

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

**SaoTomeAndPrincipe - week 53**

**N. Yanev, V. Stoimenova, D. Atanasov**

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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 SaoTomeAndPrincipe. 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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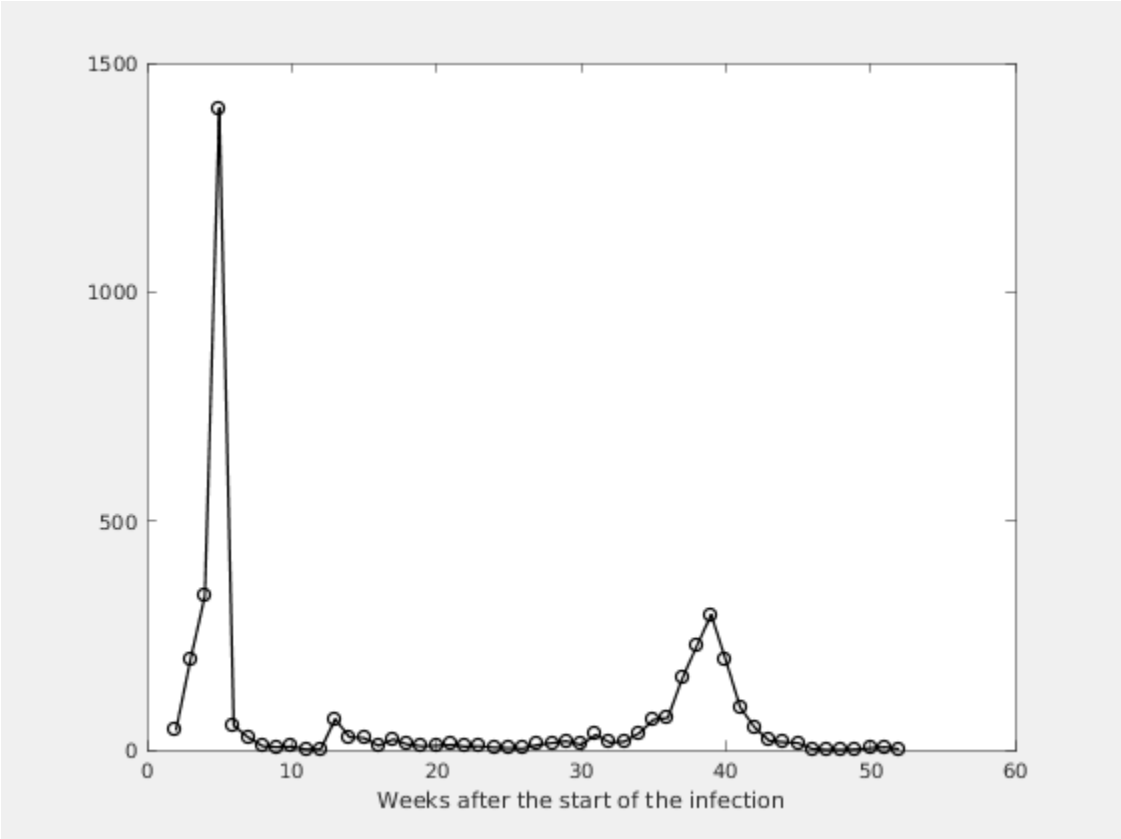
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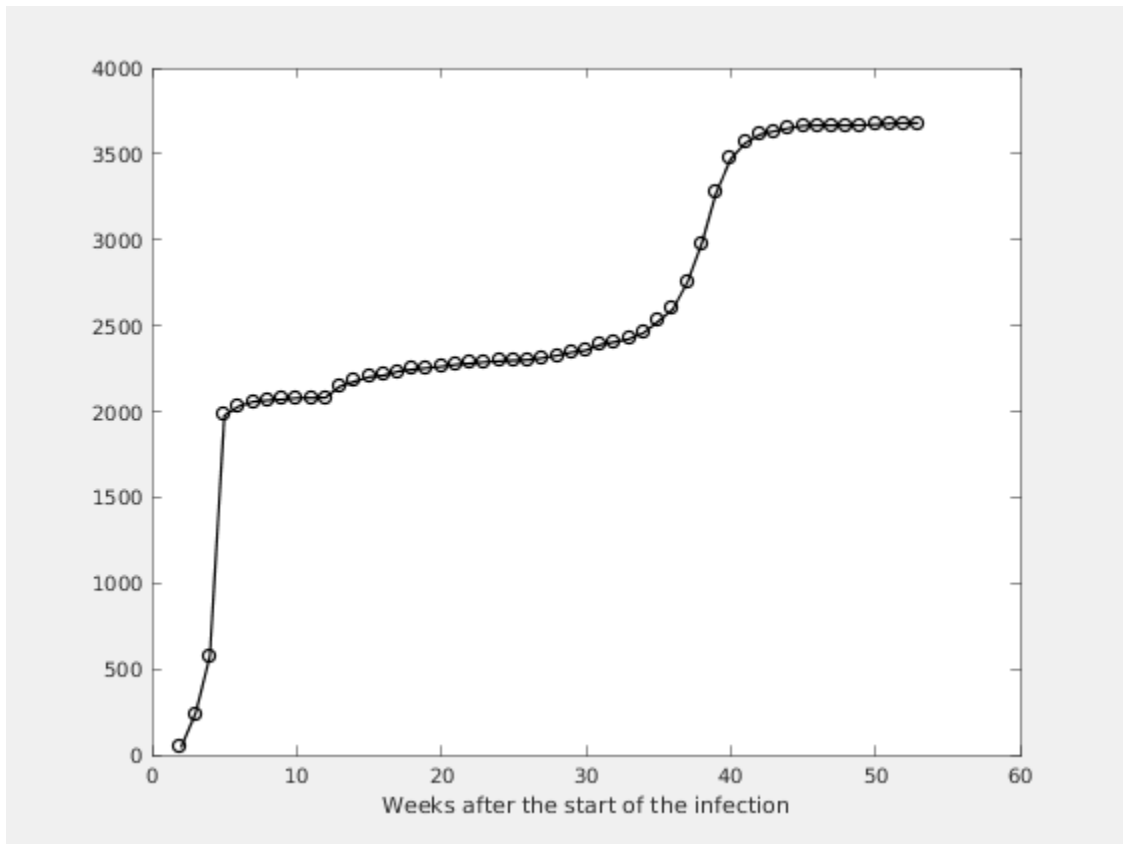
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# Chapter 1. Observed Infection data

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

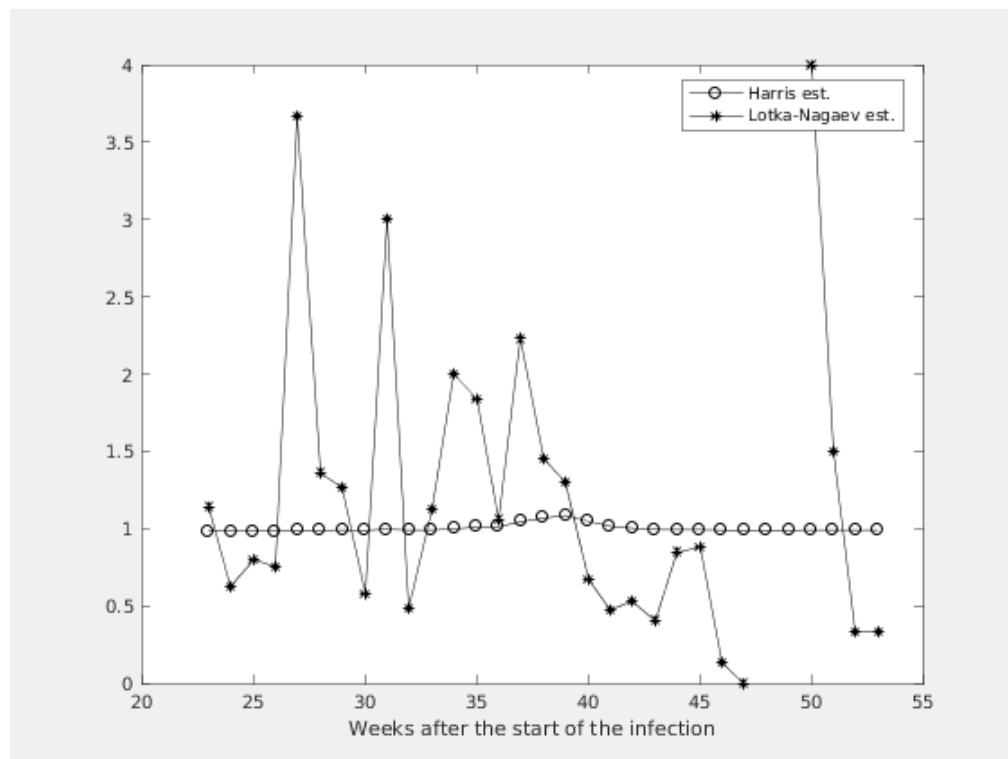
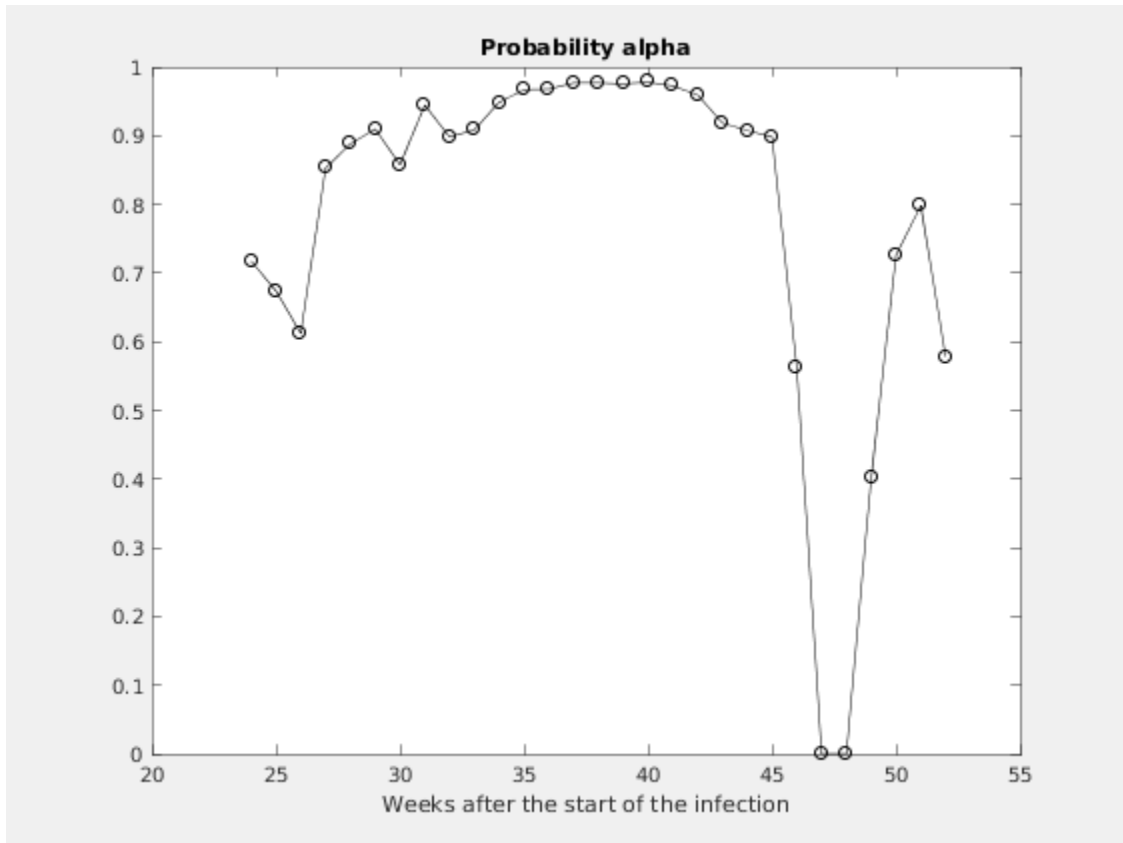
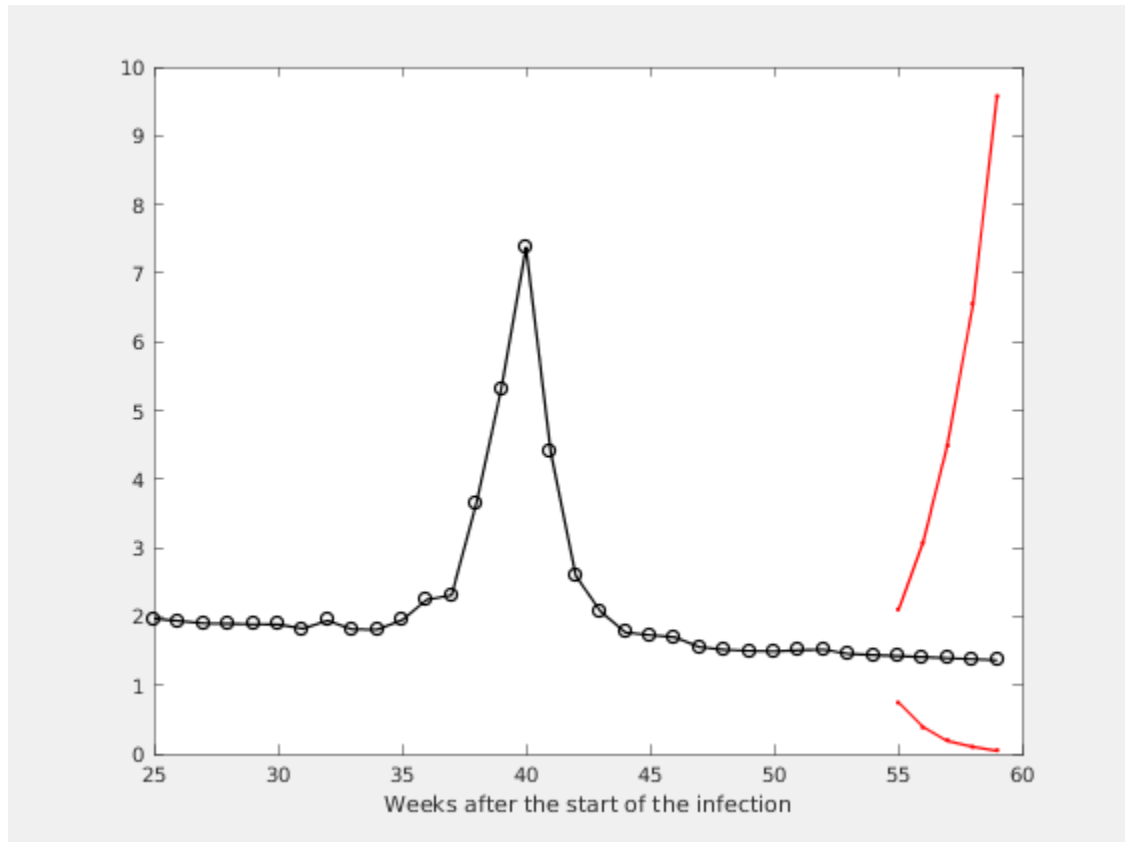


Figure 2.2. Figure

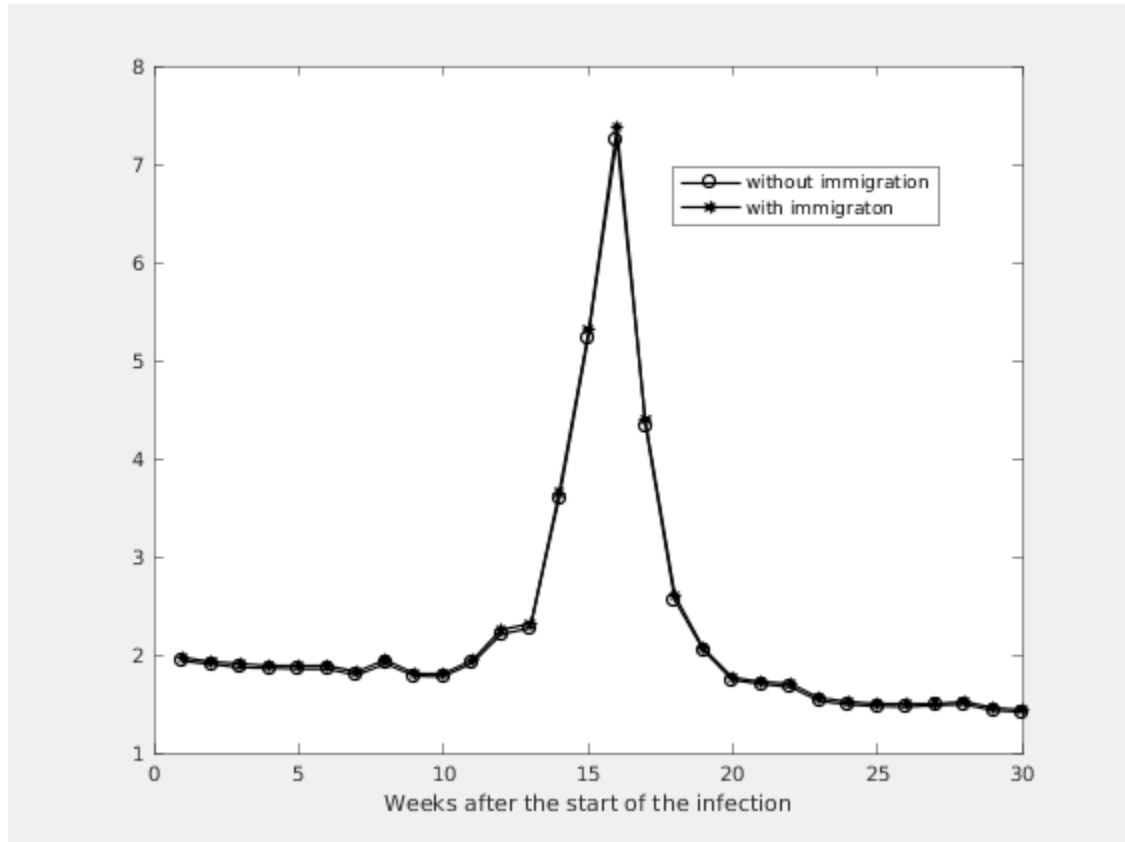




**Figure 2.3. Expected number of the nonregistered infected individuals without immigration**



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



**Estimation of the model parameters.**

| k | m      | ci              | alpha  | A1 | M1 |
|---|--------|-----------------|--------|----|----|
| 4 | 0.9888 | 0.4959 - 1.4817 | 0.0000 | 2  | 1  |
| 3 | 0.9896 | 0.4968 - 1.4825 | 0.0000 | 1  | 1  |
| 2 | 0.9902 | 0.5028 - 1.4776 | 0.4011 | 1  | 1  |
| 1 | 0.9891 | 0.5069 - 1.4713 | 0.7260 | 2  | 1  |
| 0 | 0.9891 | 0.5123 - 1.4659 | 0.7981 | 2  | 1  |