

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

**Thailand - week 53**

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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 Thailand. 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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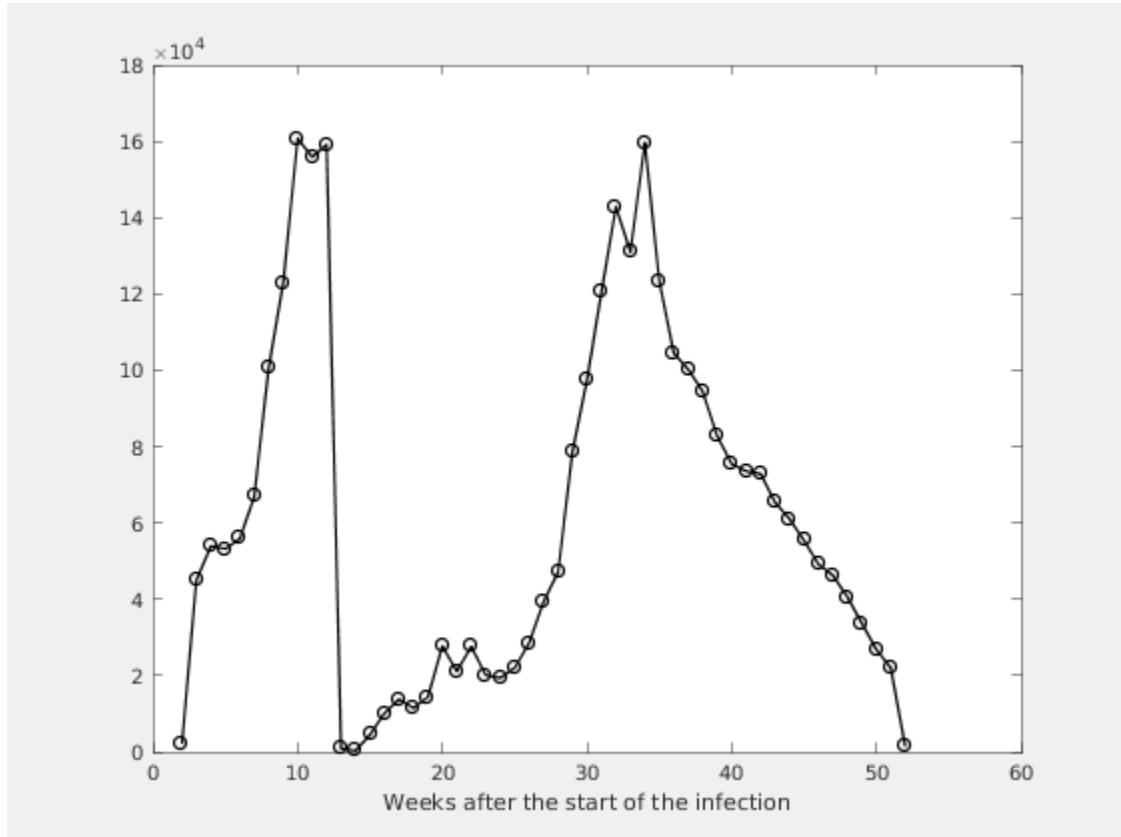
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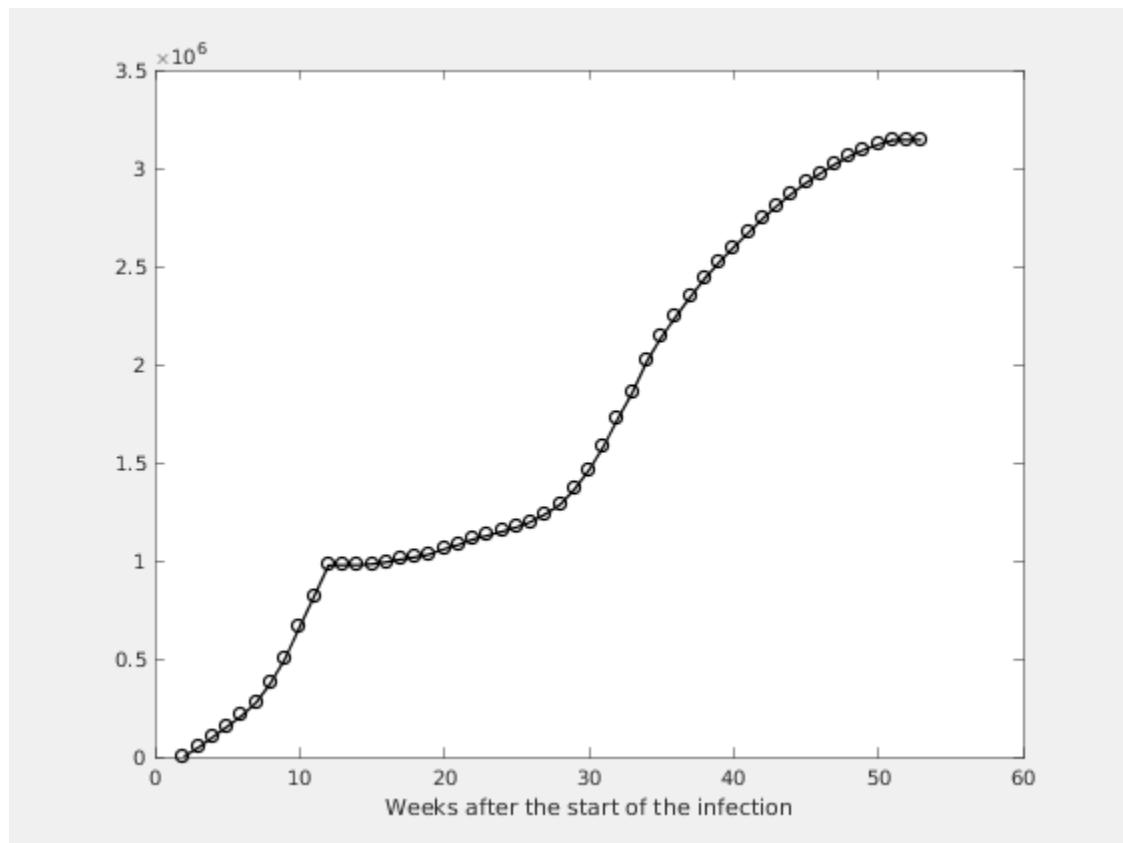
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# Chapter 1. Observed Infection data

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



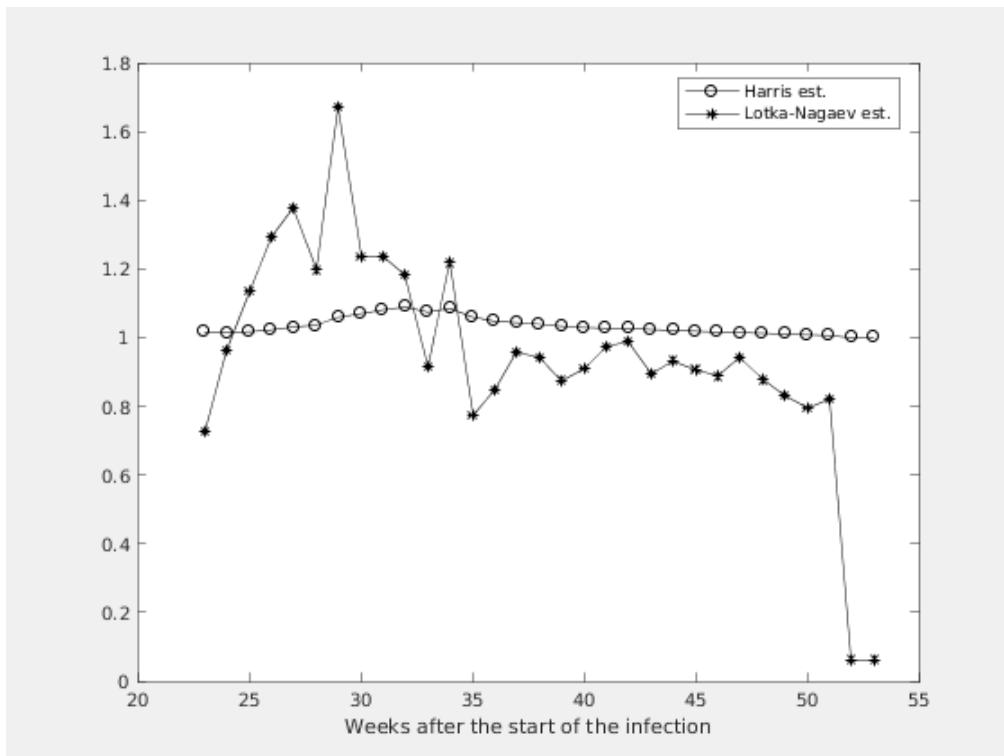
**Figure 1.2. Number of the total registered cases**



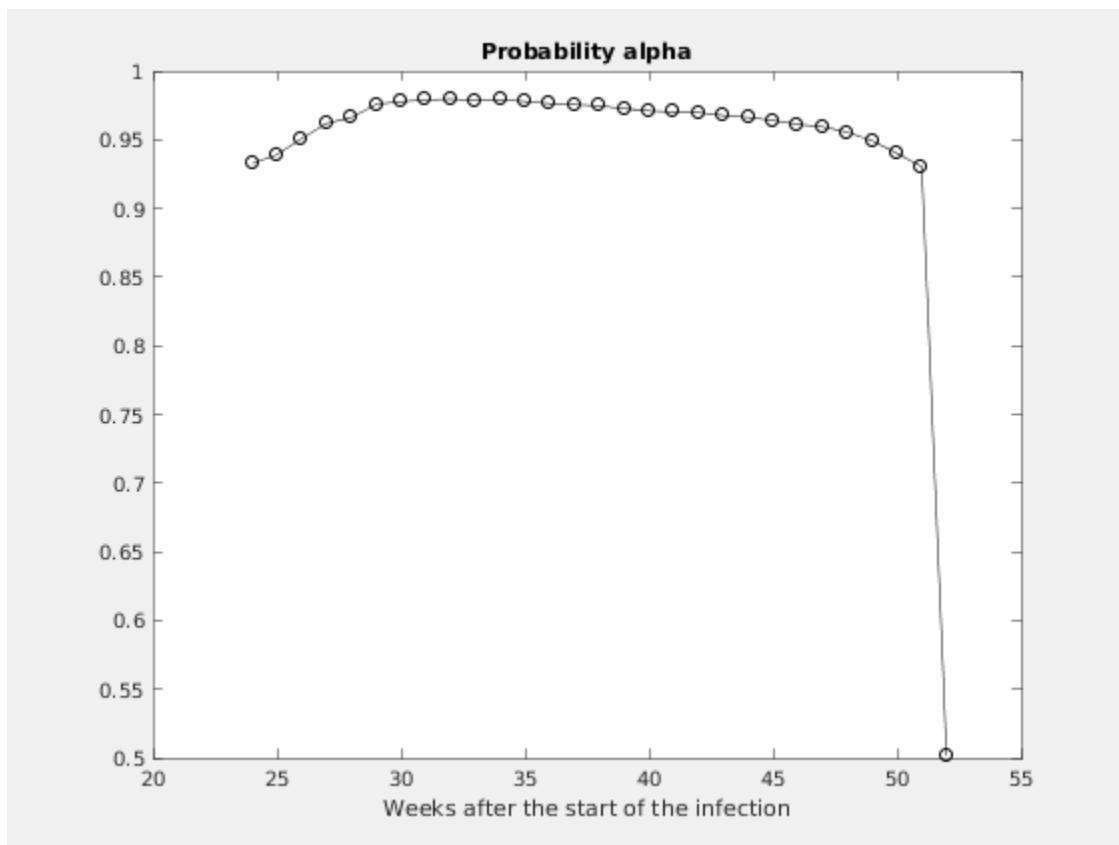
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# Chapter 2. Estimating of the main parameter and some predictions

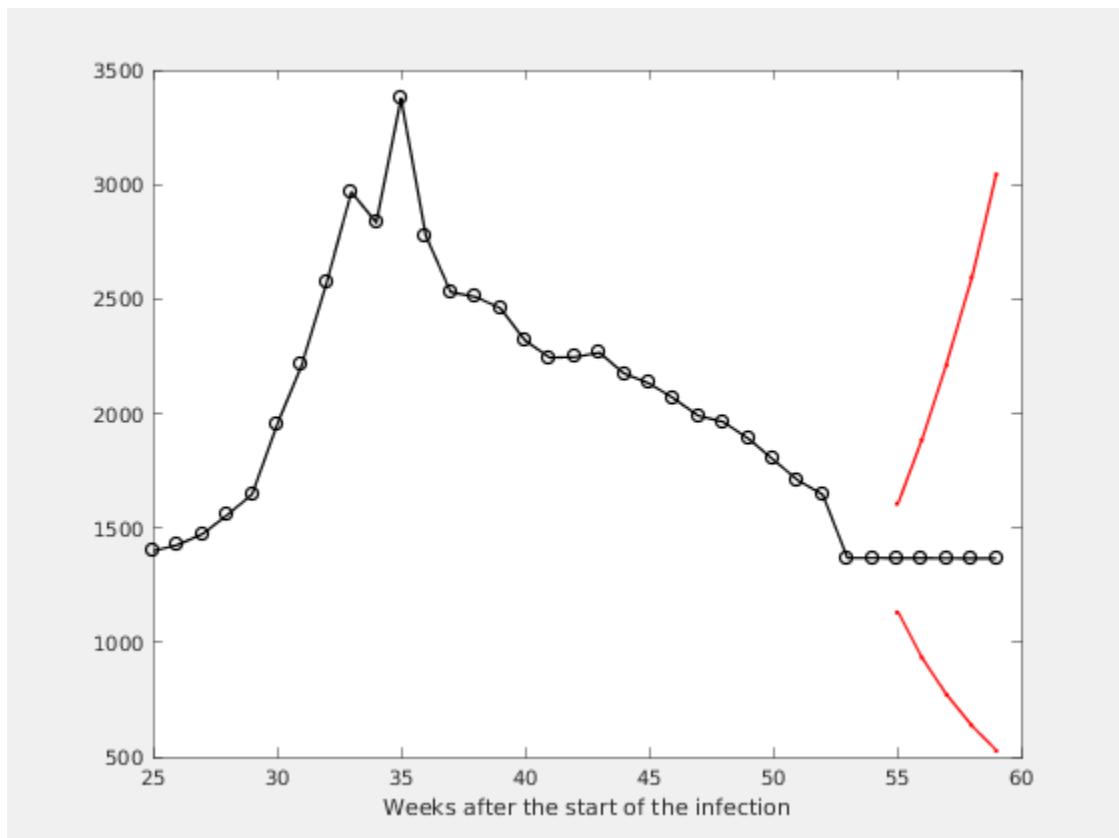
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



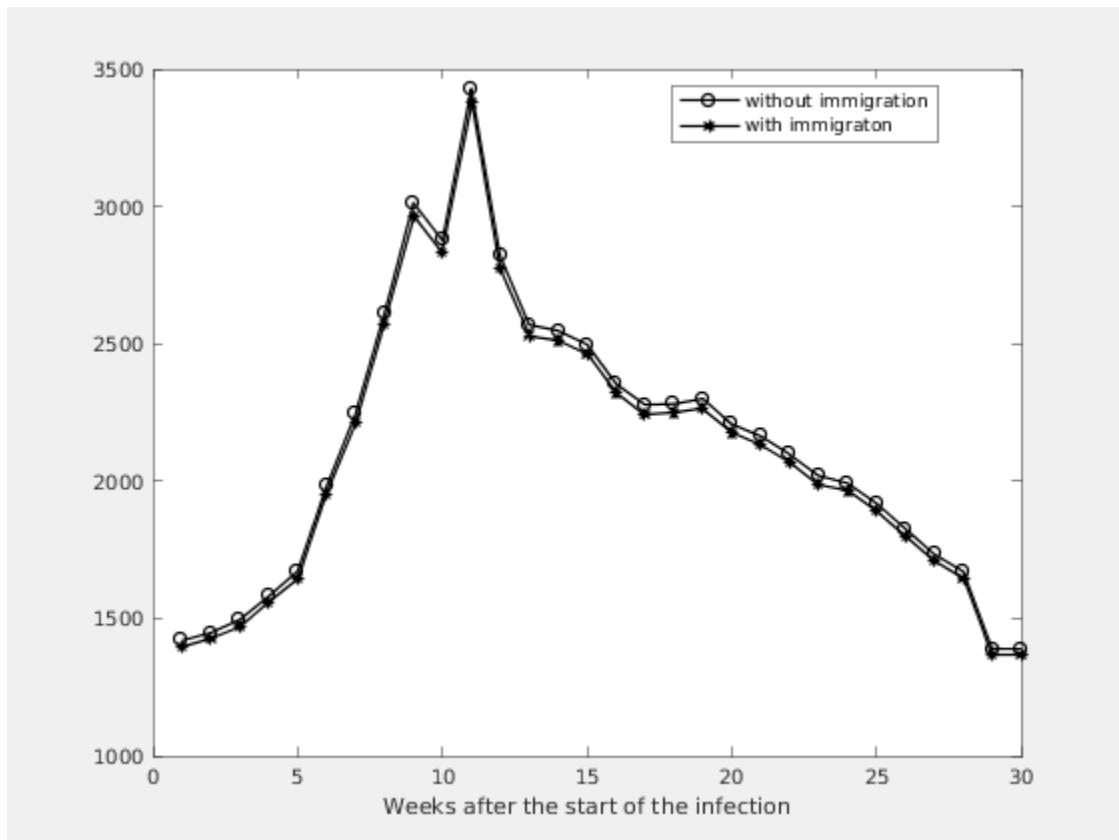
**Figure 2.2. Figure**



**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	1.0103	0.8240	- 1.1967	0.9593	1963
3	1.0080	0.8250	- 1.1909	0.9555	1891
2	1.0064	0.8265	- 1.1863	0.9493	1799
1	0.9998	0.8226	- 1.1769	0.9401	1708
0	0.9998	0.8251	- 1.1744	0.9304	1646
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