## **Table of Contents**

```
% CSCE 666
% COVID-19 Project
% Iterative parameter estimation
% The SIRD model estimates deaths from an epidemic using an ODE.
% The solution to the ODE is parameterized by:
% r0 : reproduction number
% This script finds the transmission rate which best fits the SIRD
% to real data for 210 counties in the USA using an iterative
approach.
% The resulting r0 for each county is saved in the file 'r0_list.csv'
% for further analysis and feature correlation. Counties with poor
% statistics are omitted.
clear; clc;
```

## **Load Files, Prepare Data**

## **Evaluate Transmission Rates, Find Optimum for Each County**

Create a variable to store parameter values for each county

```
params = zeros(4, num_counties);
params(1,:) = FIPS;
```

```
r0_list = linspace(1.0, 3.5, 50);
trials = zeros(length(r0 list), num counties);
for county = 1:num_counties-1
    % Population of county
    d = mortality_rates(2, county);
    pop = population(2, county);
    init_cases = cases_data.(county);
    Iinit = init cases(2);
    fprintf('\nCounty: %g\nFIPS:
                                  %g\nM.R.: %f\n', [county,
 FIPS(county), d])
    % If ONLY optimizing transmission rate:
    for i = 1:length(r0 list)
        x = [r0_list(i) d county pop Iinit];
        trials(i, county) = mse_sir(x);
    end
    [~,ind] = min(trials(:, county));
    fprintf('r0:
                     %f\n', r0_list(ind))
```

## **Export Results**

```
r0_eval = zeros(3, num_counties);
r0_{eval(1, :)} = FIPS;
for county = 1:num_counties
   [mser,ind] = min(trials(:, county));
   r0 eval(2, county) = r0 list(ind);
   r0_eval(3, county) = mser;
end
tbd = [];
for county = 1:num_counties-1
    if r0_eval(2, county) == 1
        tbd = [tbd county];
    elseif r0_eval(2, county) == 2
        tbd = [tbd county];
    end
end
r0_eval(:,tbd) = [];
csvwrite('r0_list.csv', r0_eval)
```

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