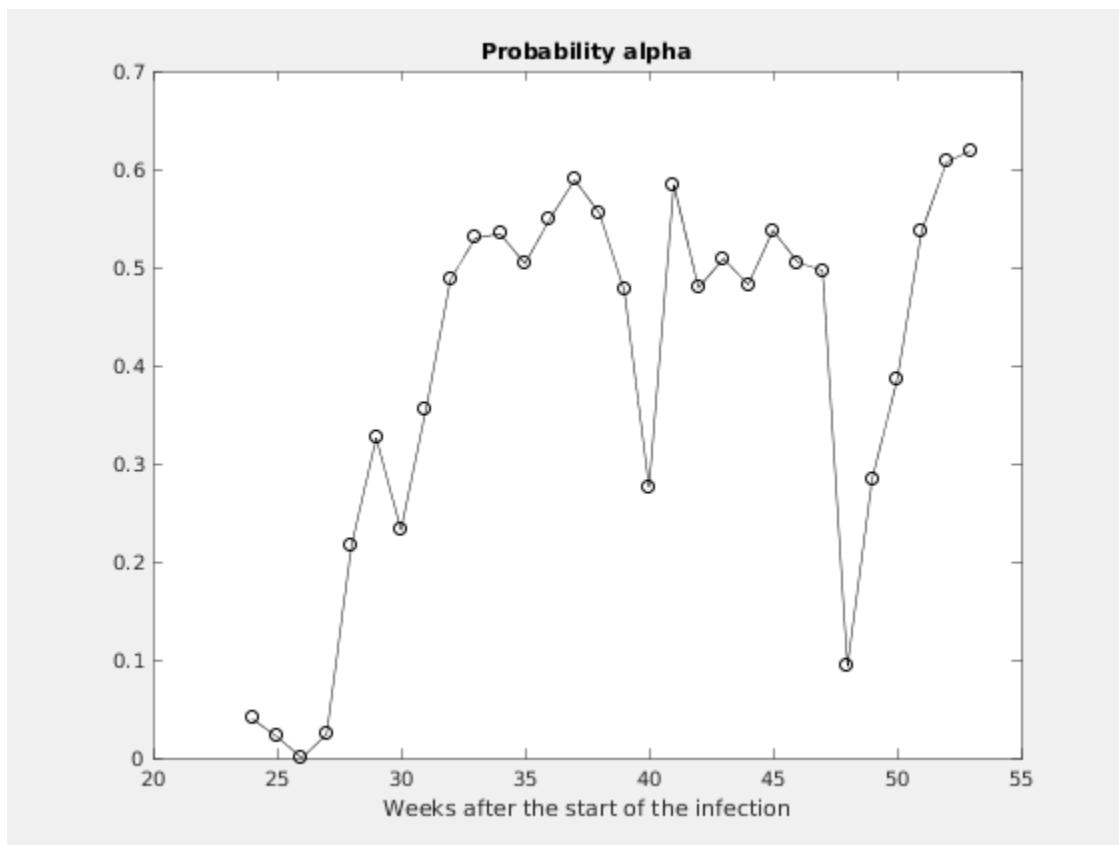
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# Chapter 2. Estimating of the main parameter and some predictions

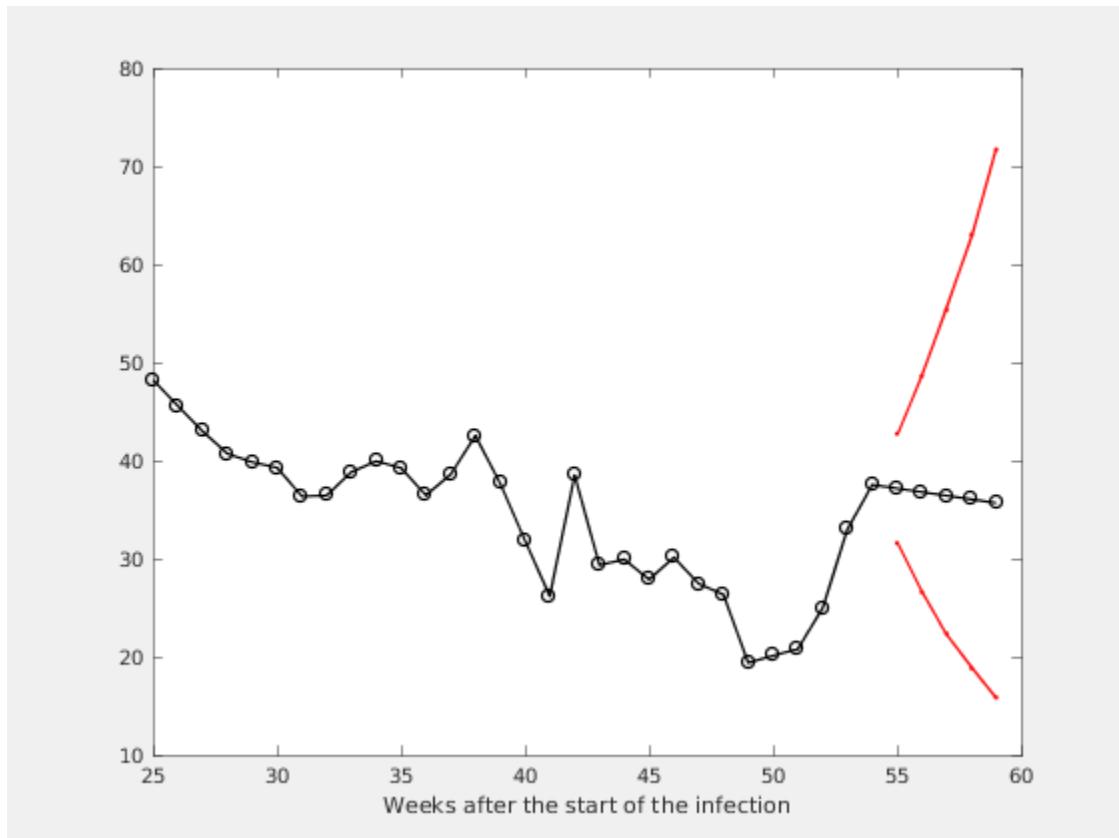
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



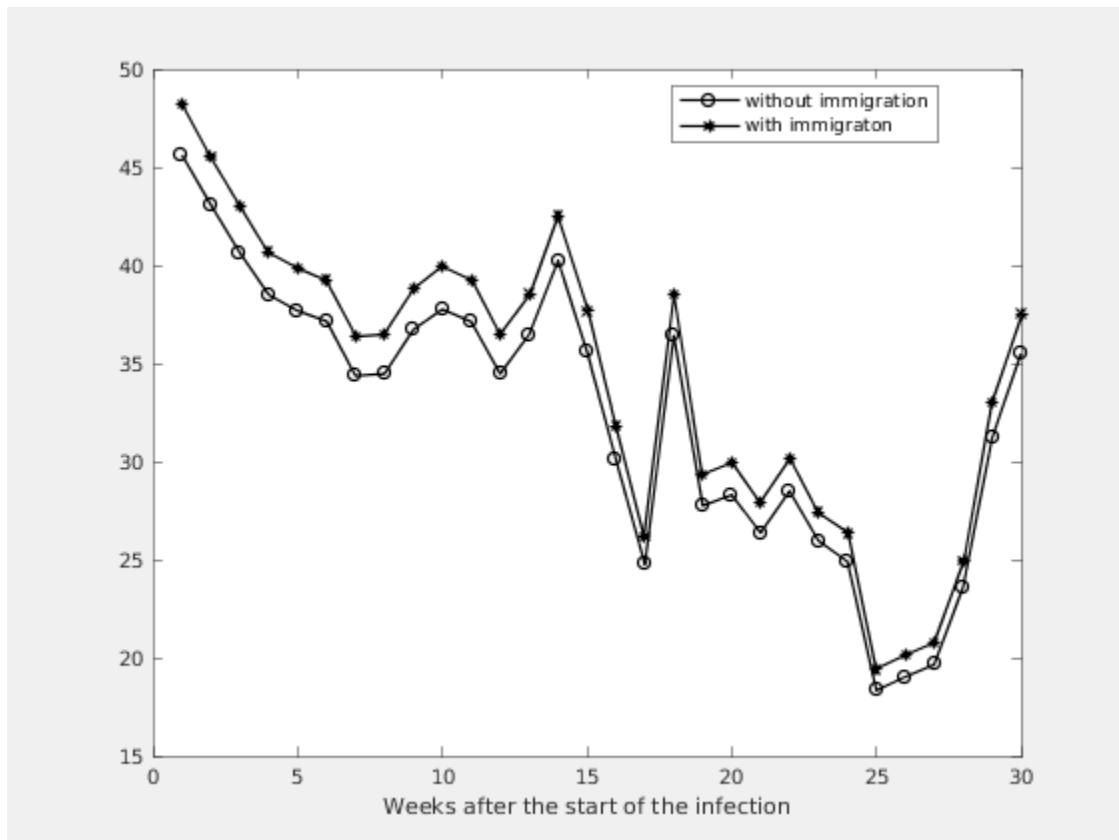
**Figure 2.2. Figure**



**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	
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4	0.9649	0.8128	- 1.1169	0.4968	26	25
3	0.9673	0.8152	- 1.1194	0.0935	19	18
2	0.9748	0.8220	- 1.1276	0.2845	20	19
1	0.9851	0.8337	- 1.1364	0.3848	21	20
0	0.9898	0.8380	- 1.1417	0.5373	25	24