

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

**Brazil - week 53**

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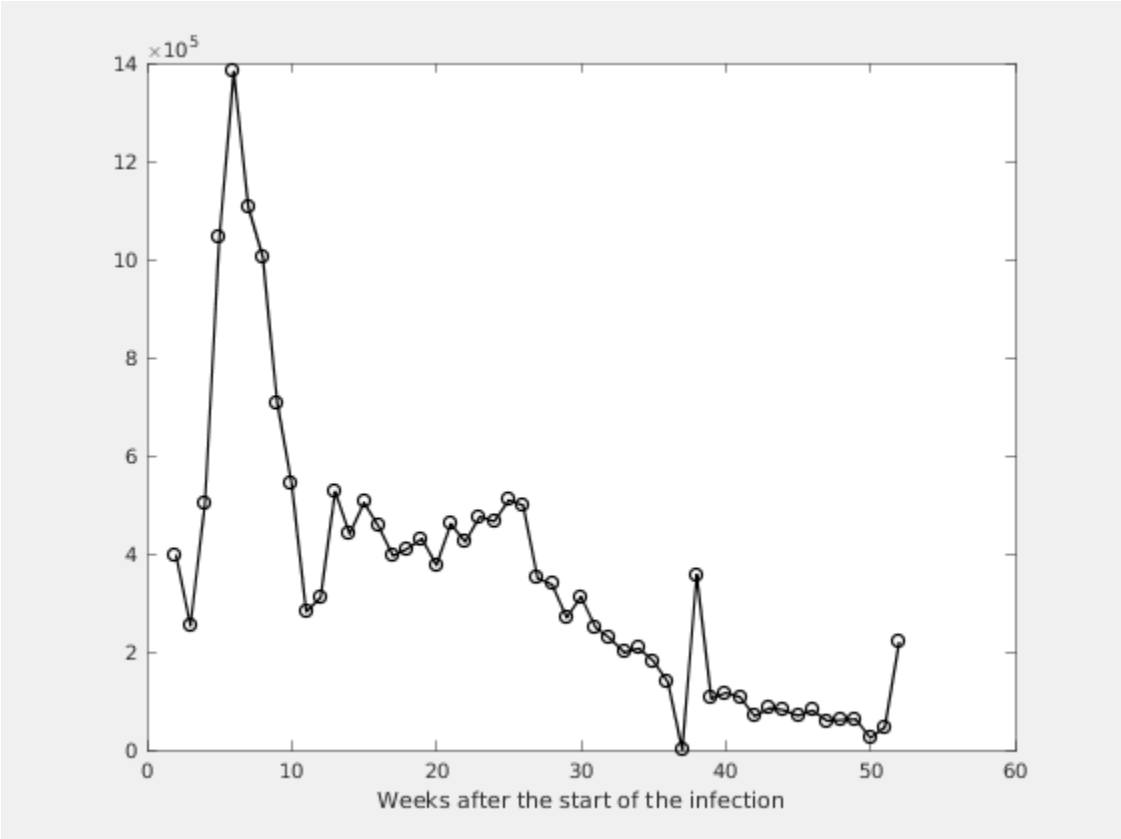
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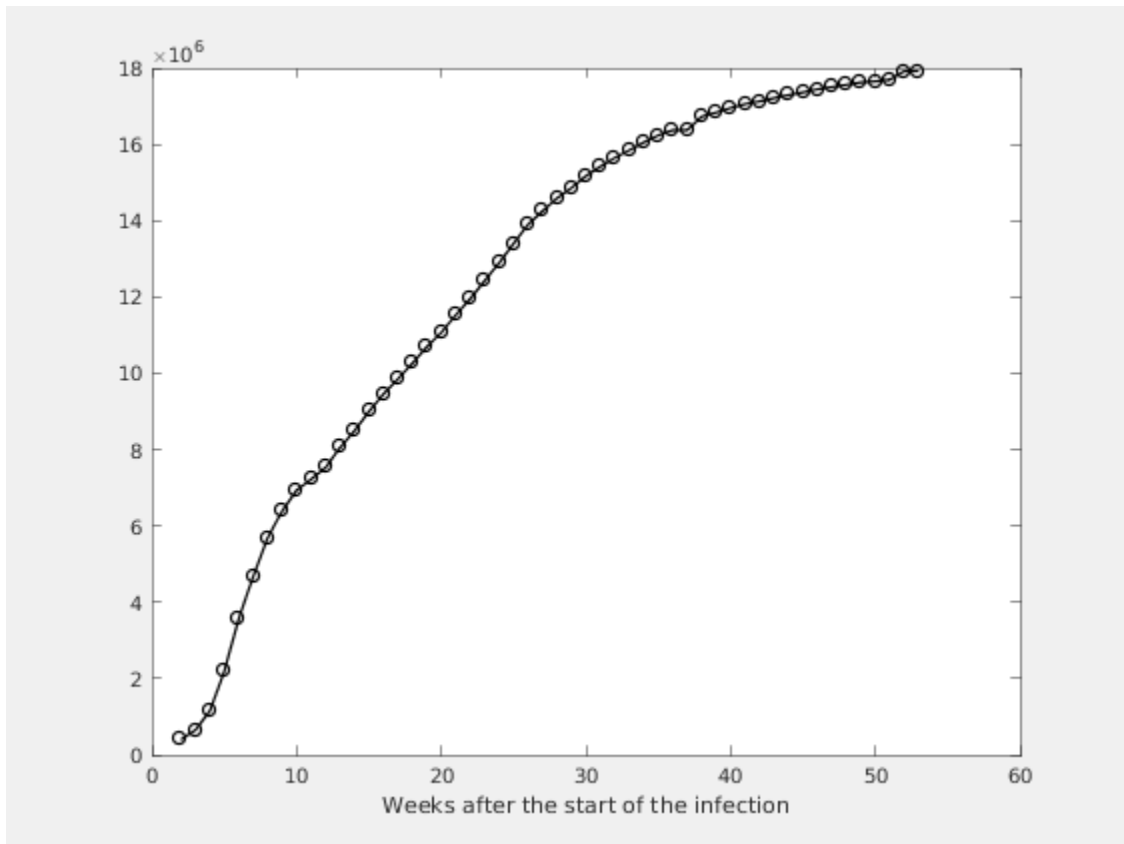
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# Chapter 1. Observed Infection data

Figure 1.1. Number of the 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

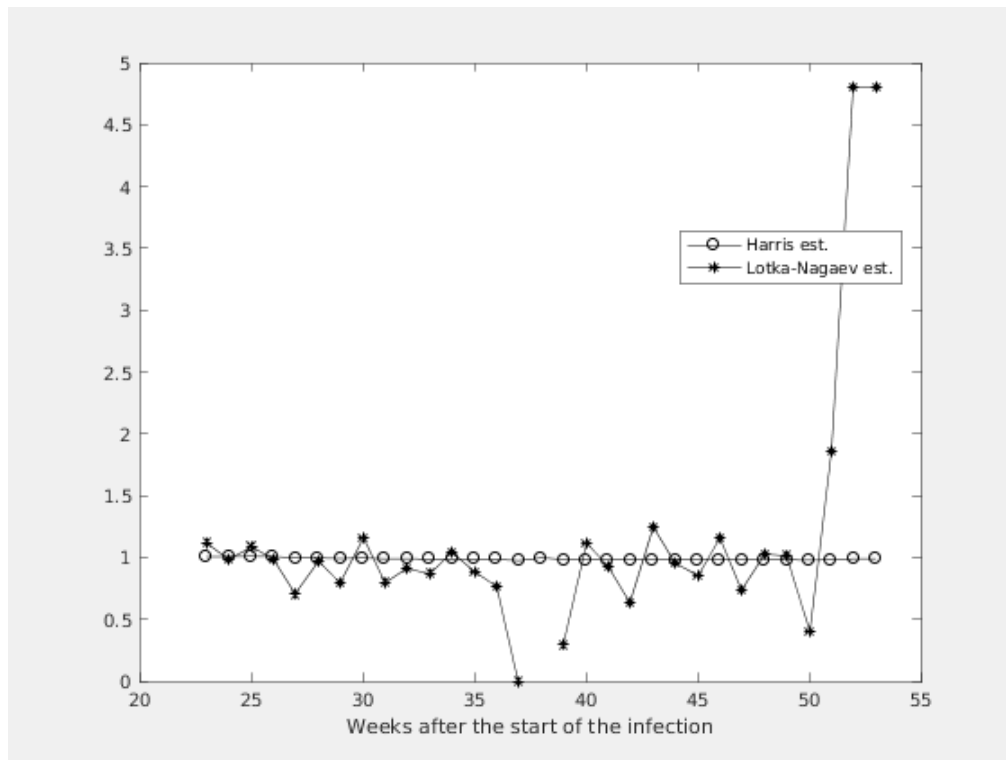
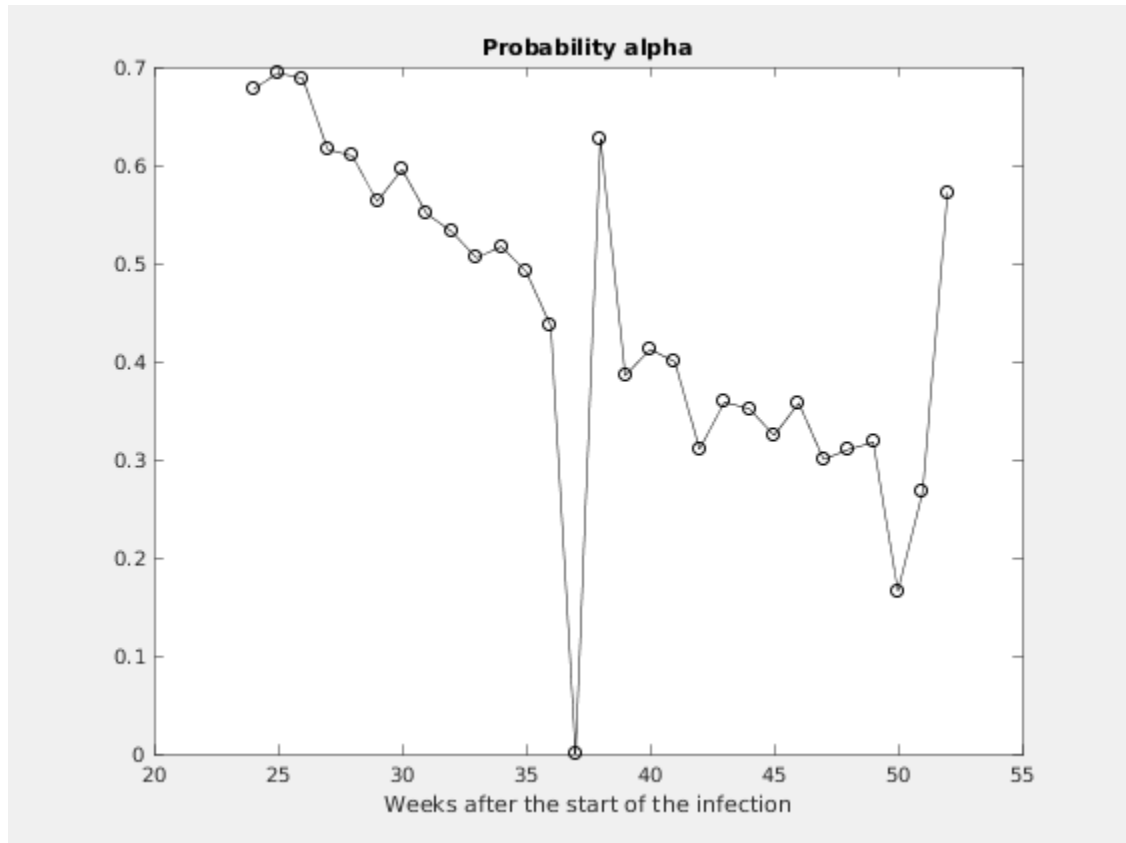
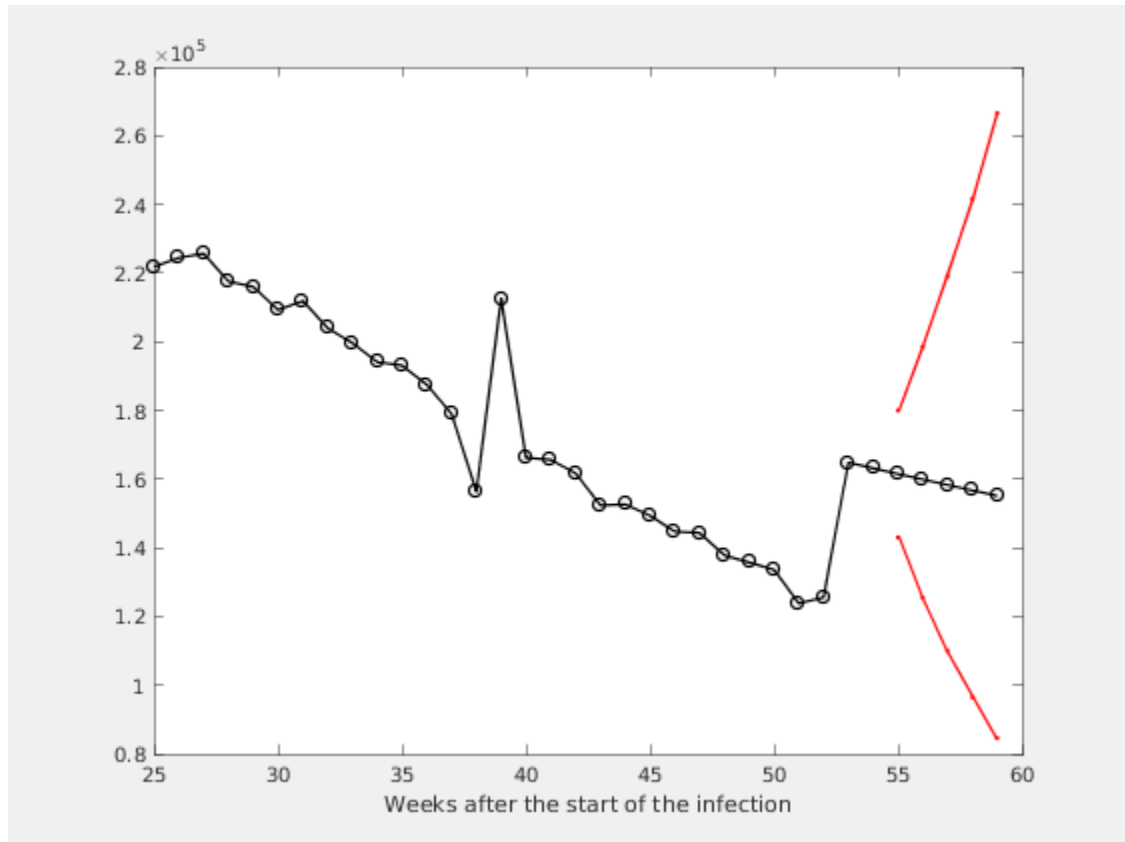


Figure 2.2. Figure

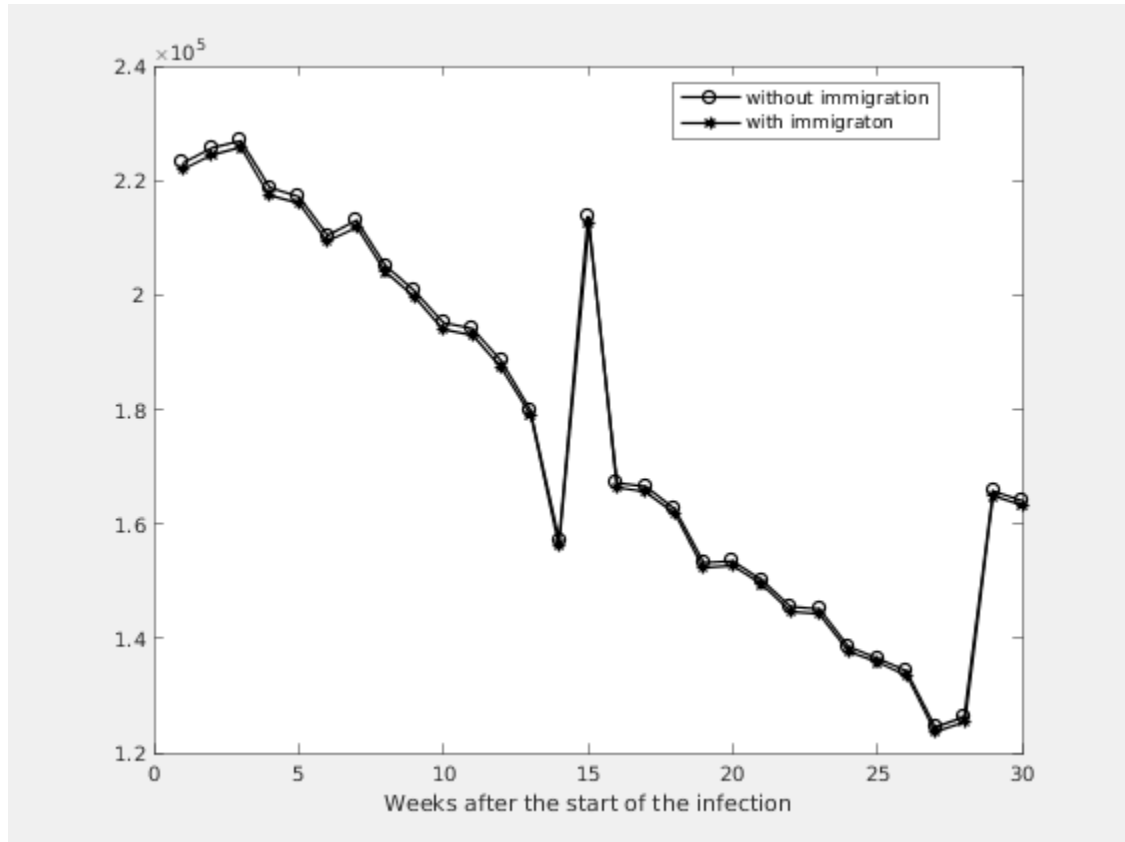




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

k	m	ci	alpha	A1	M1
4	0.9809	0.8764 - 1.0854	0.3001	137828	138581
3	0.9788	0.8757 - 1.0820	0.3101	135760	136500
2	0.9801	0.8782 - 1.0819	0.3176	133615	134344
1	0.9900	0.8889 - 1.0911	0.1662	123830	124506
0	0.9900	0.8896 - 1.0904	0.2680	125567	126252