

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

**Senegal - 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 Senegal. 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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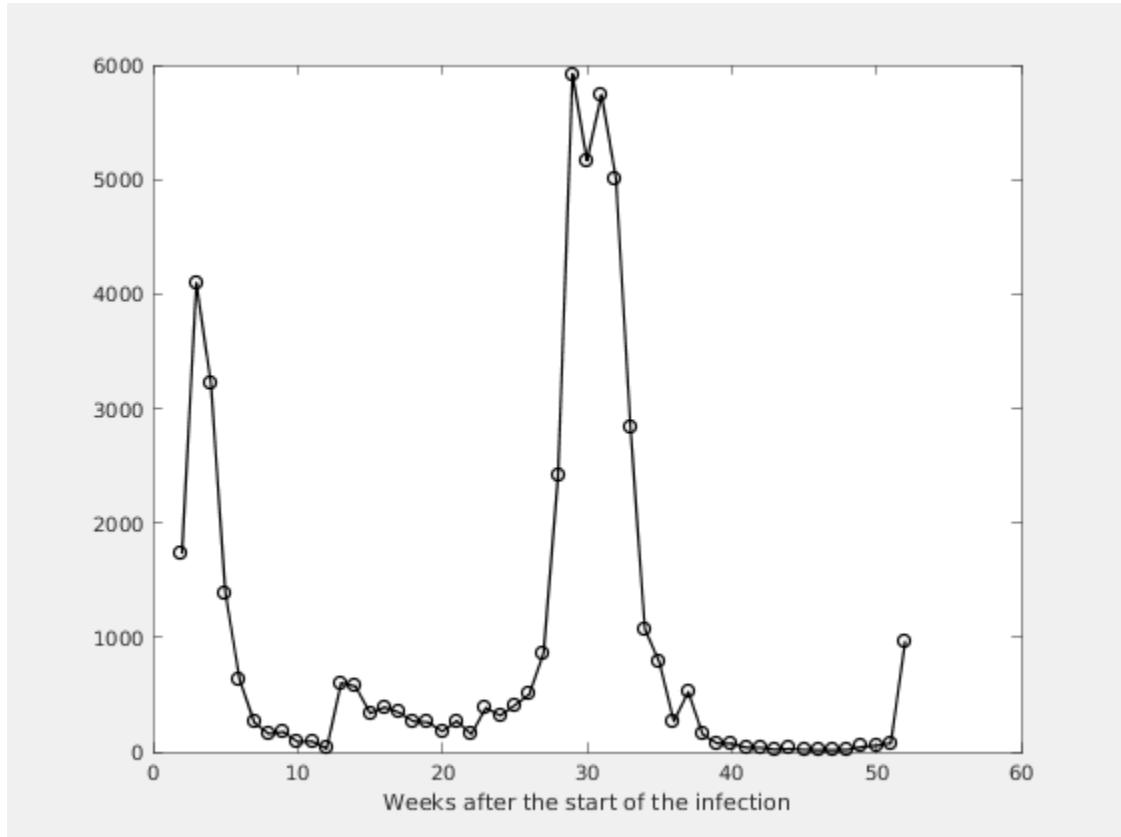
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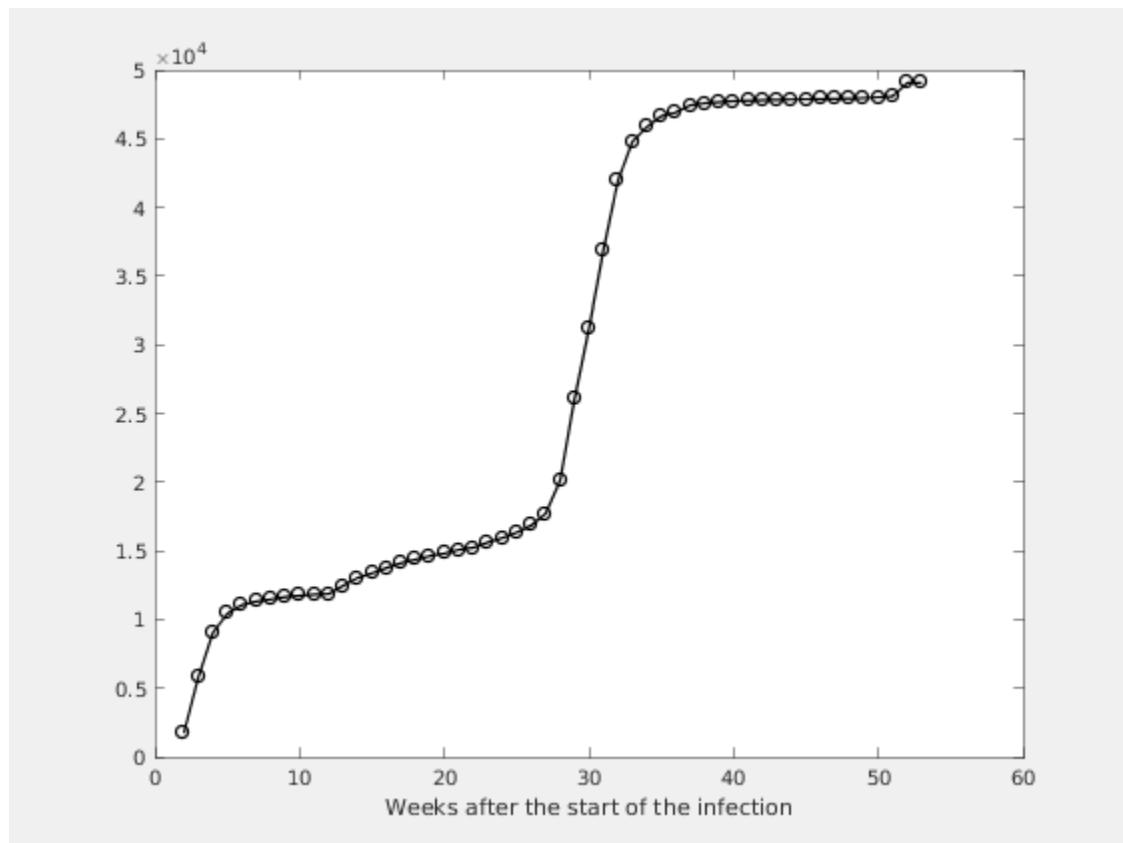
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# Chapter 1. Observed Infection data

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



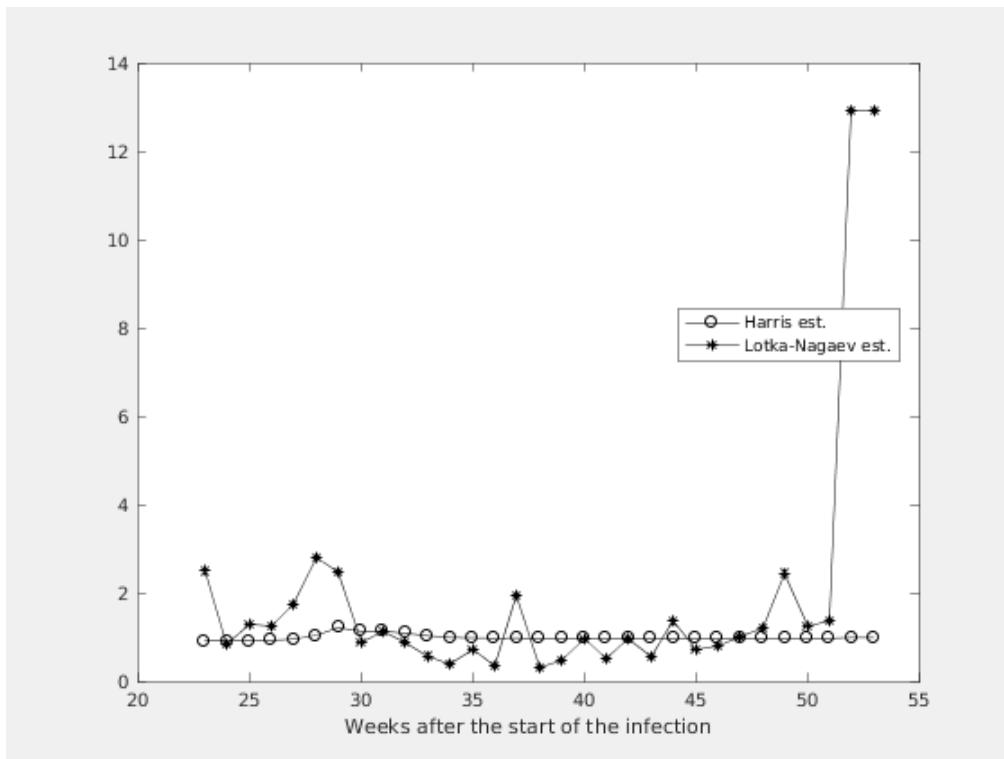
**Figure 1.2. Number of the total registered cases**



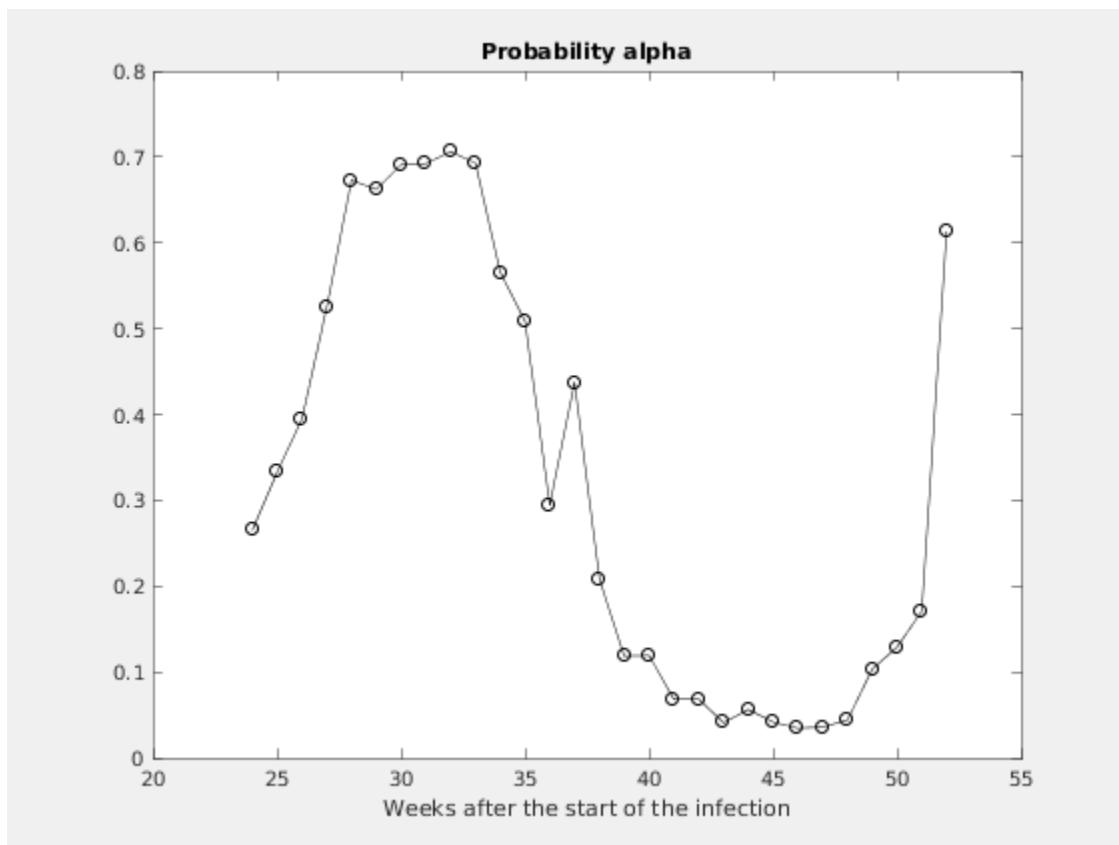
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# Chapter 2. Estimating of the main parameter and some predictions

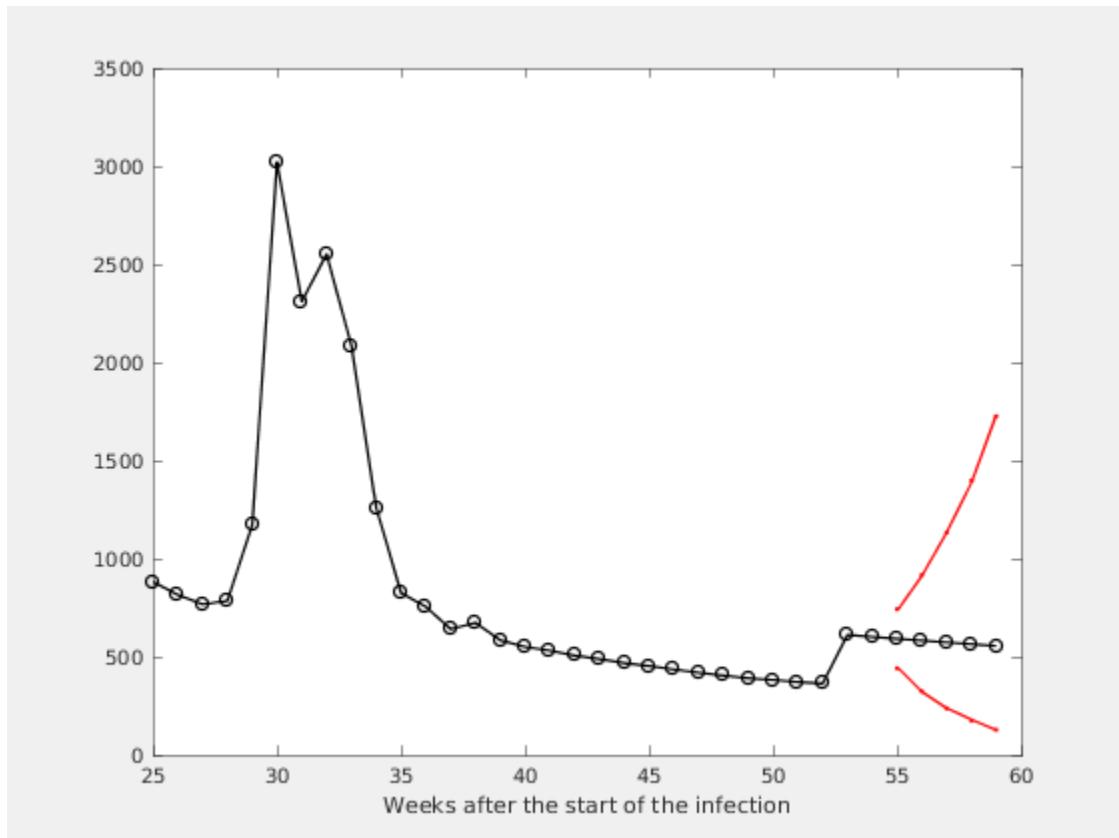
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



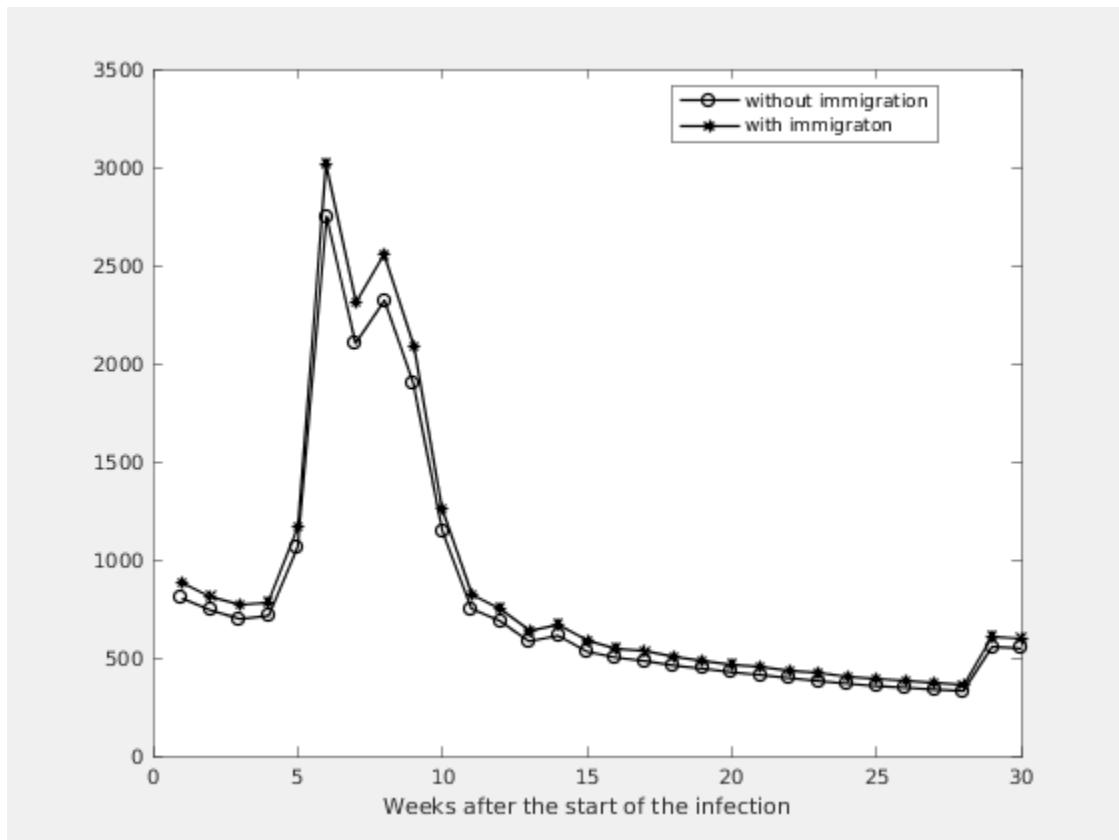
**Figure 2.2. Figure**



**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
<hr/>					
4	0.9648	0.7395	- 1.1902	0.0358	404
3	0.9651	0.7422	- 1.1879	0.0441	391
2	0.9655	0.7449	- 1.1861	0.1032	382
1	0.9842	0.7659	- 1.2025	0.1290	372
0	0.9842	0.7683	- 1.2001	0.1711	363