

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

**Brazil - 20201214**

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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 Brazil. 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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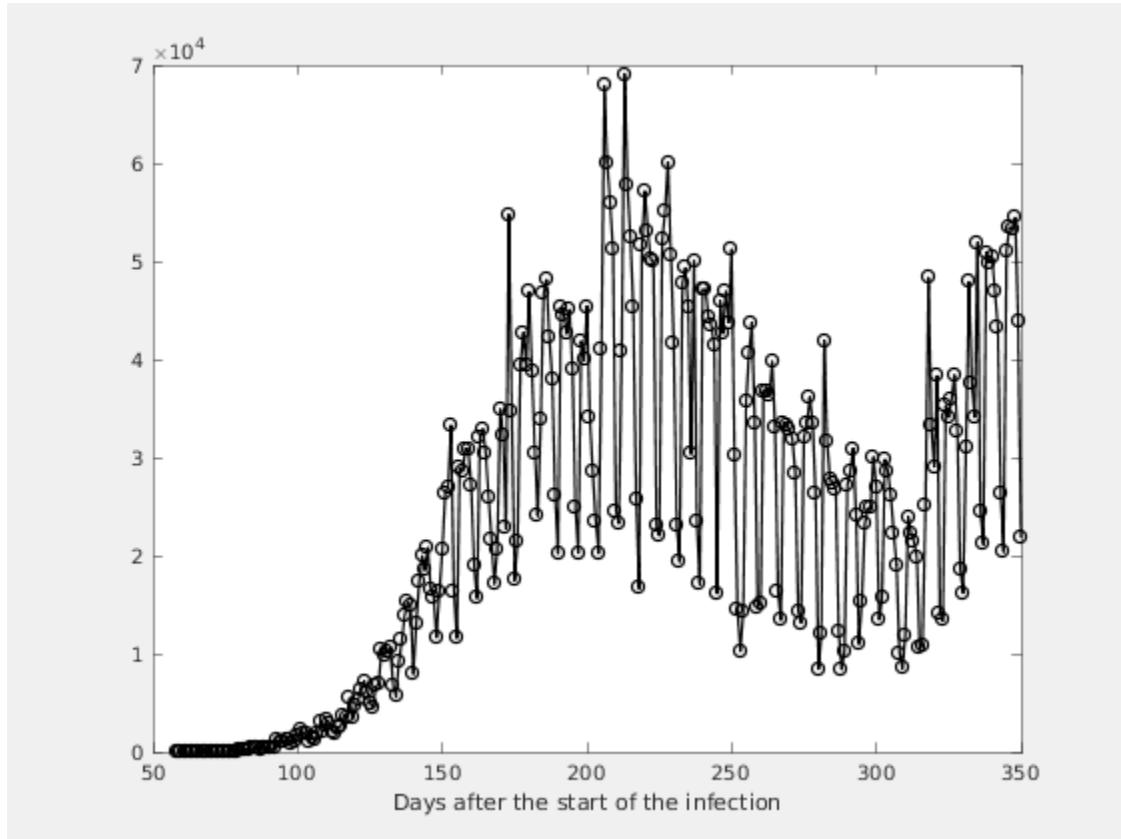
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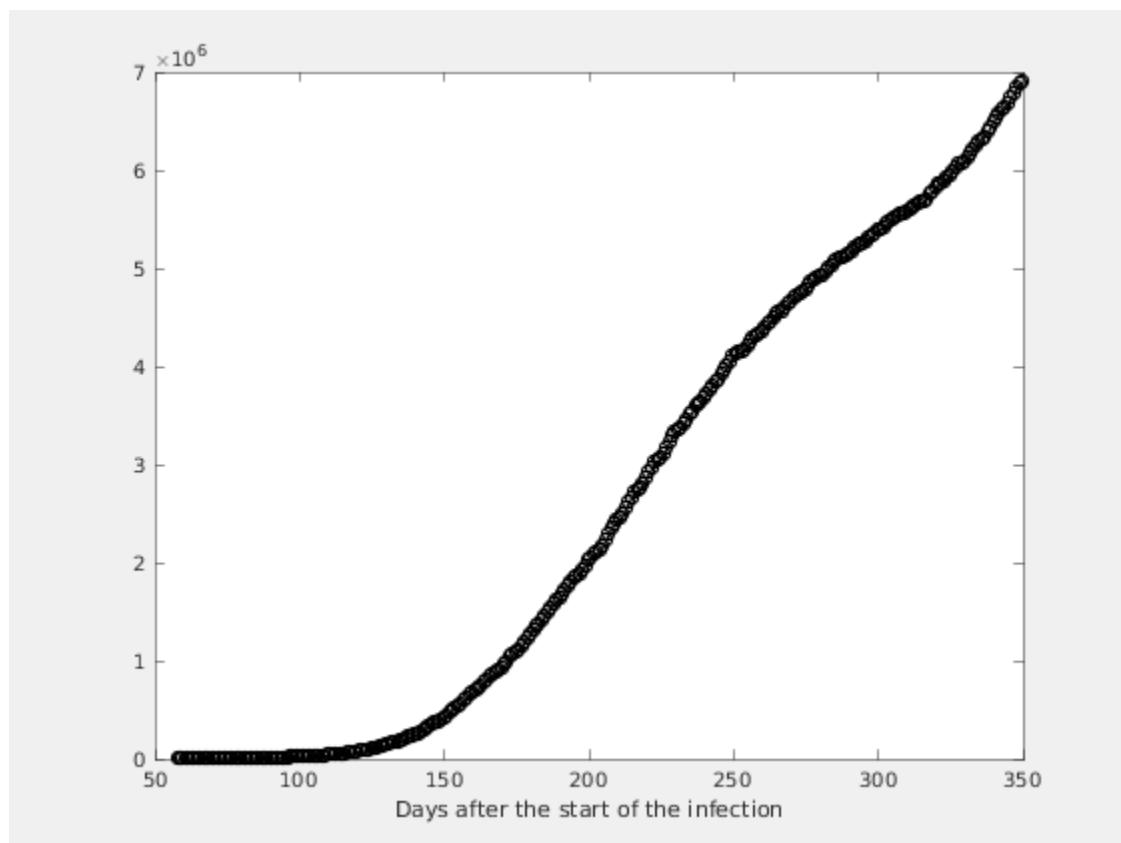
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# Chapter 1. Observed Infection data

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



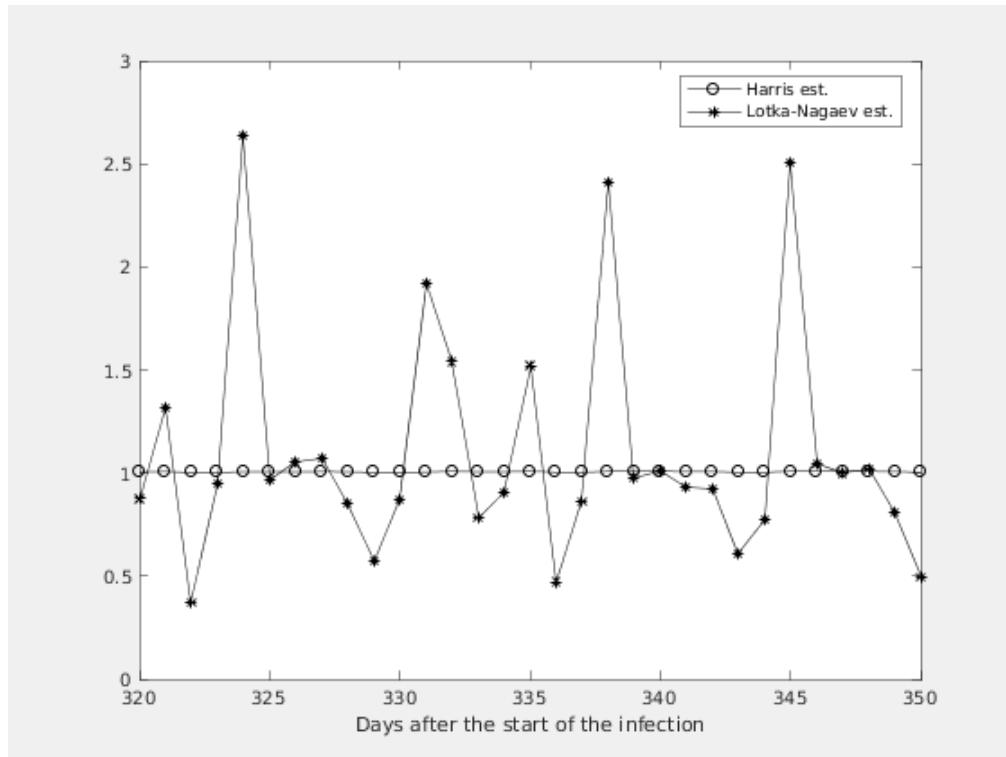
**Figure 1.2. Number of the total registered cases**



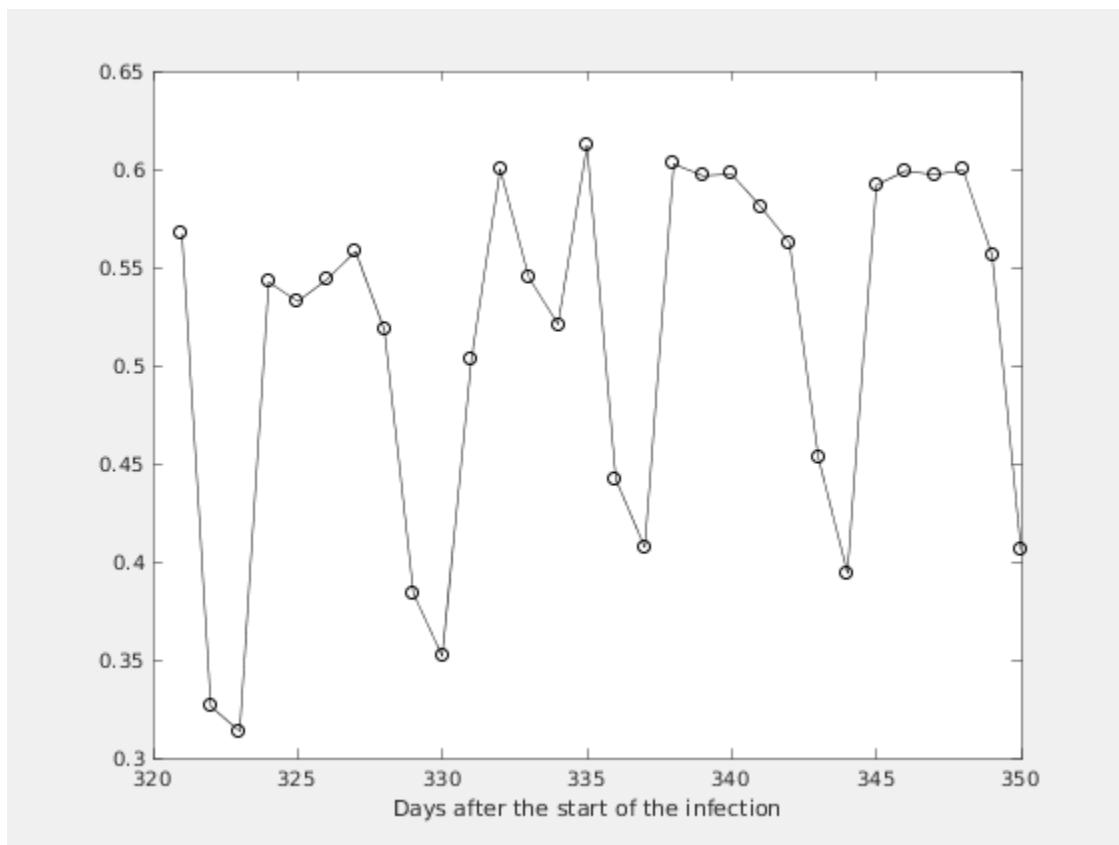
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# Chapter 2. Estimating of the main parameter and some predictions

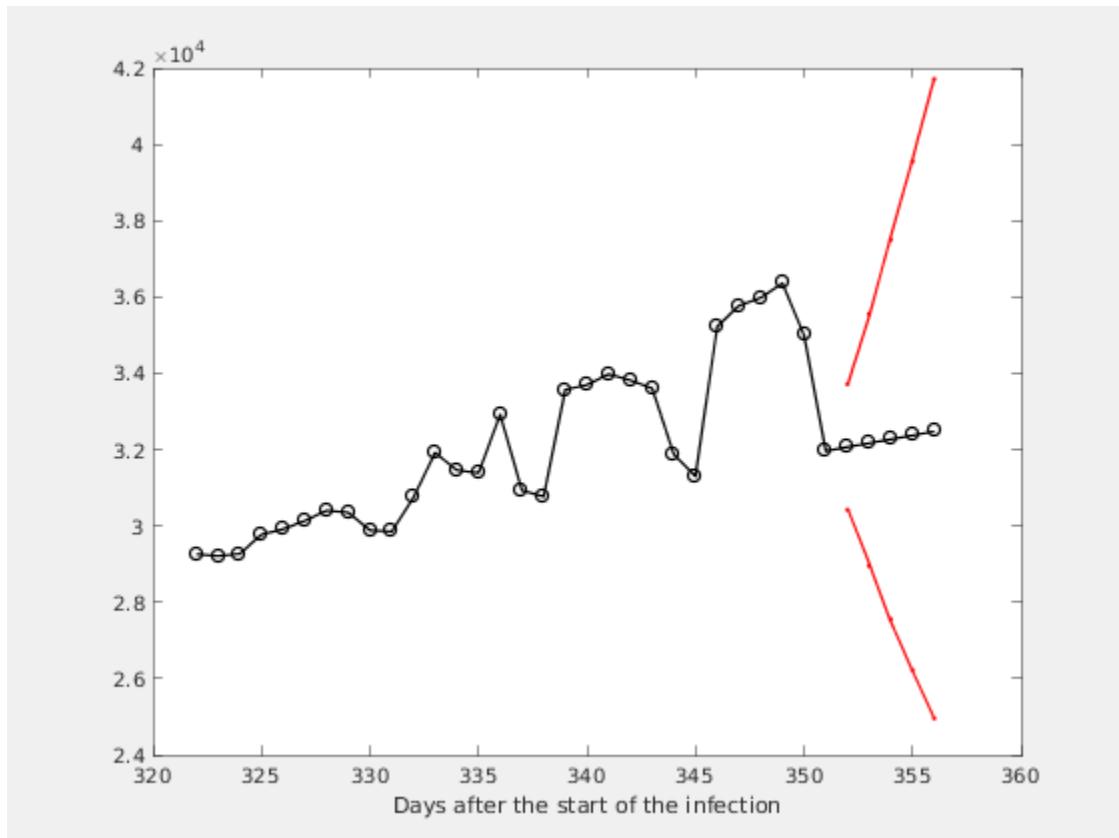
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



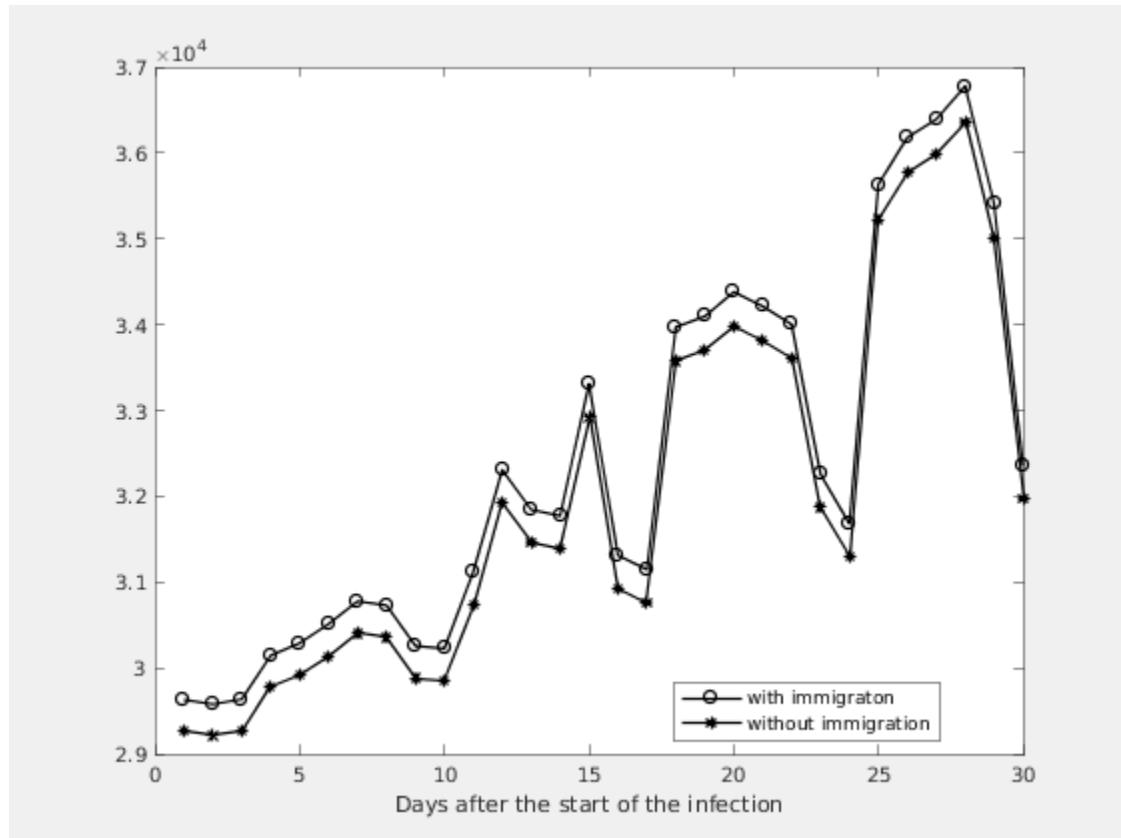
**Figure 2.2. Figure**



**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	M1	A1
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4	1.0080	0.9561	- 1.0600	0.3942	31300
3	1.0079	0.9552	- 1.0606	0.5919	35226
2	1.0080	0.9556	- 1.0604	0.5991	35769
1	1.0064	0.9543	- 1.0585	0.5972	35979
0	1.0032	0.9514	- 1.0550	0.5995	36362
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