

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

**Guernsey - week 53**

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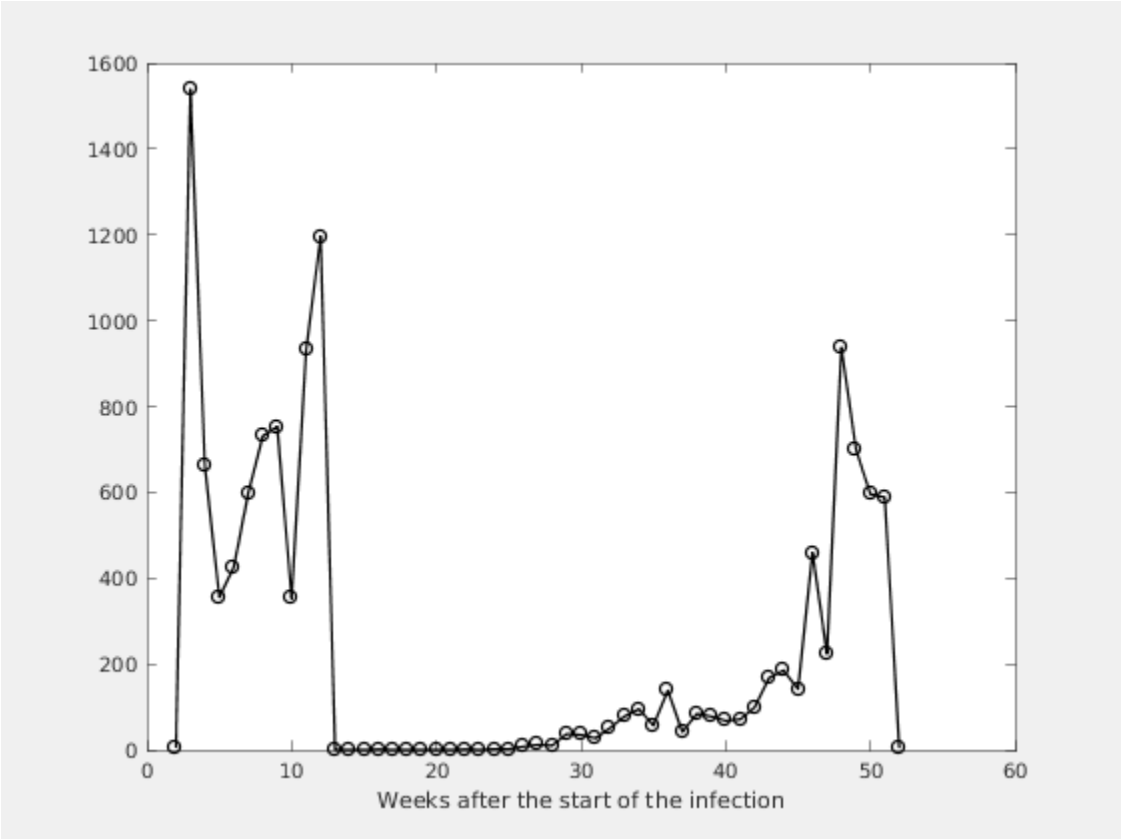
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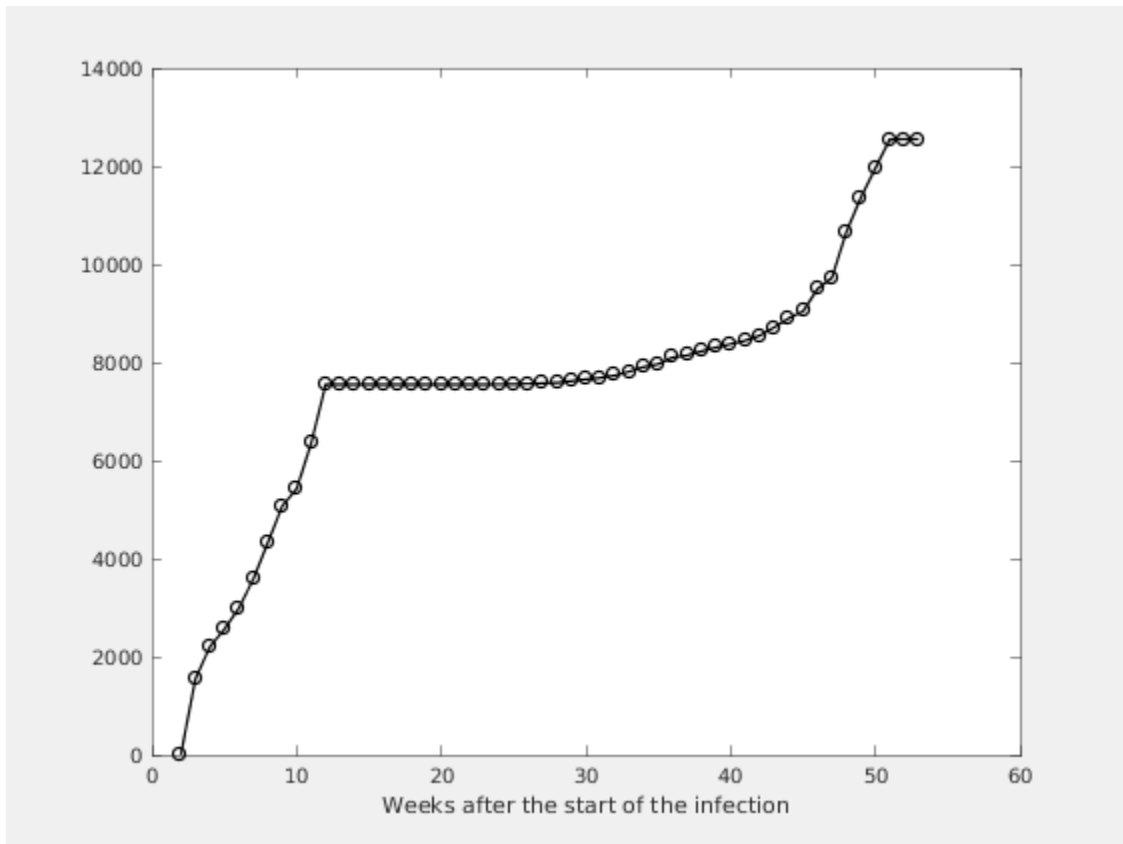
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# Chapter 1. Observed Infection data

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

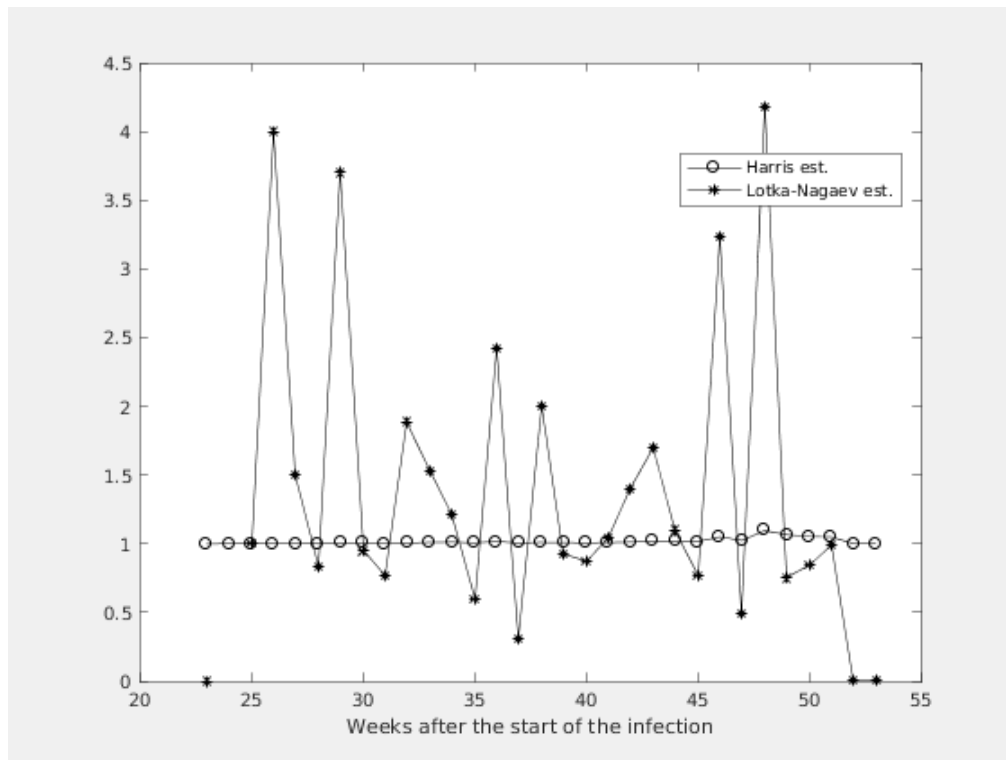
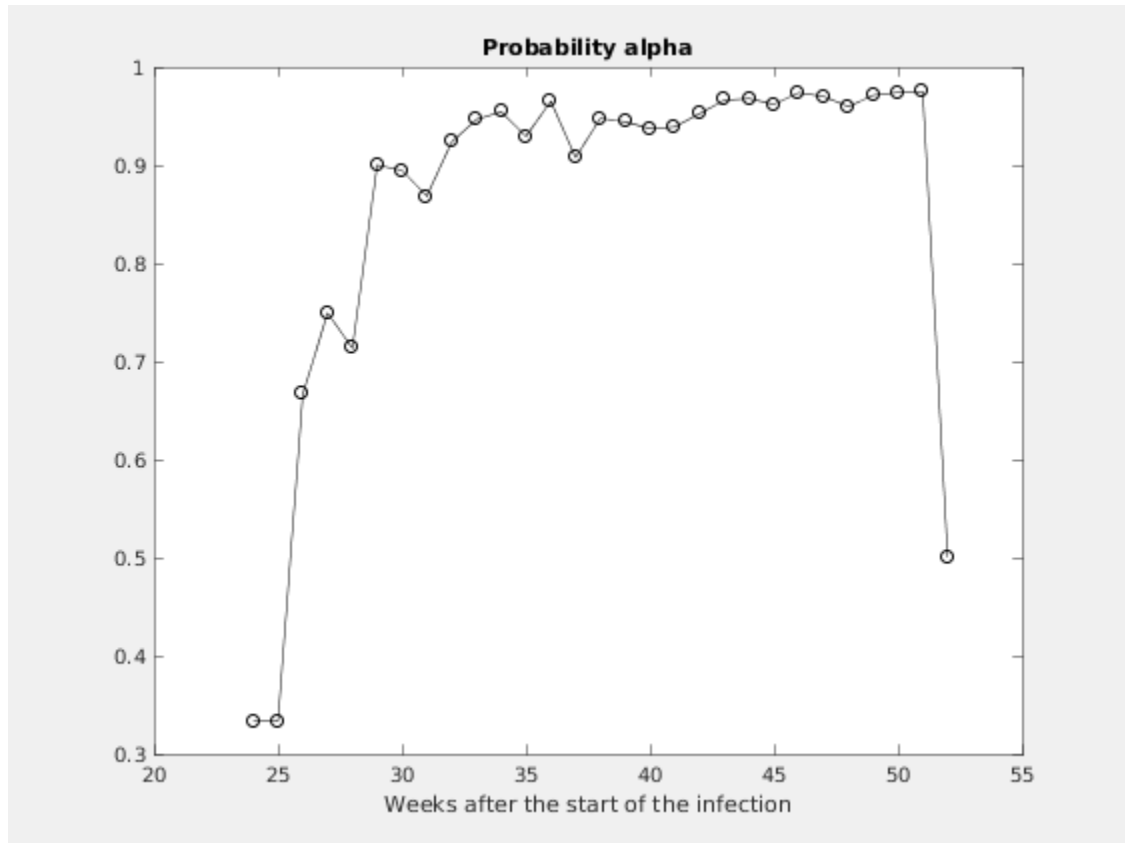
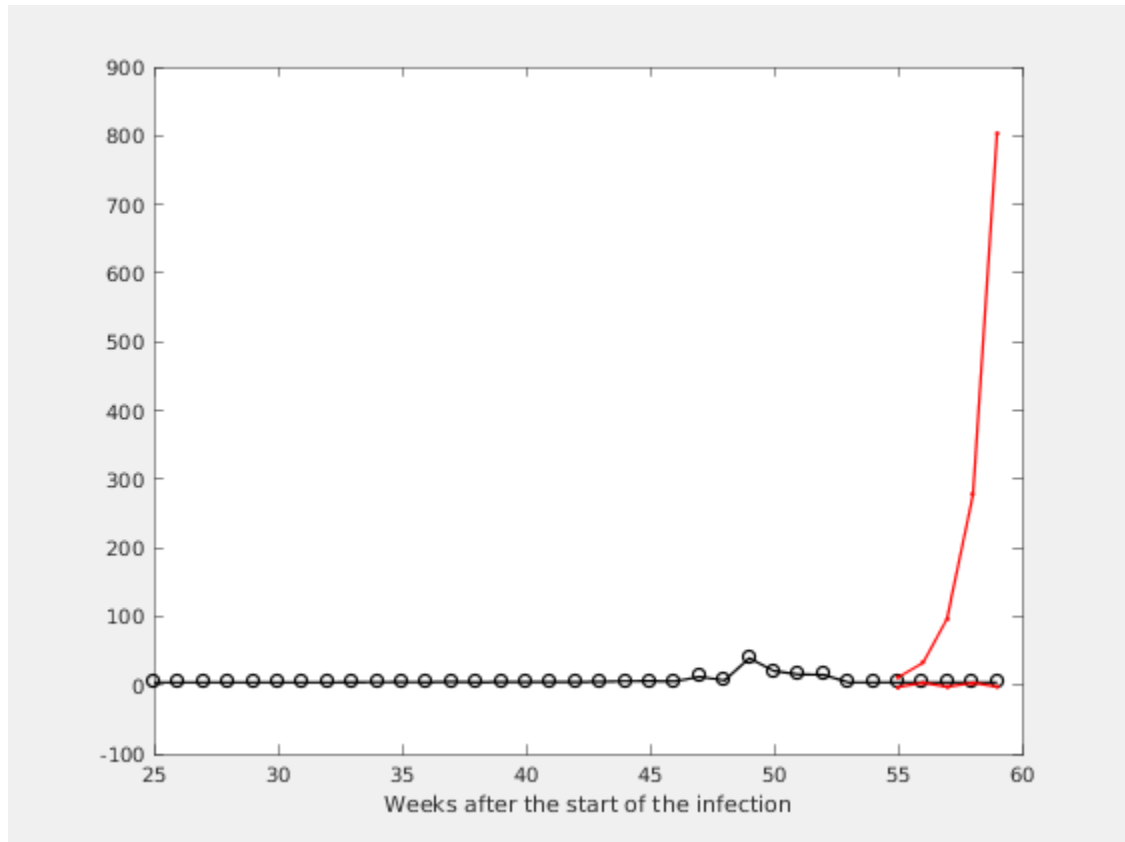


Figure 2.2. Figure

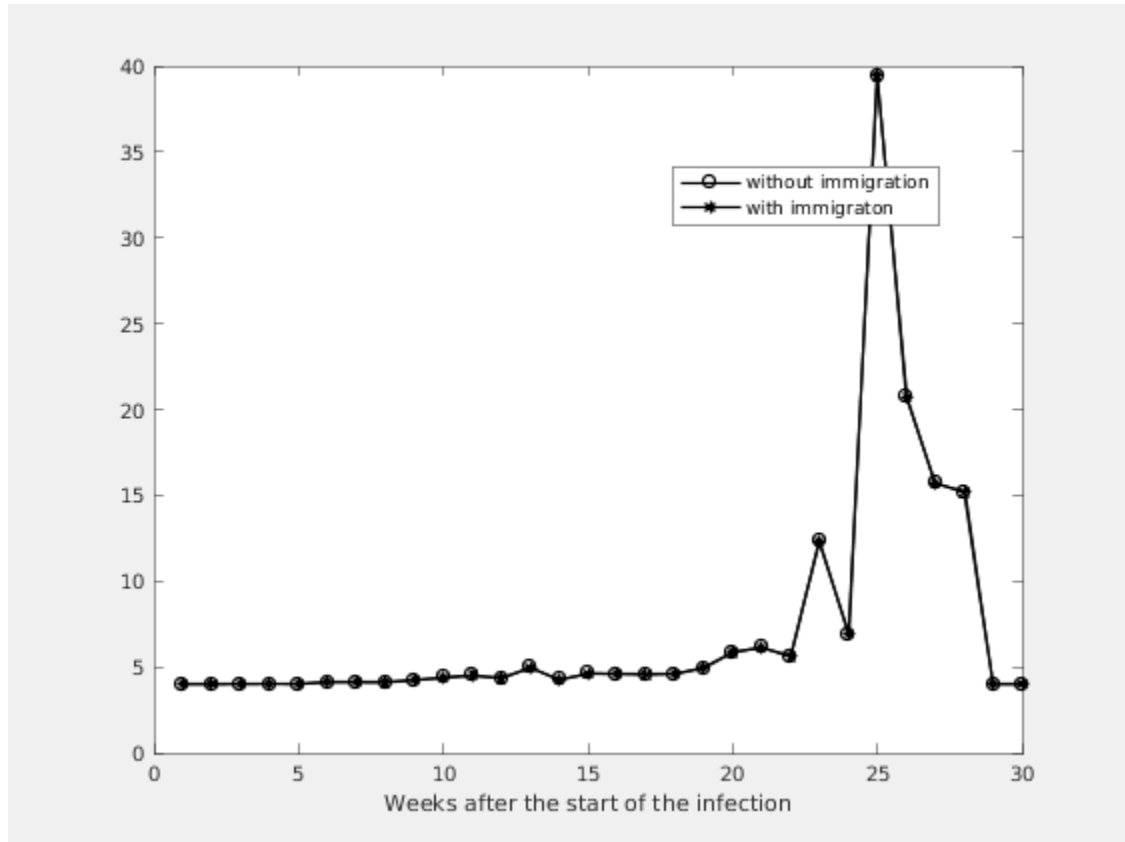




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

k	m	ci	alpha	A1	M1
4	1.0653	-1.2448 - 3.3754	0.9701	7	7
3	1.0520	-1.2054 - 3.3095	0.9596	39	39
2	1.0488	-1.0788 - 3.1763	0.9713	21	21
1	0.9999	-1.0345 - 3.0344	0.9743	16	16
0	0.9999	-0.9582 - 2.9581	0.9749	15	15