

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

**Var104 - week 53**

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## **Branching stochastic processes as models of Covid-19 epidemic development : Var104 - 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 Var104. 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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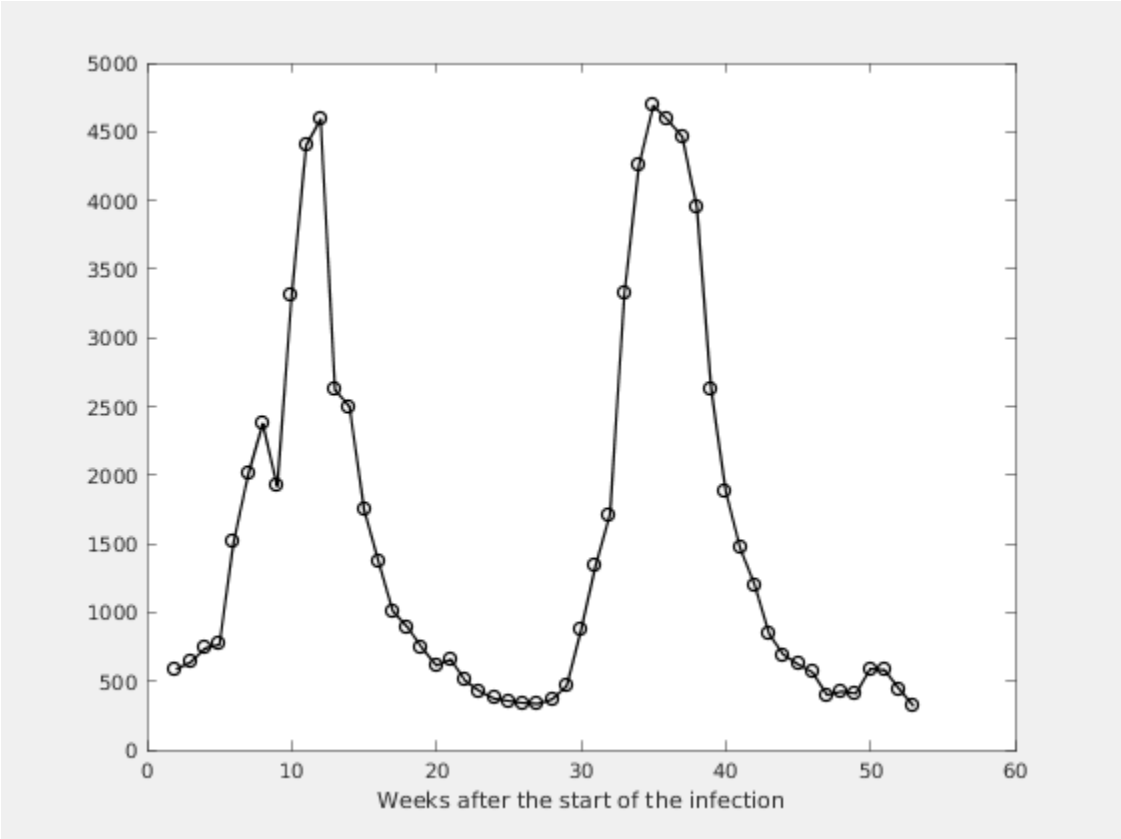
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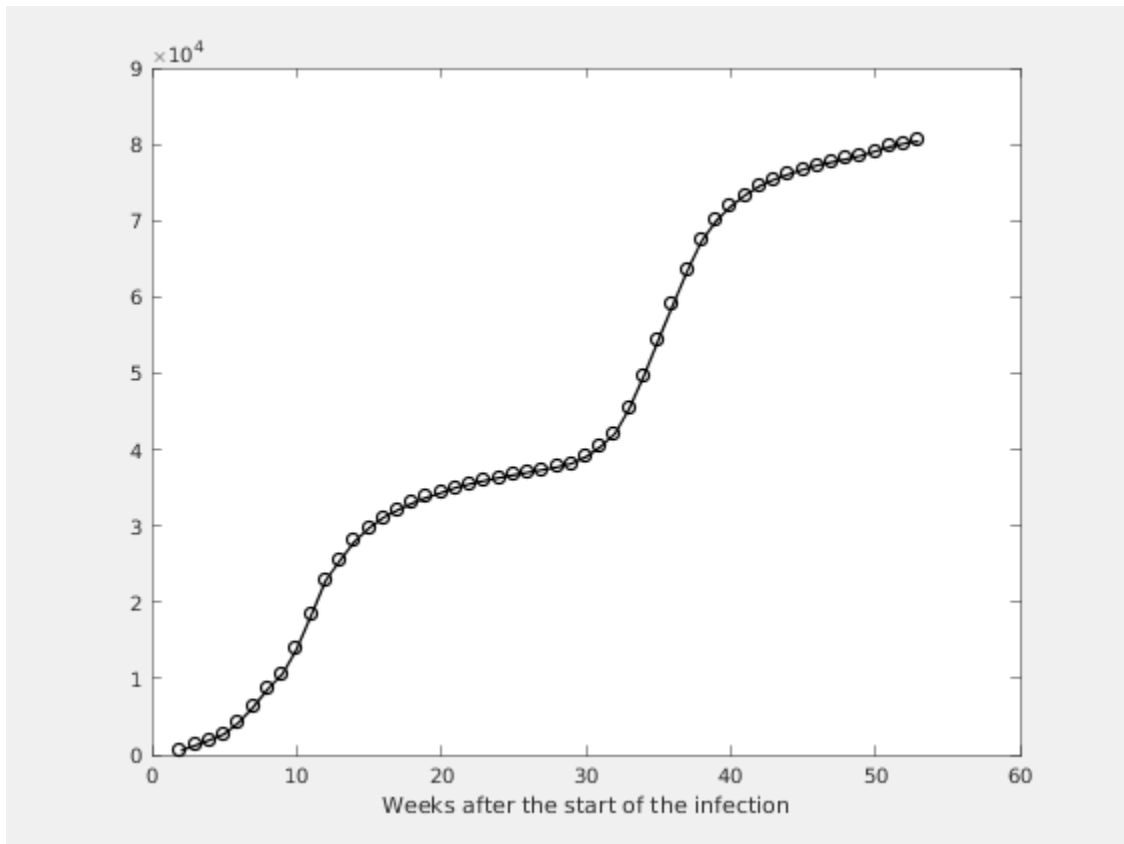
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# Chapter 1. Observed Infection data

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

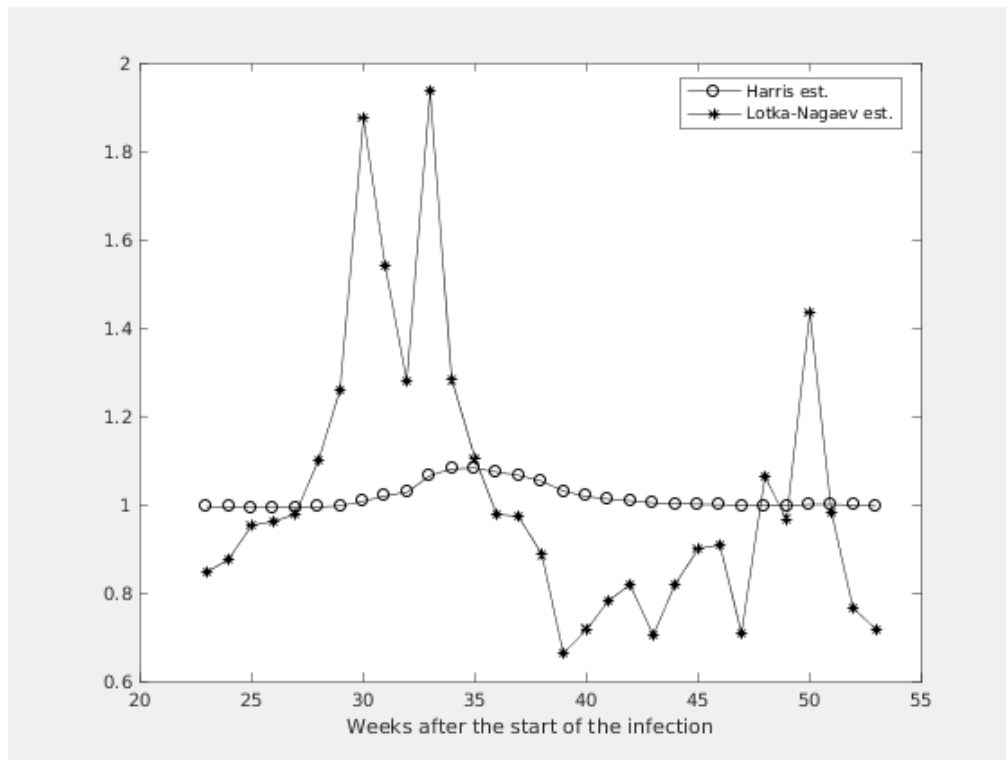
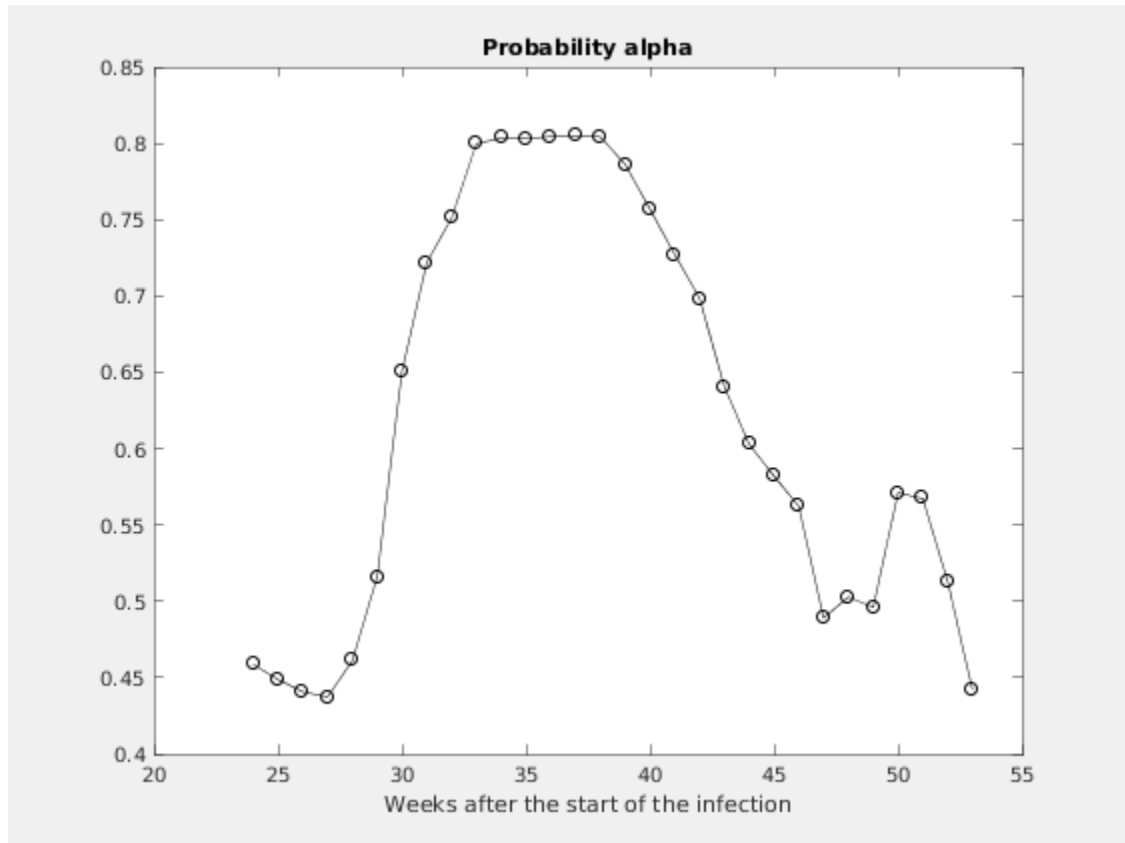
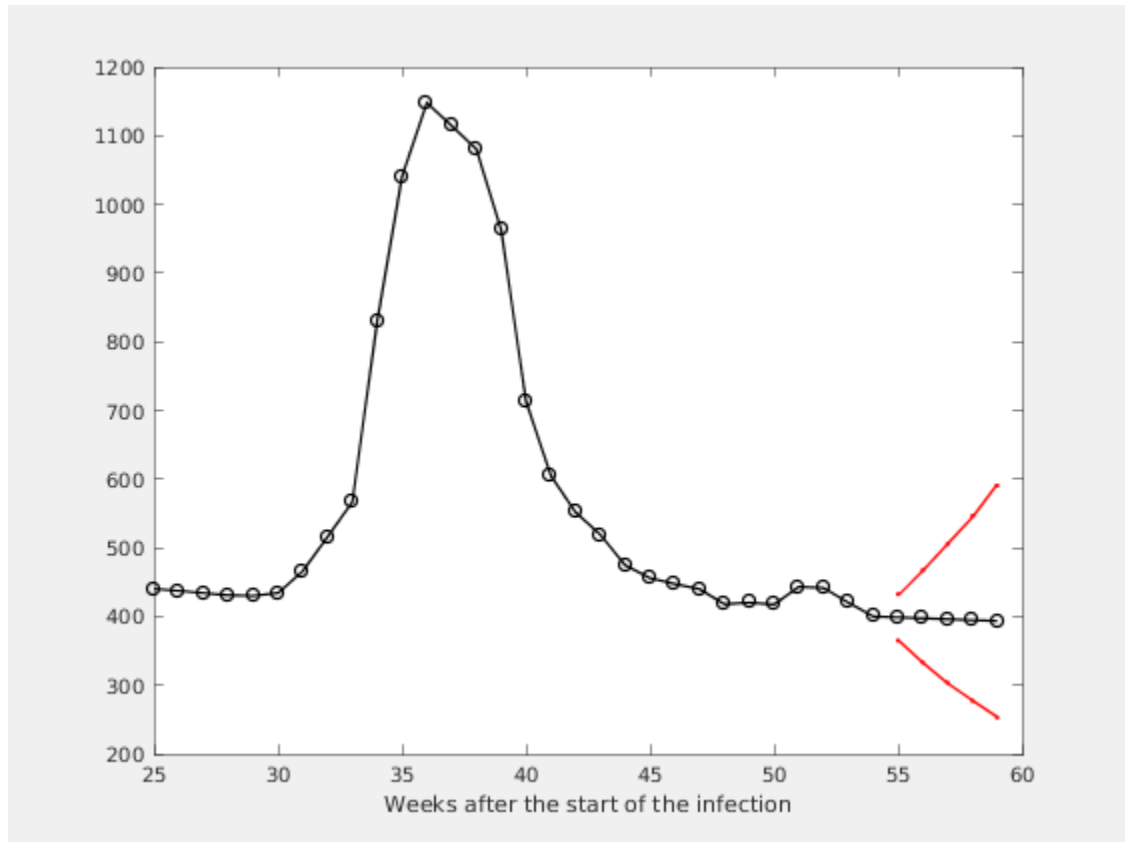


Figure 2.2. Figure

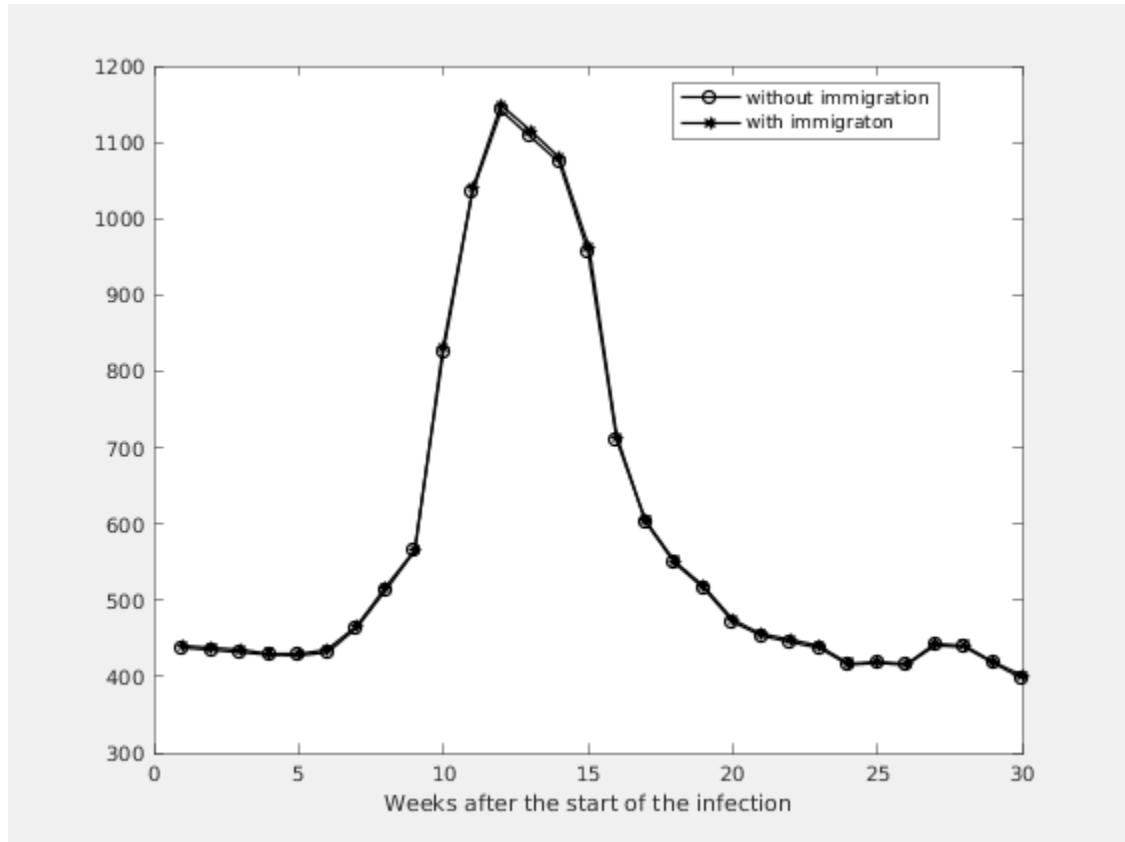




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

k	m	ci	alpha	A1	M1
4	0.9977	0.9071 - 1.0883	0.4891	418	415
3	1.0000	0.9106 - 1.0894	0.5028	420	418
2	0.9999	0.9117 - 1.0881	0.4960	418	415
1	0.9982	0.9107 - 1.0857	0.5710	443	441
0	0.9966	0.9103 - 1.0829	0.5673	442	439