

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

**Var115 - week 53**

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## **Branching stochastic processes as models of Covid-19 epidemic development : Var115 - 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 Var115. 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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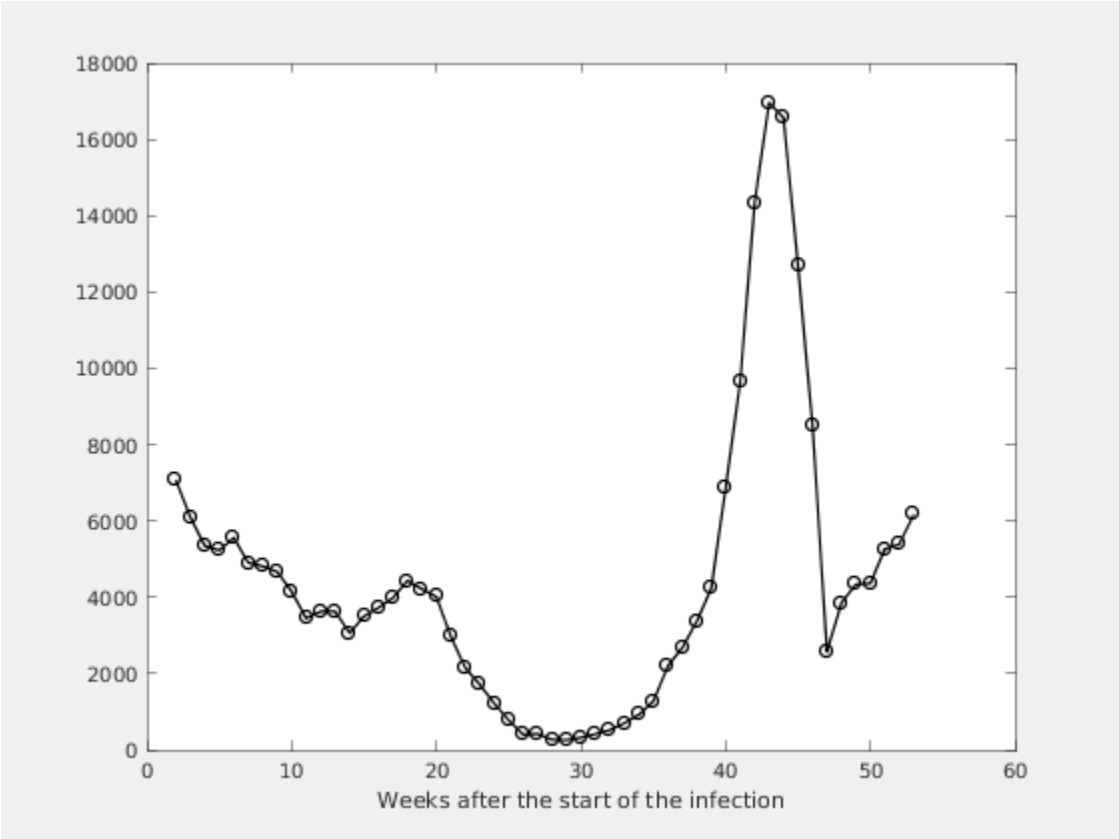
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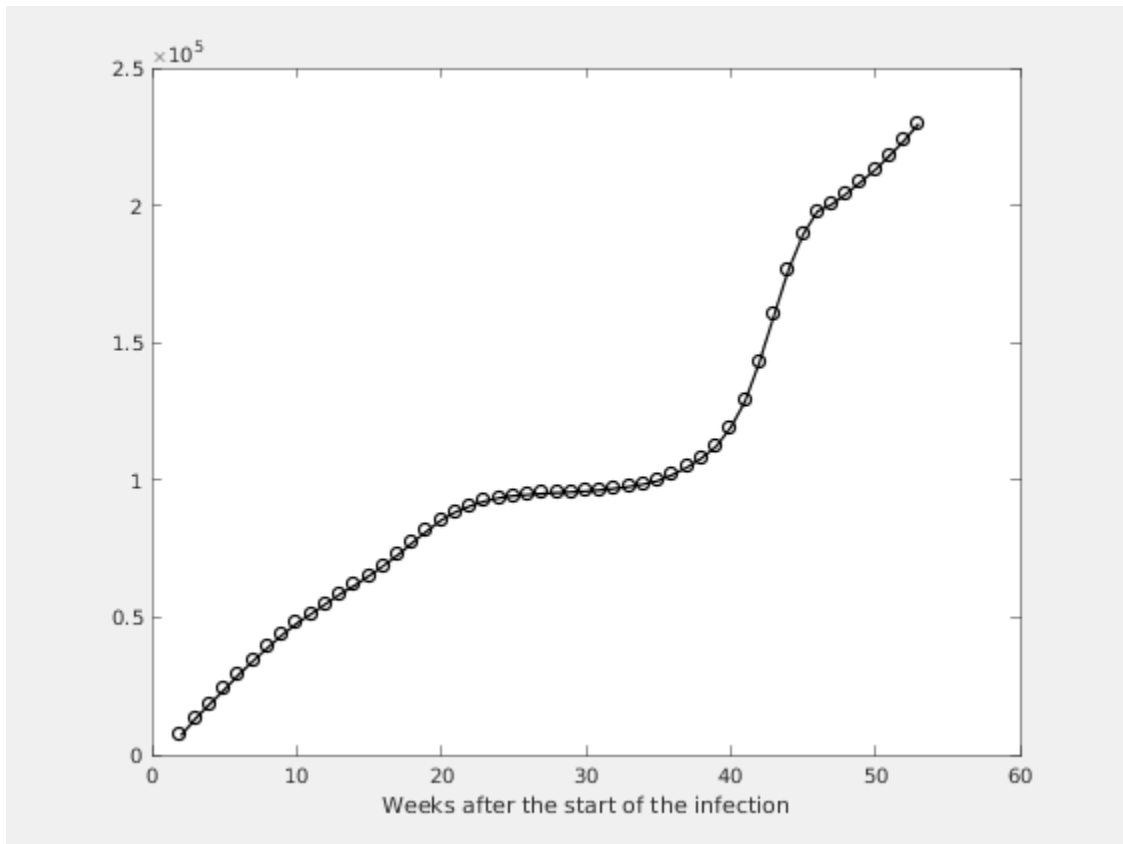
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# Chapter 1. Observed Infection data

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

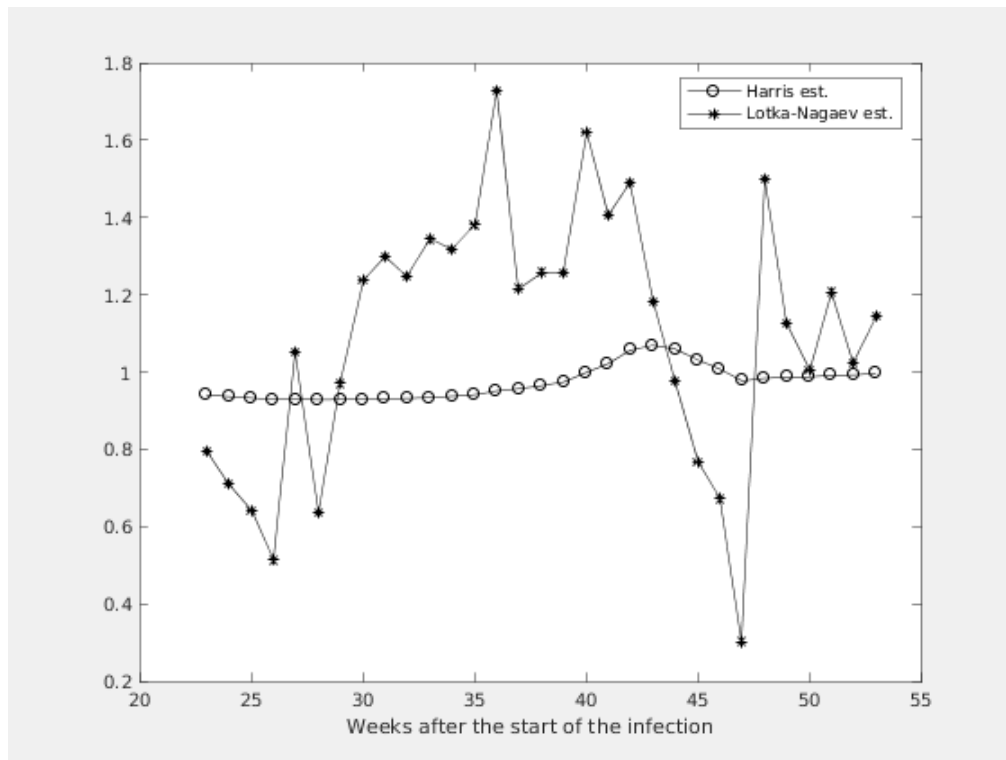
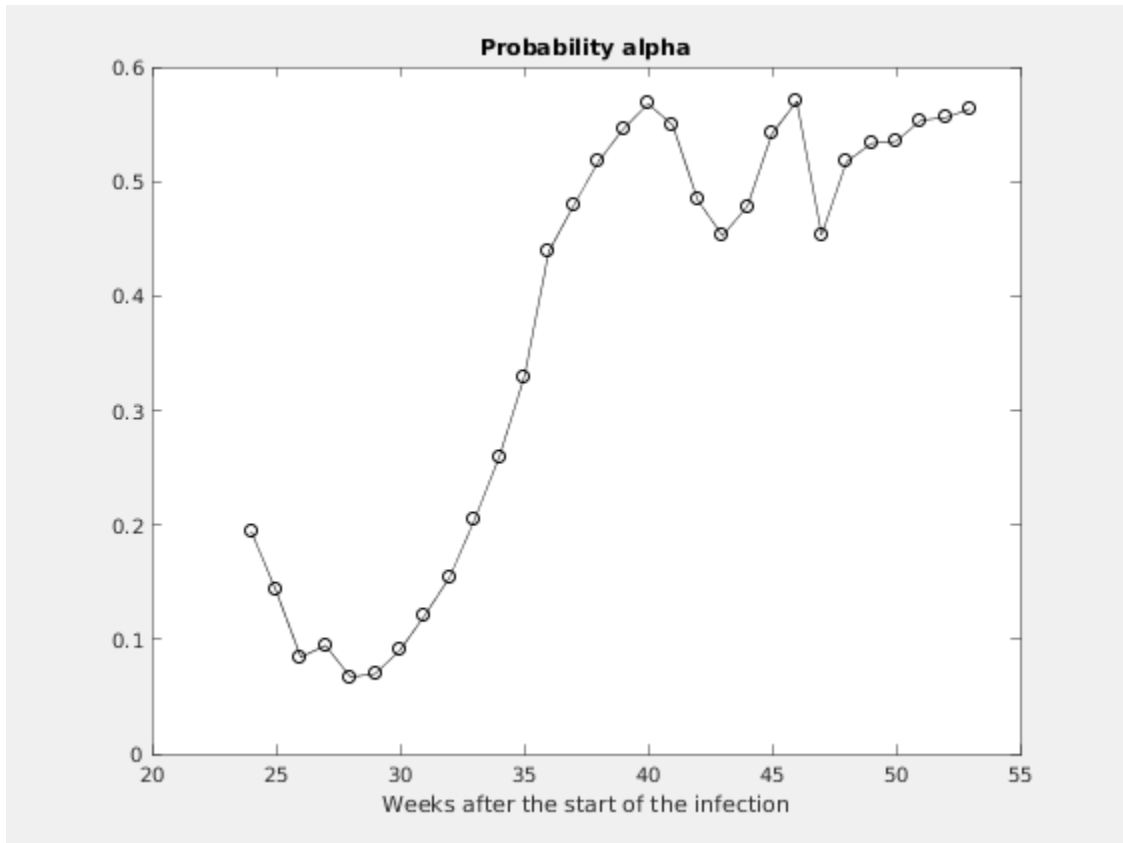
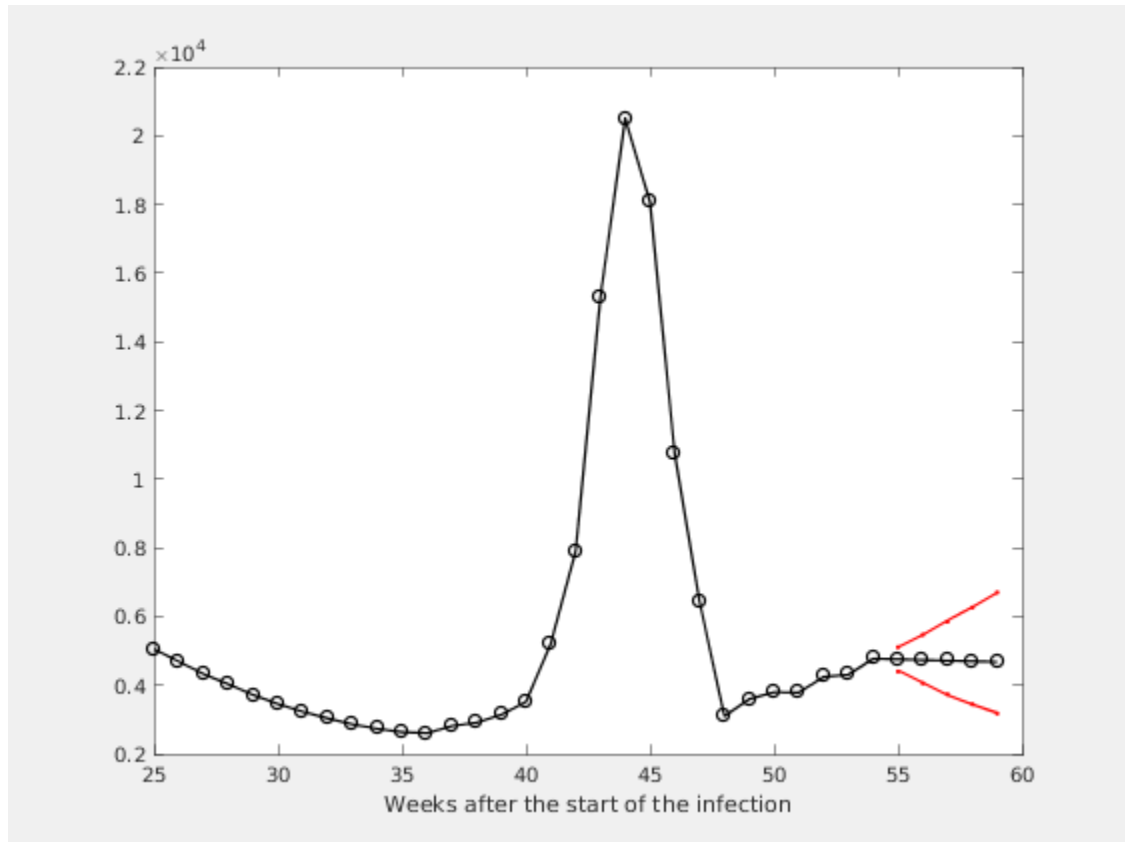


Figure 2.2. Figure

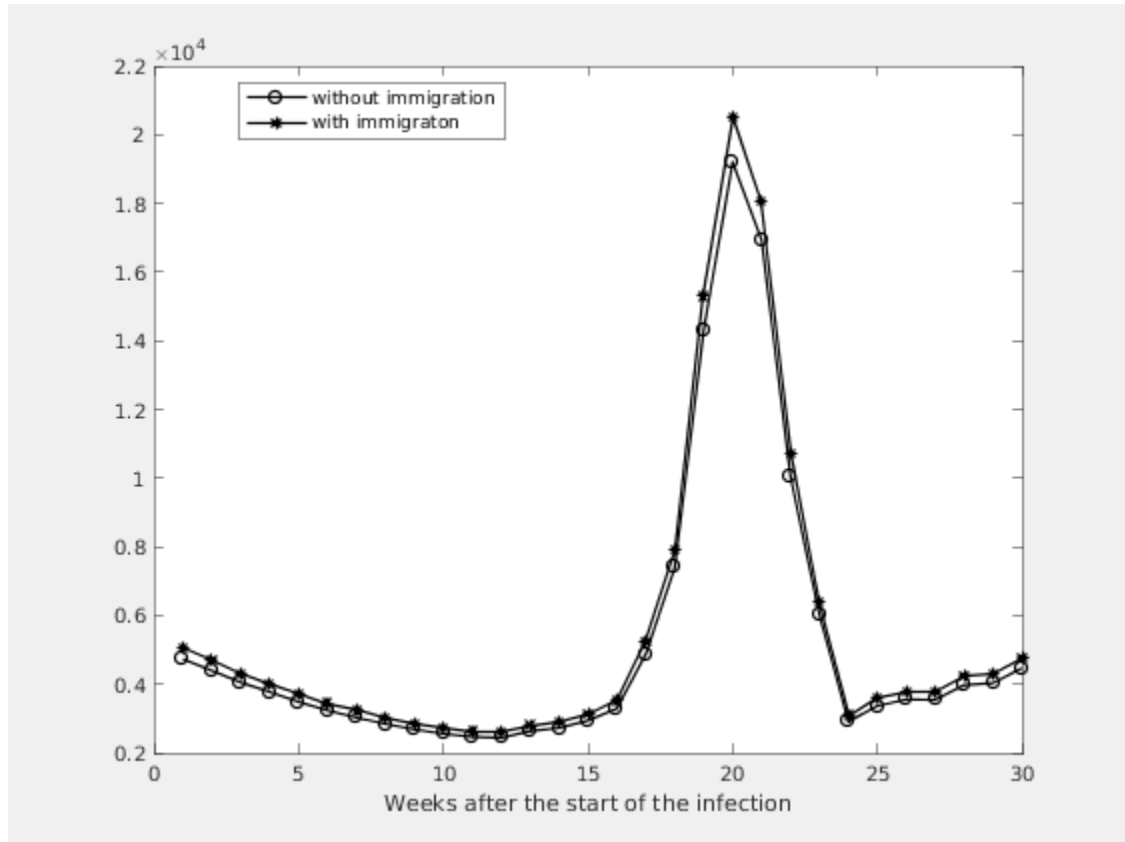




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

k	m	ci	alpha	A1	M1
4	0.9866	0.9054 - 1.0679	0.4533	3104	2907
3	0.9870	0.9054 - 1.0686	0.5173	3597	3369
2	0.9915	0.9114 - 1.0717	0.5332	3797	3557
1	0.9923	0.9138 - 1.0708	0.5353	3787	3547
0	0.9960	0.9186 - 1.0733	0.5535	4246	3977