

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

**Argentina - week 53**

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### **Abstract**

The results presented here are obtained using the methodology proposed in the paper <https://arxiv.org/abs/2004.14838> for the country Argentina. 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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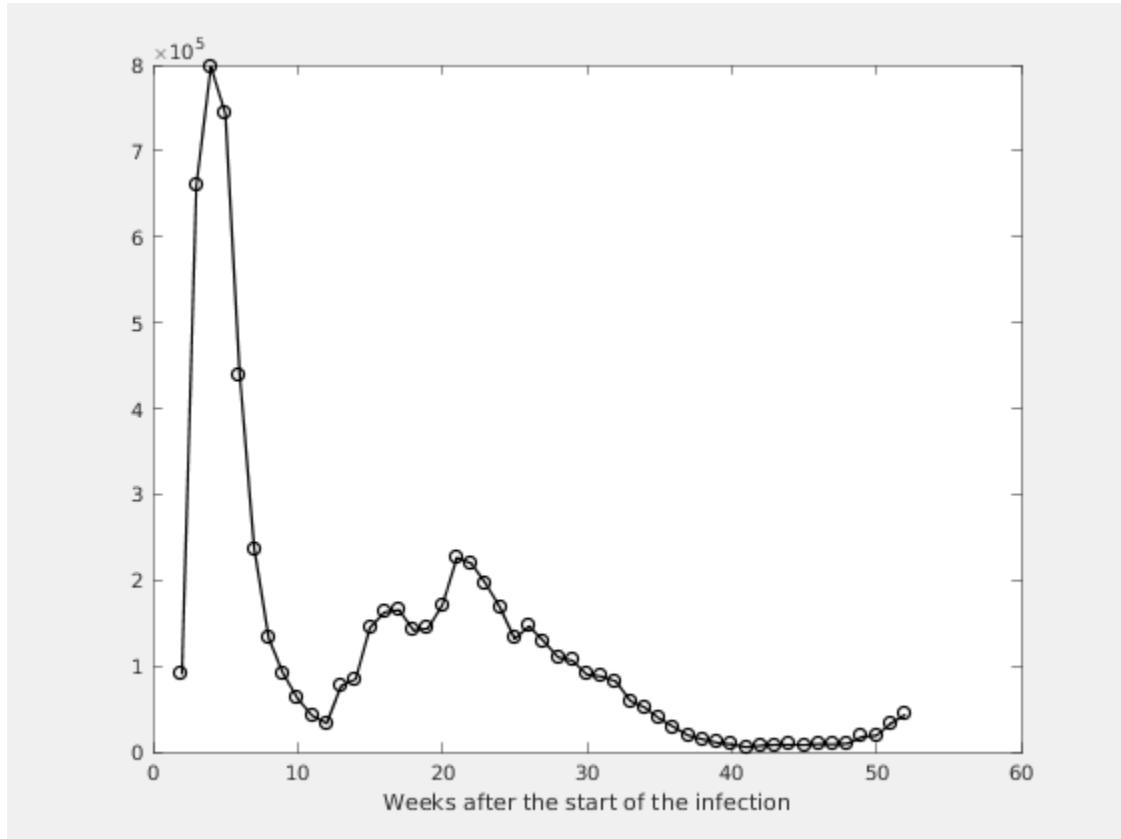
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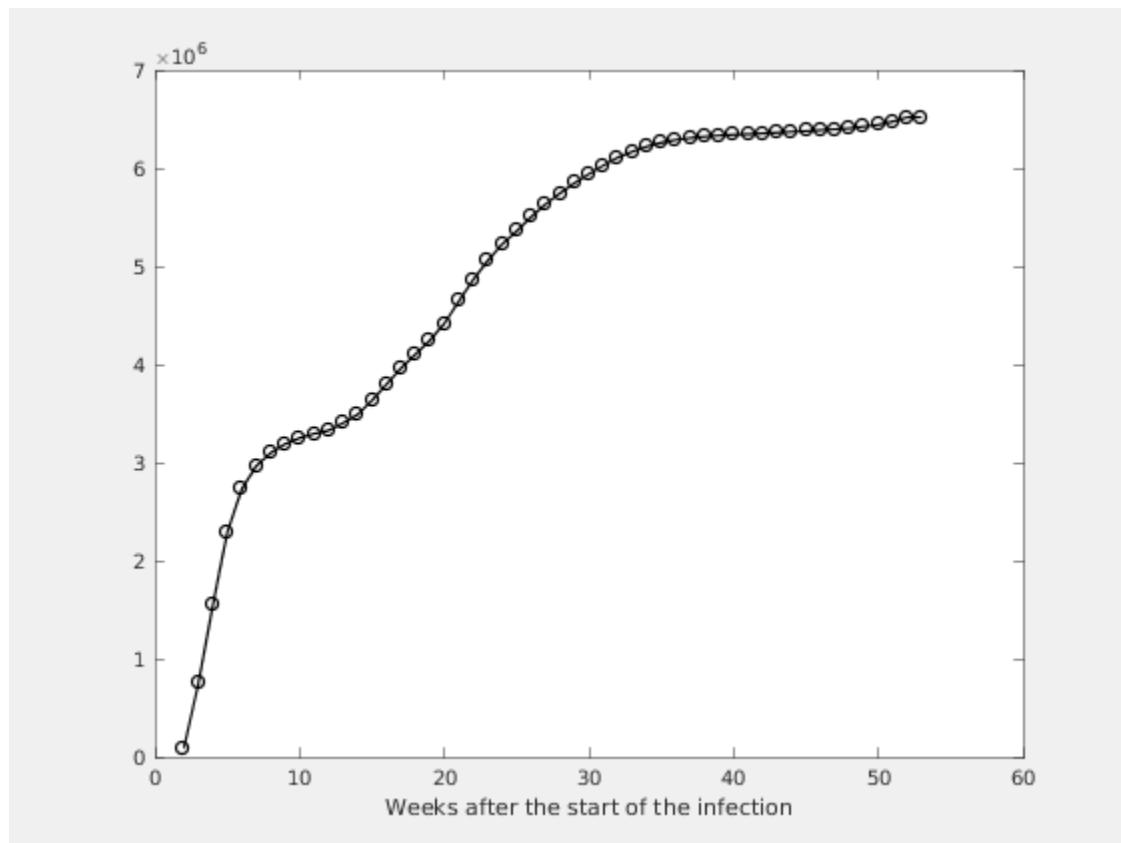
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# Chapter 1. Observed Infection data

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



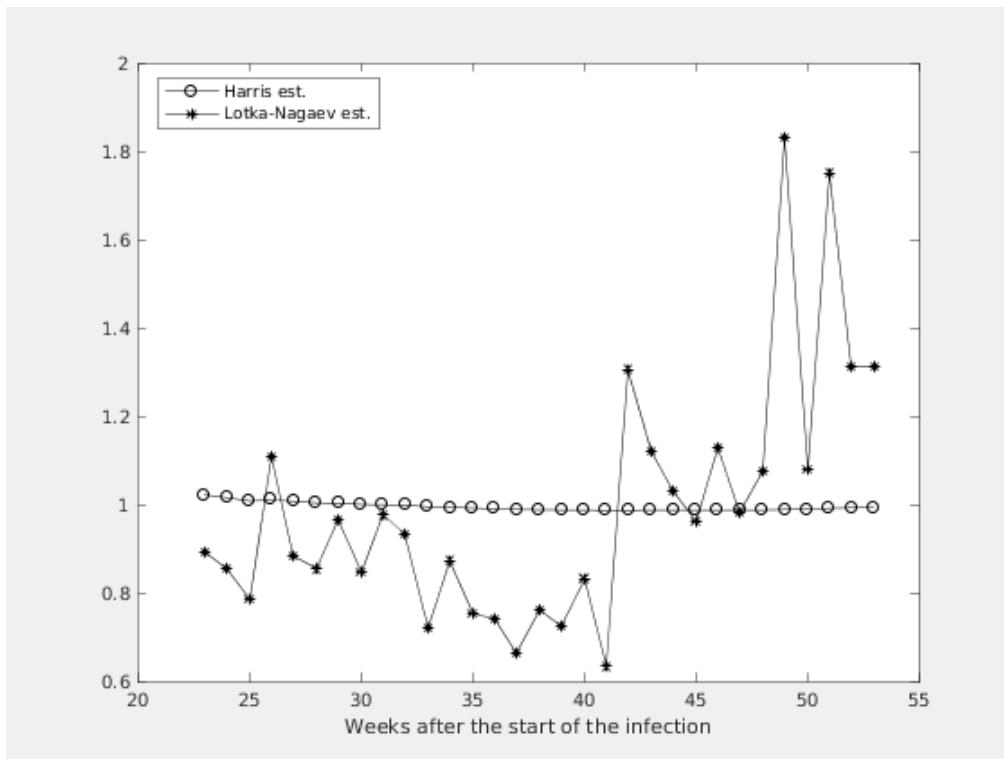
**Figure 1.2. Number of the total registered cases**



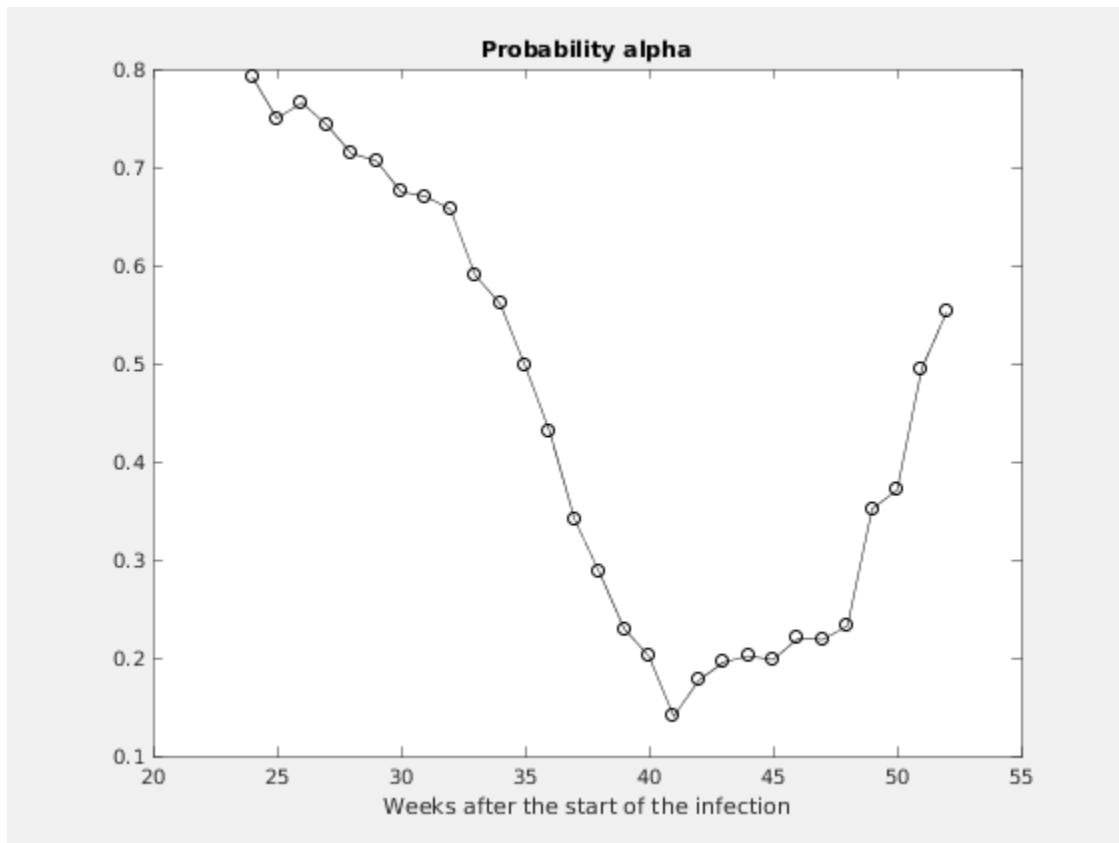
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# Chapter 2. Estimating of the main parameter and some predictions

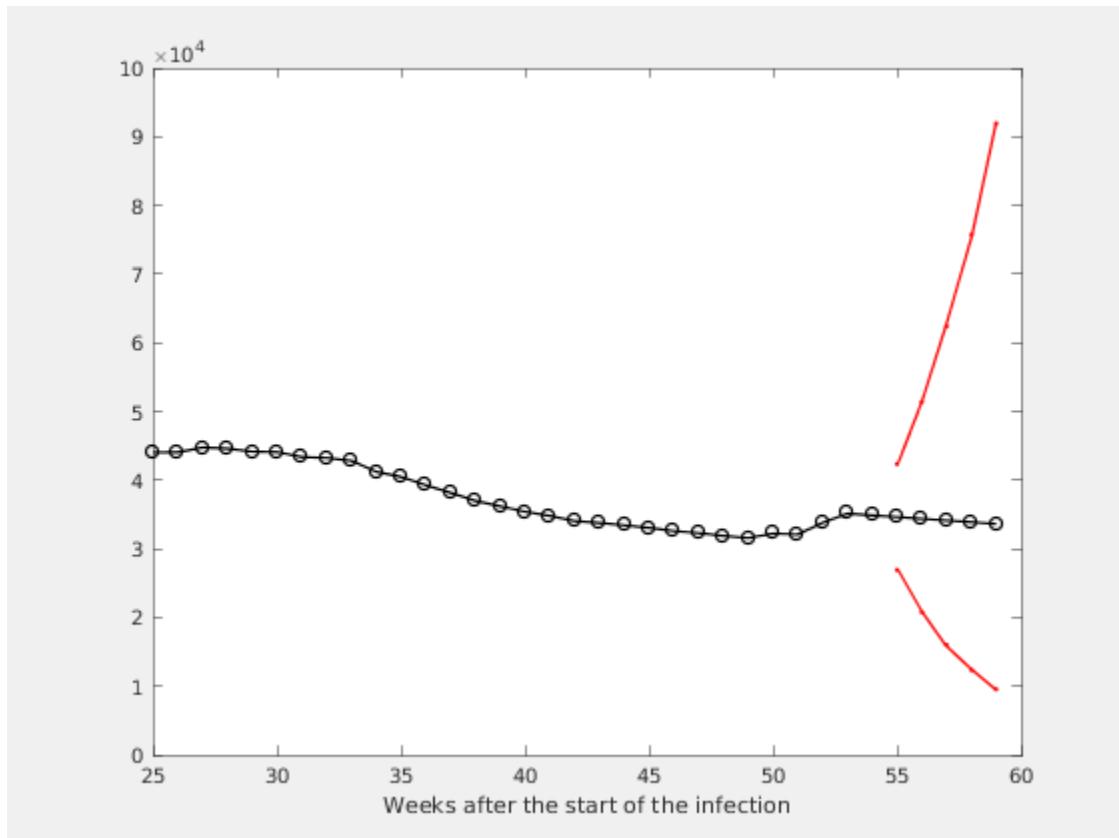
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



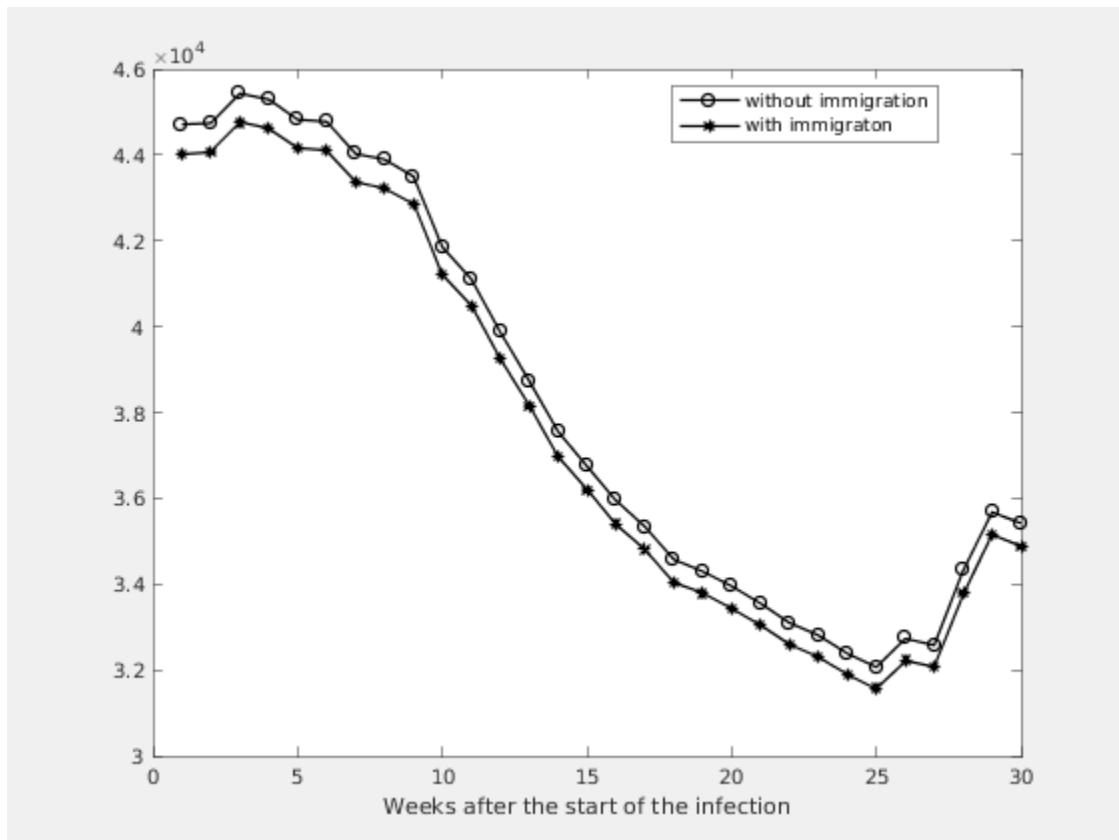
**Figure 2.2. Figure**



**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	0.9887	0.7546	- 1.2227	0.2183	31893
3	0.9889	0.7576	- 1.2202	0.2326	31584
2	0.9911	0.7623	- 1.2200	0.3520	32236
1	0.9928	0.7666	- 1.2189	0.3706	32085
0	0.9928	0.7689	- 1.2166	0.4946	33798
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