

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

**Var100 - week 53**

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## **Branching stochastic processes as models of Covid-19 epidemic development : Var100 - 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 Var100. 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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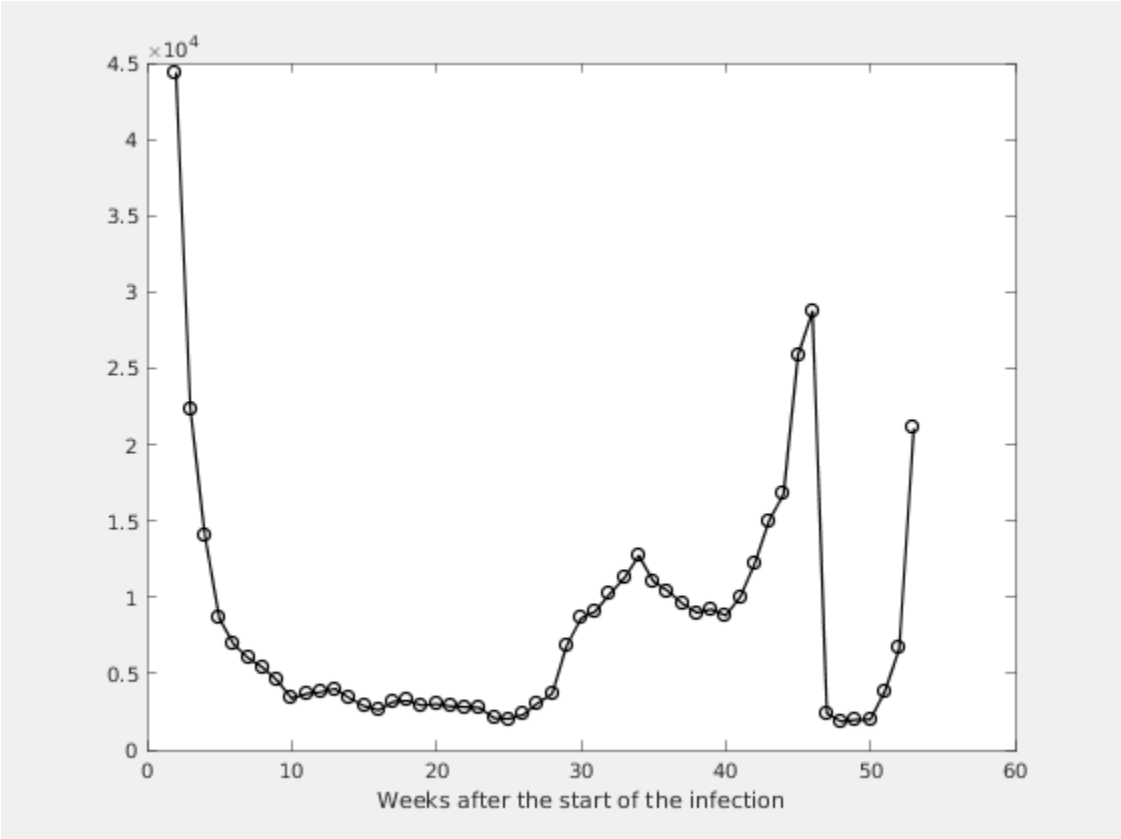
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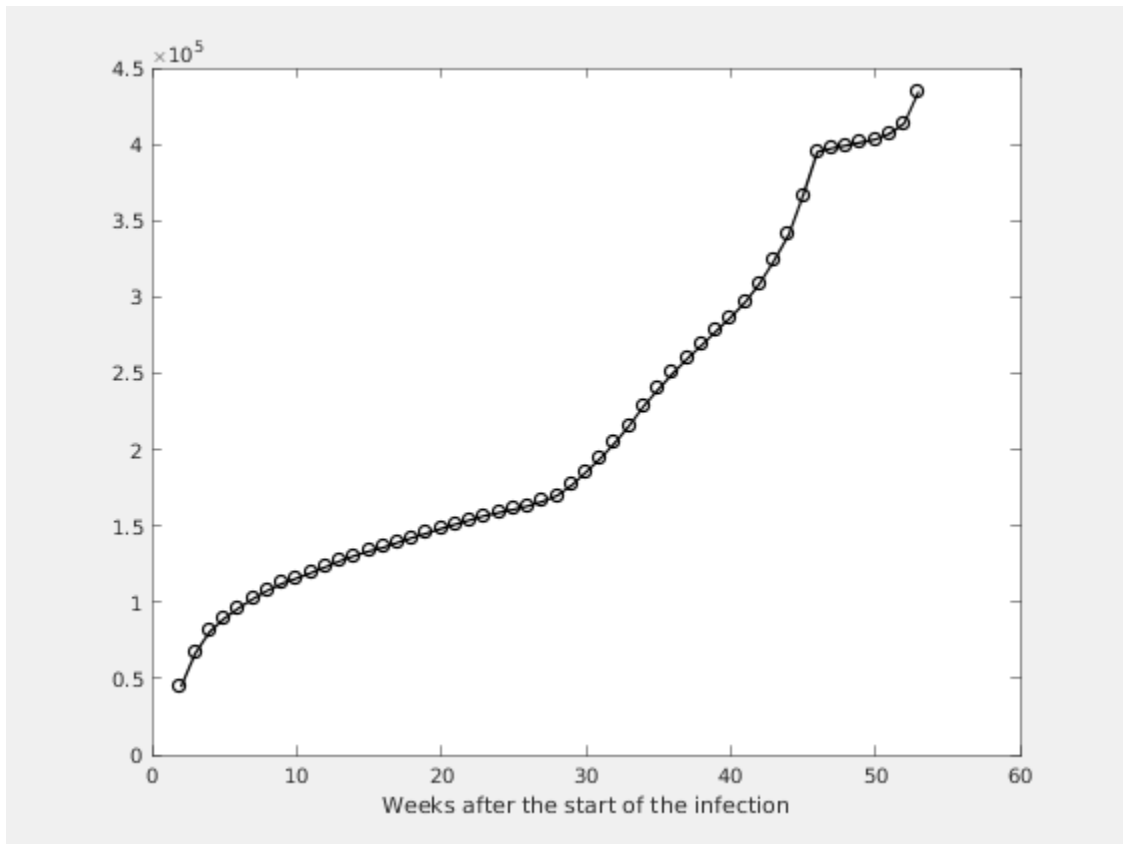
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# Chapter 1. Observed Infection data

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

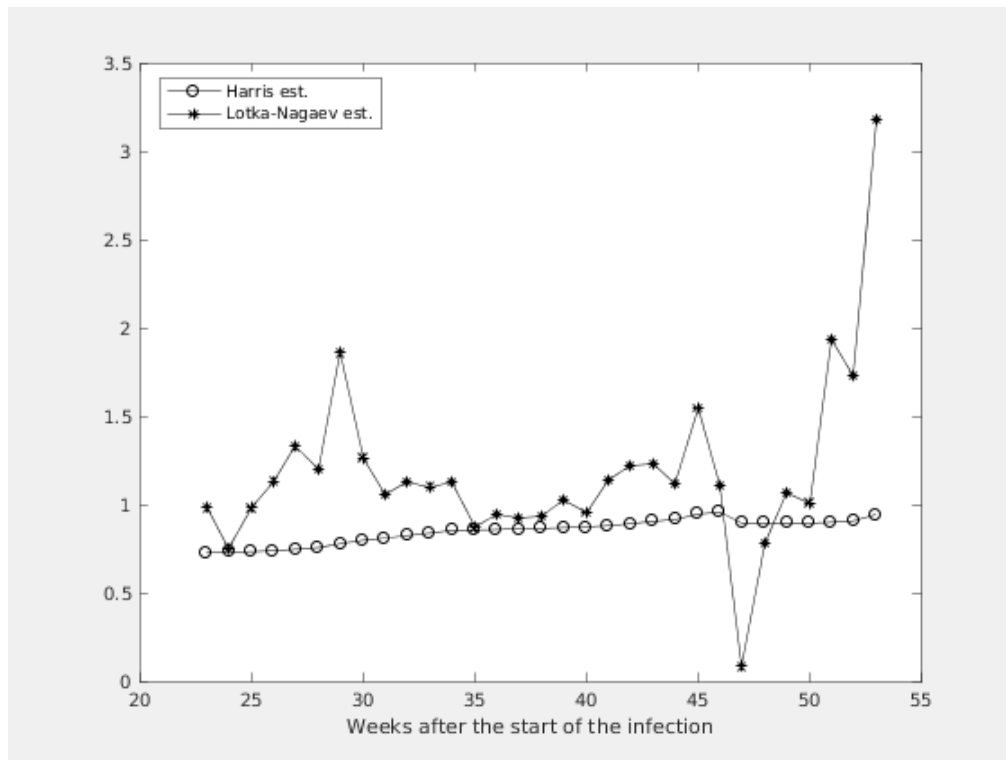
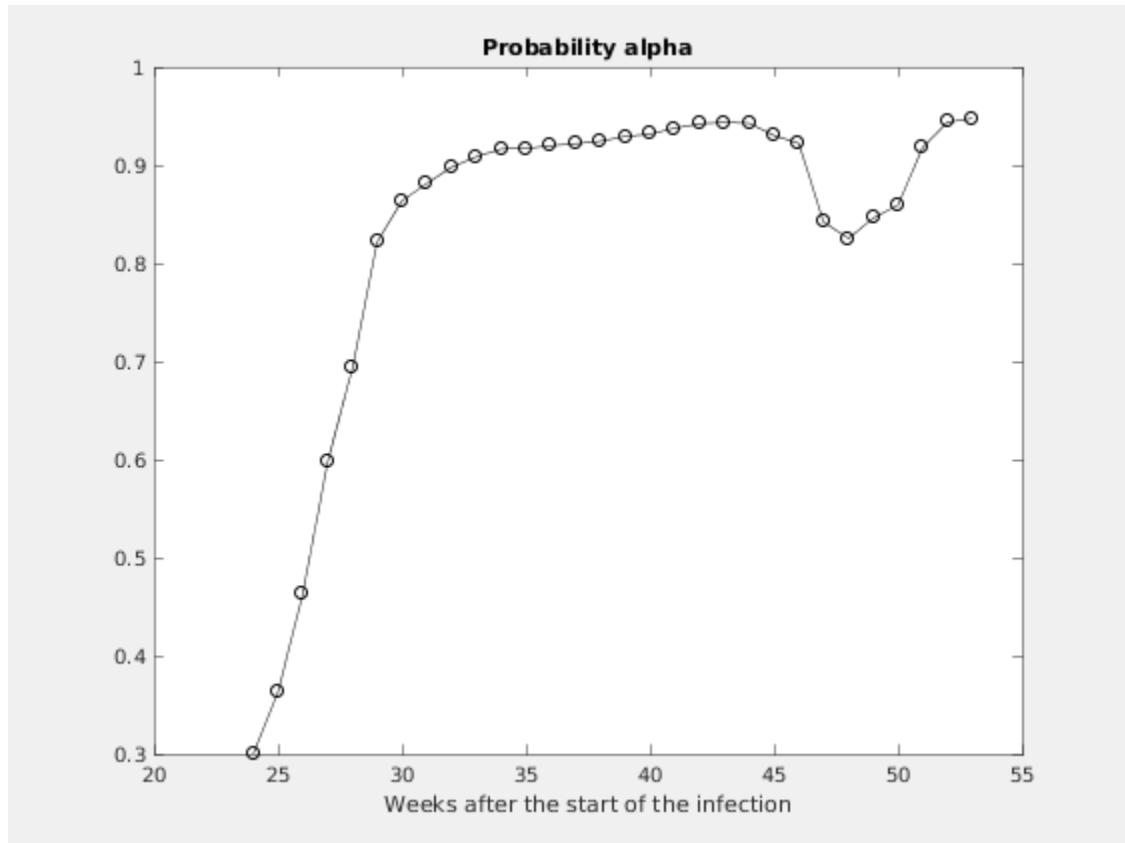
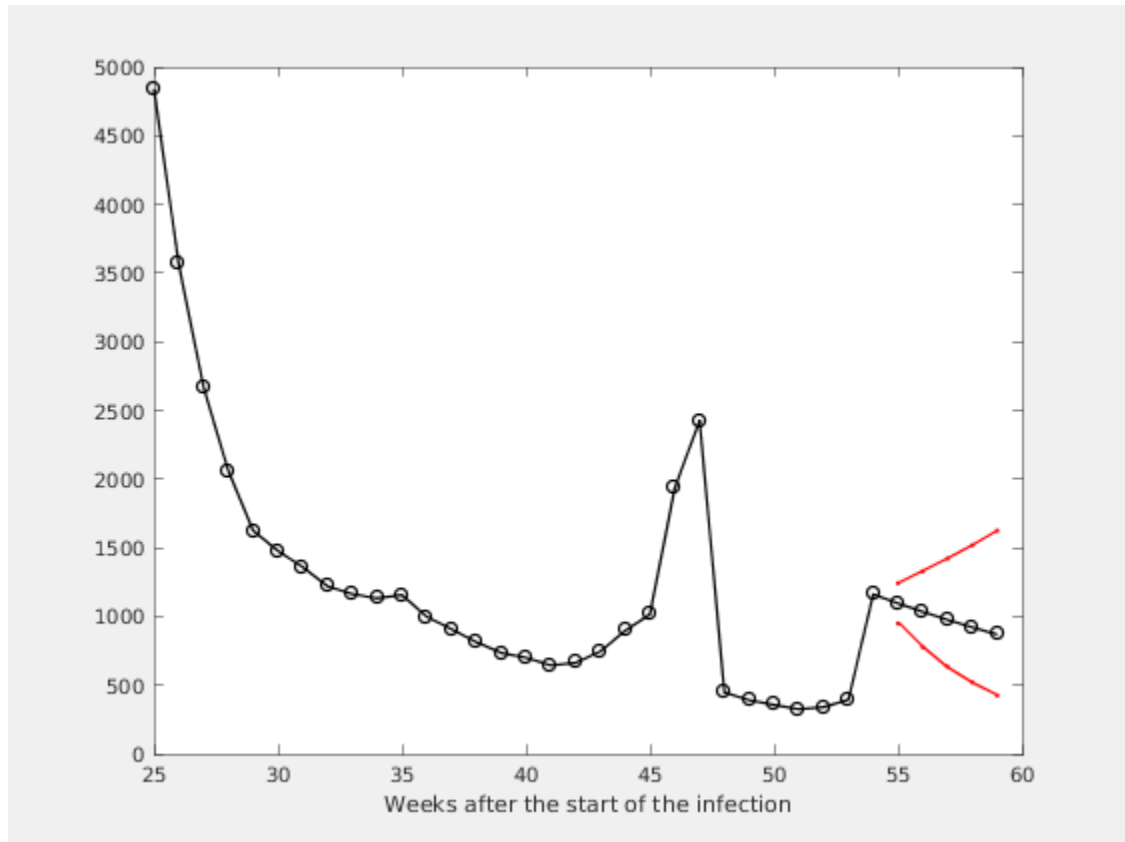


Figure 2.2. Figure

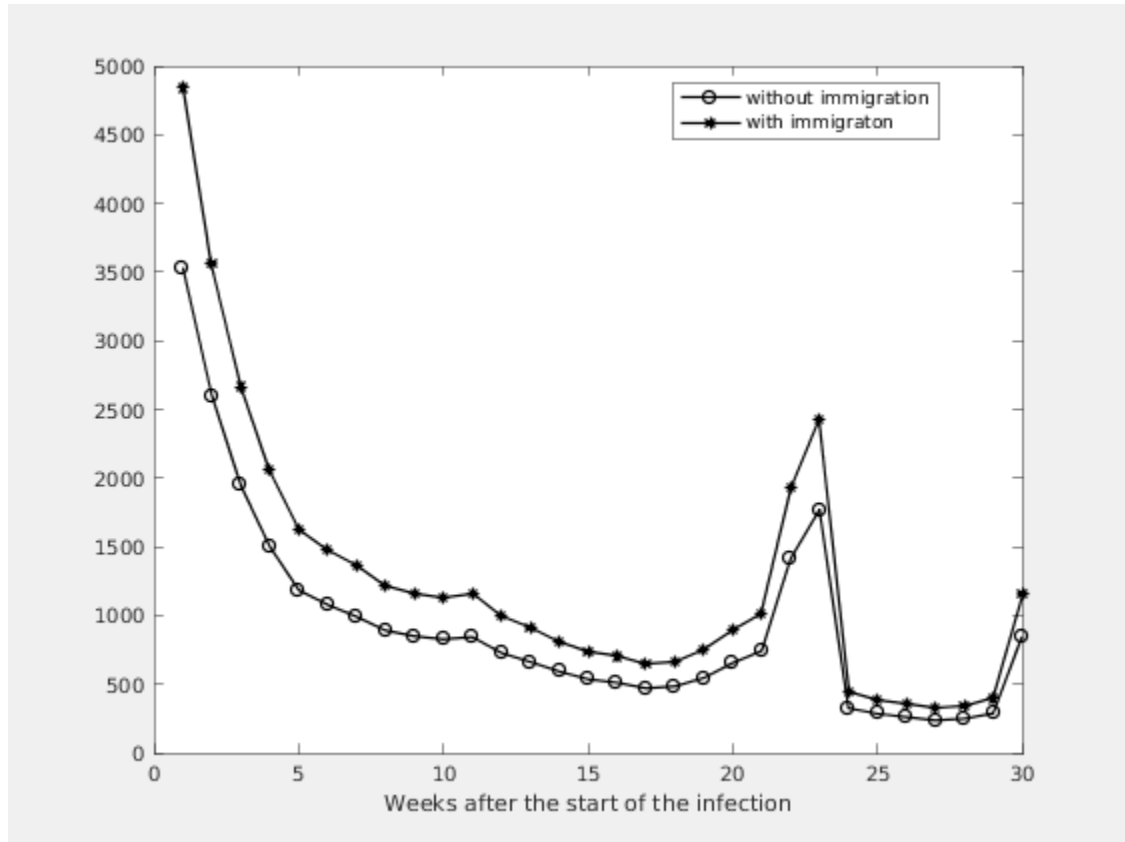




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

k	m	ci	alpha	A1	M1
4	0.8937	0.7913 - 0.9962	0.8420	446	326
3	0.8943	0.7933 - 0.9953	0.8254	391	285
2	0.8994	0.7996 - 0.9992	0.8461	357	261
1	0.9072	0.8086 - 1.0058	0.8593	325	237
0	0.9436	0.8435 - 1.0437	0.9183	341	249