

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

**TrinidadAndTobago - 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 TrinidadAndTobago. 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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## **Table of Contents**

1. Observed Infection data .....	1
2. Estimating of the main parameter and some predictions .....	3

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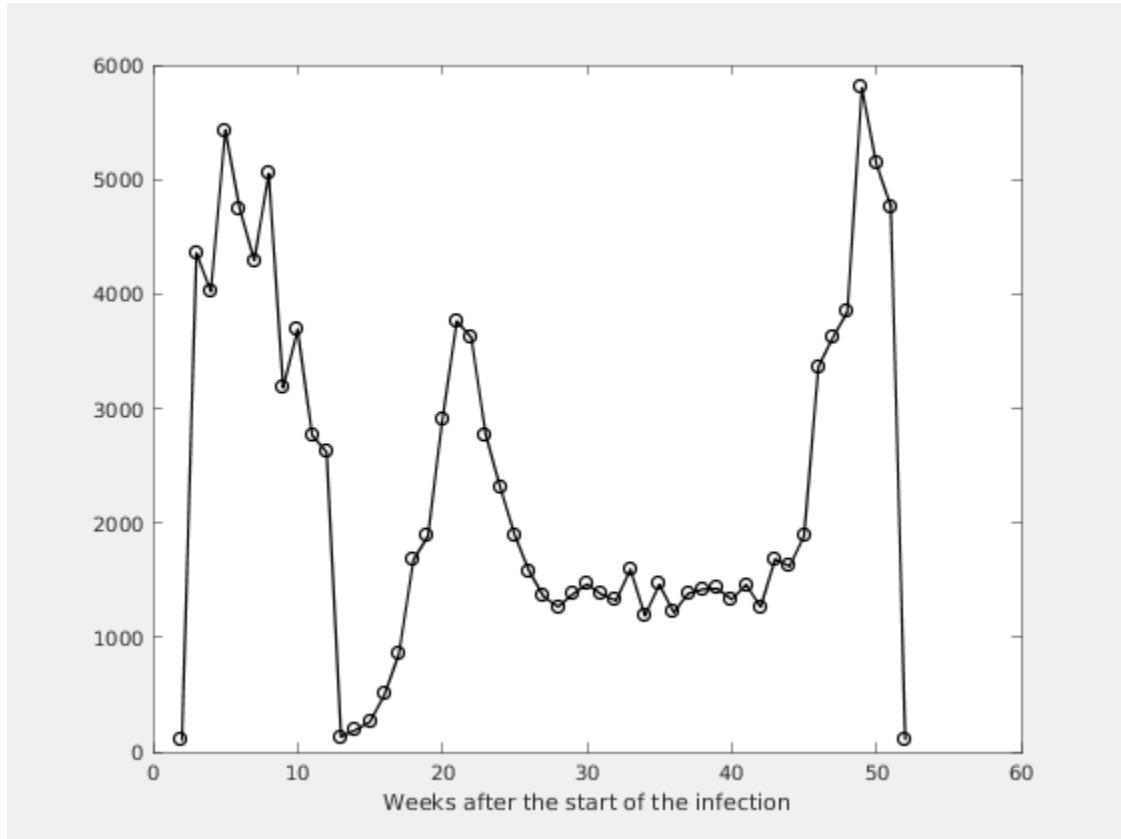
## List of Figures

1.1. Number of the weekly reported laboratory-confirmed cases .....	1
1.2. Number of the total registered cases .....	2
2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate .....	3
2.2. Figure .....	4
2.3. Expected number of the nonregistered infected individuals without immigrati-	
on .....	5
2.4. Expected number of the nonregistered infected individuals with immigration .....	6

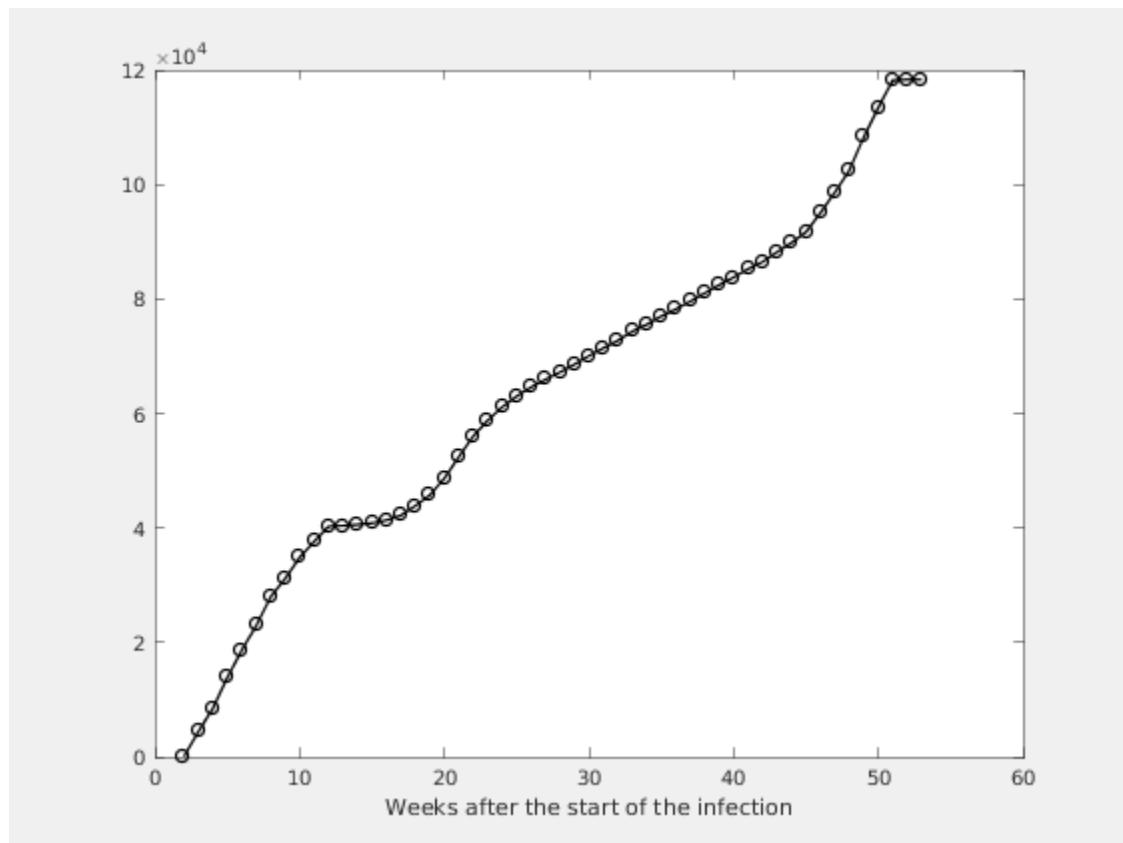
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# Chapter 1. Observed Infection data

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



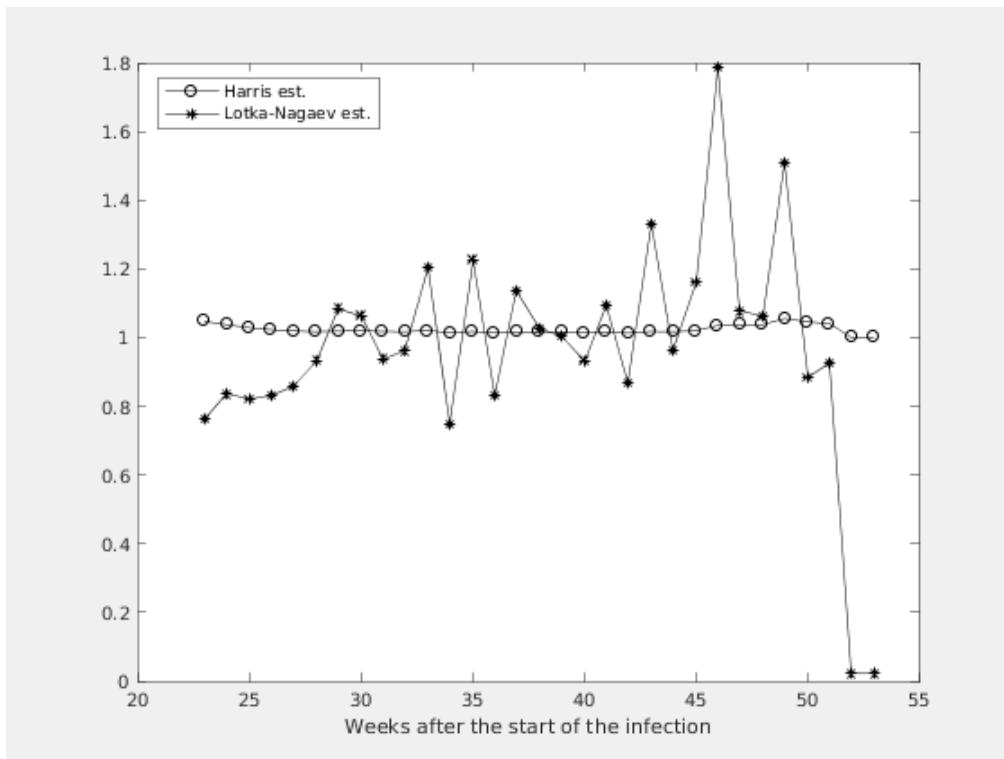
**Figure 1.2. Number of the total registered cases**



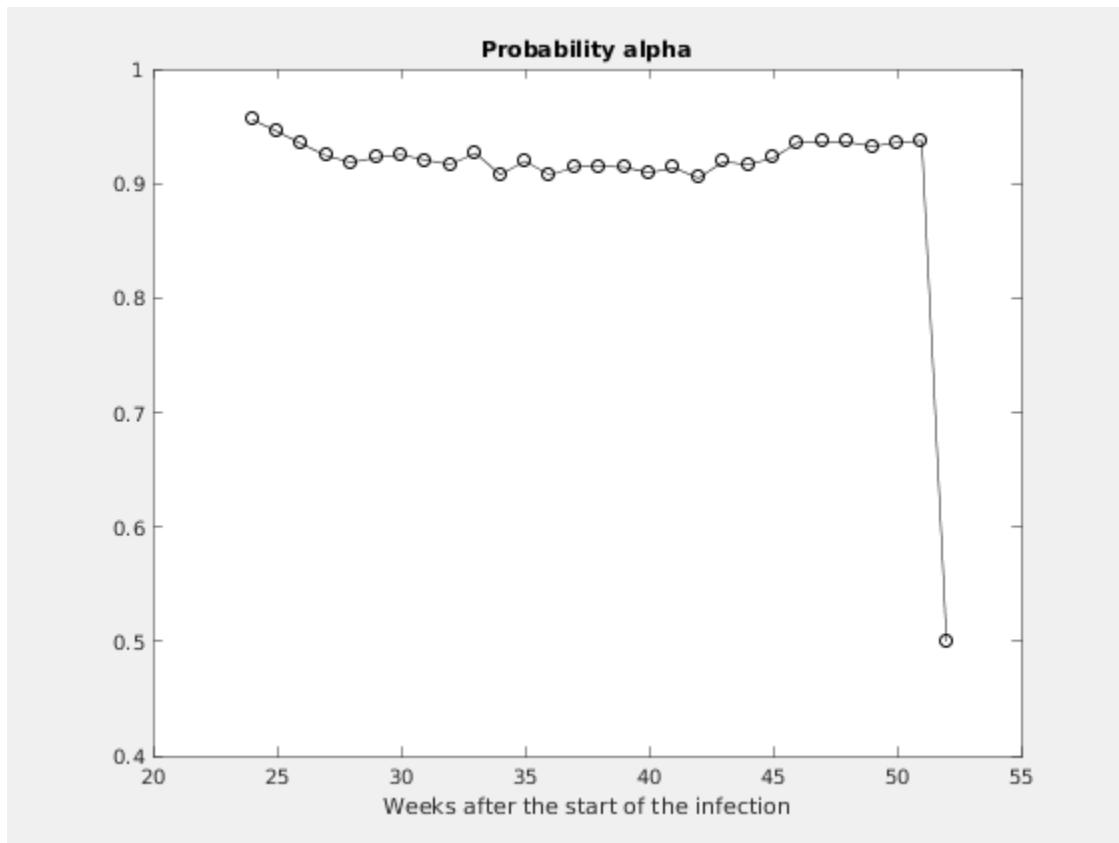
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# Chapter 2. Estimating of the main parameter and some predictions

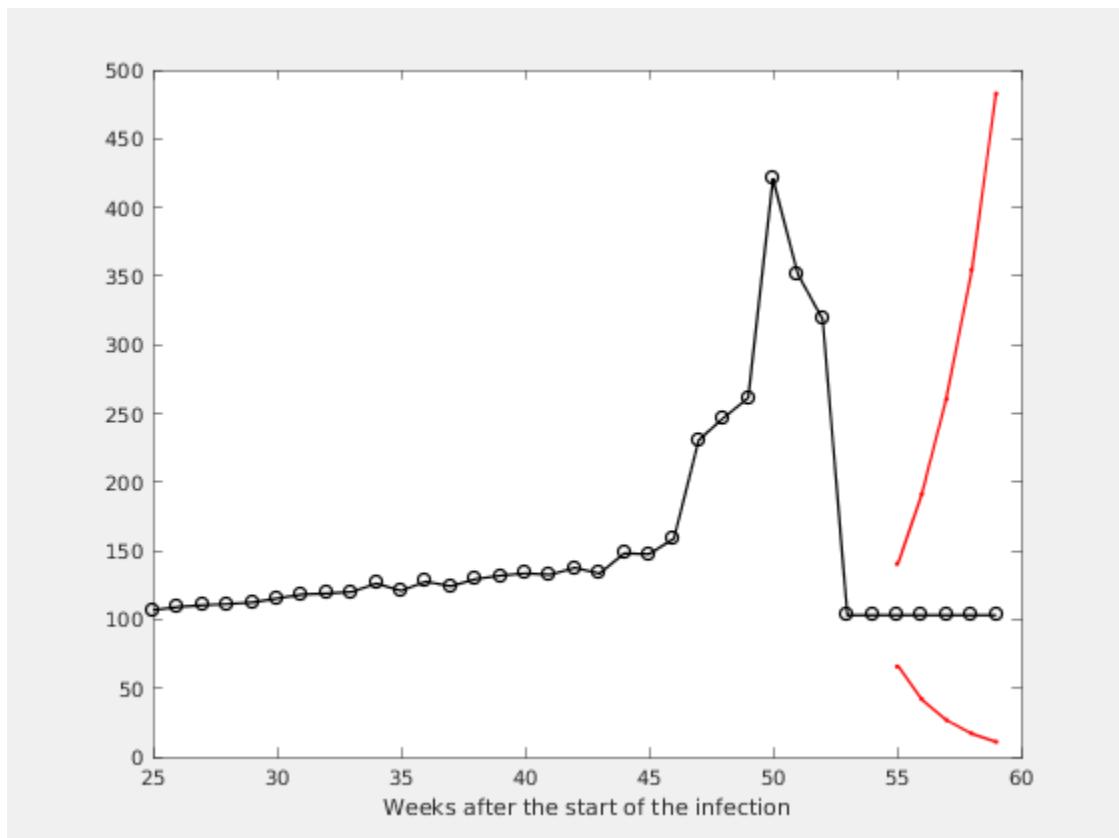
Figure 2.1. The Lotka-Nagaev and the Harris type estimator of the growth rate



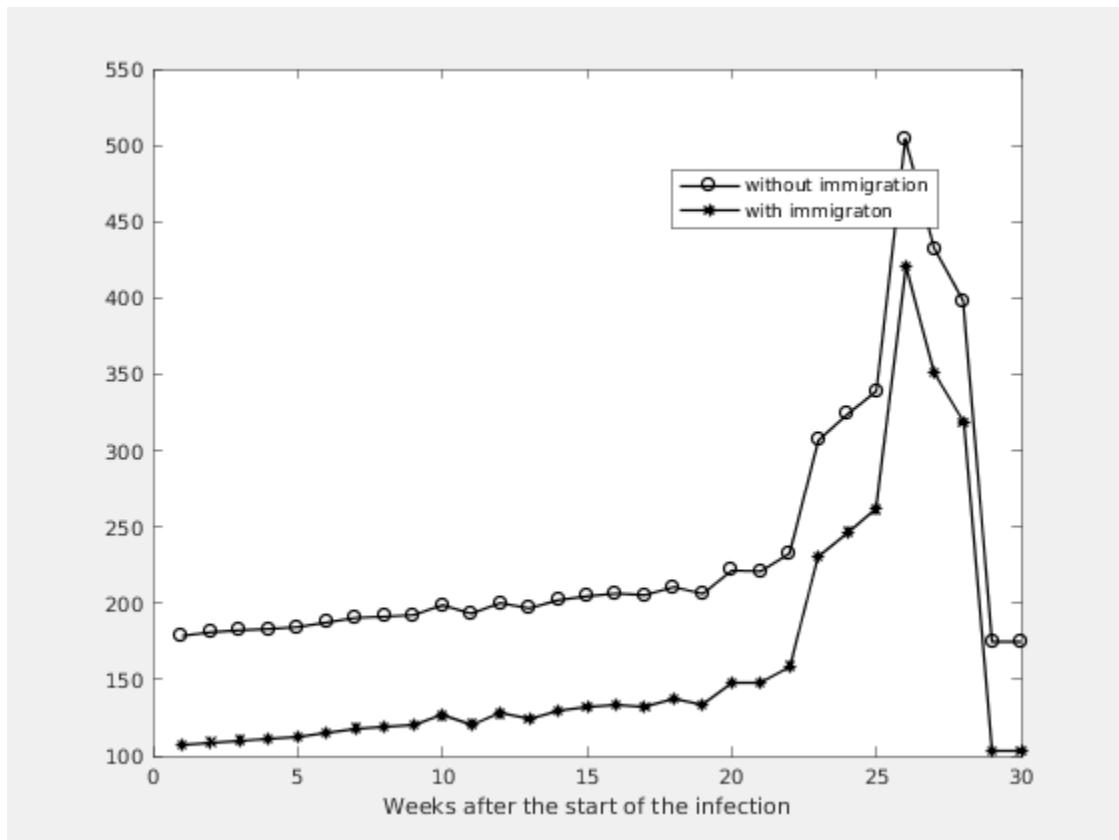
**Figure 2.2. Figure**



**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	1.0556	0.6367 - 1.4745	0.9362	247	324	
3	1.0465	0.6399 - 1.4532	0.9363	261	339	
2	1.0411	0.6456 - 1.4367	0.9324	421	504	
1	1.0001	0.6189 - 1.3812	0.9359	352	432	
0	1.0001	0.6314 - 1.3687	0.9373	319	398	