Title: Epidemiologic Study of Prince George’s County COVID-19 Data over the period February 1, 2020 – September 30, 2020

Author: Robert Bull and Alexander Klein

University of Maryland

Date: March 11, 2023

Venue: MATH 420 Mid-semester Project Presentation

1. **Introduction**

Our project focuses on the Prince George’s County infection and death rates during the COVID-19 pandemic. A predominantly democratic county, it faced relatively typical infection rates for the virus, but also had an interestingly high mortality rate.

We implemented the SIR and SEIR models described in class; we optimized these models to more effectively model and subsequently predict the infection rates and fatalities of COVID within the county.

1. **Background/Methods**

The SIR (Susceptible, Infected, Recovered) Model is a deterministic model that is intended to simulate states of an individual who has or has not been exposed to the disease we wish to model. It uses the three states listed above and presents the transitions as differential equations that we solved using the Euler method for initial value problems.

(Talk about SEIR here)

We used these models primarily because they are particularly inclined to model this exact situation: they are relatively simplistic and allow for the information gained from them to be optimized easily. By using the Euler method on the equations, we can quickly redo simulations when we need to change coefficients to better fit the measured data.

We parameterized the data using four general parameters: alpha (a), beta (b), gamma (c), and rho (p). Alpha is used to represent the infection period of the disease, or how long it stays transmissible after a patient becomes contagious. Beta we used to represent the number of daily contacts an infected individual has with susceptible people. In our data, we used R0 to represent b/a, or the number of individuals one infected person interacts with over their infectious period. Gamma was used to represent the fatality rate after infection. That is, what percentage of people died after being infected instead of recovering after their infectious period. Finally, we used rho to determine the undercounting factor of infections. Often, people would become infected with COVID and not report the illness to authorities or counting officials. Thus, they would be data points we would not know about, and we would have to scale the modelled infections down to represent that. We parameterized this model by creating an overall accuracy function, J, which includes differences between our predictions and measured data points.

(Presents Models and Algorithms designed for this problem.

Explanation of why the models/algorithms are the way they are.

Explain how these models are parametrized. What do these parameters mean.)

1. **Results**

Describe what results you obtained.

1. **Discussion**

Describe what these results mean.

**References**

Cite any source you used for your report.

PG Demographics: <https://elections.maryland.gov/pdf/vrar/2022_10.pdf>