## Math 420, Spring 2024 Second Team Homework: SIR and SEIR Models

Consider the dataset associated to your team. The problems are stated with Matlab indexing convention: from 1 to the length of the vector. The algorithms presented in class use the natural convention, where time  $t_0 = 0$  may be indexed by 0. Be careful to implement algorithms correctly, regardless of indexing convention on your platform.

**Exercise 1.** This problem asks you to implement and run the Meta-Algorithm for SIR Calibration using the time series of *cumulative number of detected infection* V(t) and the cumulative number of deaths Y(t). Use initialization S(0) = N, I(0) from the data set, R(0) = 0. Use the following parameters:  $T_{max} = 119$  (i.e., a total of 120 time samples),  $V_{min} = 5$ ,  $\tau_0 = 7$ ,  $p = 1, 2, \infty$  (check all three values),  $(c_I, c_Y) = (0, 1)$  and  $(C_I, c_Y) = (1, 1)$  (check both values). For the set  $\Omega$  write nested loops over  $\alpha$  and  $R_0$  by searching over the following intervals:

 $\alpha \in [0.05, 0.4] \ stepsize = 0.01 \ , \ R_0 \in [0.8, 2.2] \ stepsize = 0.05$ 

Recall  $\beta = R_0 \alpha$ , and N = Population is extracted from your data set.

- (1) Find and print the minimum of the objective function J. Print the optimal values of  $\hat{\alpha}, \hat{\beta}, \hat{R_0}$  and  $\hat{\gamma}, r\hat{h}o$  ( $\hat{\gamma}$  is obtained by implementing the minimization of  $||Y \gamma R_{sim}||_p$  over  $\gamma$ , whereas  $r\hat{h}o$  is obtained by minimization of  $||I \rho I_{sim}||_p$  over  $\rho$ ).
- (2) Visualize the 2D surface (function)  $(\alpha, \beta) \mapsto J = J(\alpha, \beta; \hat{\gamma}, \hat{\rho})$  at the optimal values  $\hat{\gamma}$  and  $\hat{rho}$  for each value of  $(\alpha, \beta)$ . You should obtain one plot for each p.
- (3) For  $(\alpha, \beta) = (\hat{\alpha}, \hat{\beta})$ , plot on the same graph the simulated  $I_{sim}$  and the preprocessed rate of detected infections  $I(t) = V(t+t_0+\tau_0) - V(t+t_0-\tau_0)$ . Plot on the same graph the predicted number of deaths  $Y_{sim} = \hat{\gamma}R_{sim}$  and the observed number of deaths Y(t)from your data set (make sure to align Y with the same first time  $t_0$ ). You should obtain two plots for each value of p.
- (4) Repeat the previous questions for each pair  $(c_I, c_Y)$ .

NOTE: the Euler scheme produces an estimate of the solution every h = 1/100 day time step. However the experimental data is measured **daily**. Hence you need to downsample the output of the Euler scheme by 1/h = 100.

**Exercise 2.** This problem asks you to implement and run the Meta-Algorithm for SEIR Calibration using the time series of cumulative number of detected infection V(t) and the cumulative number of deaths Y(t).

Use the same parameters as in Problem 1 with the following initialization: S(0) = N, E(0) = I(0) to match the data set value  $V(t_0 + \tau_0) - V(t_0 - \tau_0)$ , R(0) = 0. Use the following parameters:  $T_{max} = 119$  (i.e., a total of 120 time samples),  $V_{min} = 5$ ,  $\tau_0 = 7$ ,  $p = 1, 2, \infty$  (check all three values),  $(c_I, c_Y) = (0, 1)$  and  $(C_I, c_Y) = (1, 1)$  (check both values). For the set  $\Omega$  write nested loops over  $\alpha$ ,  $R_0$ , and  $\delta$  by searching over the following intervals:

 $\alpha \in [0.05, 0.4]$  stepsize = 0.01,  $\delta \in [0.05, 0.4]$  stepsize = 0.01,  $R_0 \in [0.8, 2.2]$  stepsize = 0.05 Recall  $\beta = R_0 \alpha$ , and N = Population is extracted from your data set.

(1) Find and print the minimum of the objective function J. Print the optimal values of  $\hat{\alpha}, \hat{\beta}, \hat{\delta}, \hat{R}_0$  and  $\hat{\gamma}, \hat{\rho}$  ( $\hat{\gamma}$  is obtained by implementing the minimization of  $||Y - \gamma R_{sim}||_p$  over  $\gamma$ , whereas  $r\hat{h}o$  is obtained by minimization of  $||I - \rho I_{sim}||_p$  over  $\rho$ ).

- (2) Visualize the 2D surfaces (functions):
  - $(\alpha, \beta) \mapsto J = J(\alpha, \beta, \hat{\delta}; \hat{\gamma}, \hat{\rho})$  at the optimal values  $\hat{\delta}, \hat{\gamma},$  and  $\hat{\rho}$  obtained before. You should obtain one plot for each p.
  - $(\alpha, \delta) \mapsto J = J(\alpha, \hat{\beta}, \delta; \hat{\gamma}, \hat{\rho})$  at the optimal values  $\hat{\beta}, \hat{\gamma}$  and  $\hat{\rho}$  obtained before. You should obtain one plot for each p.
  - $(\beta, \delta) \mapsto J = J(\hat{\alpha}, \beta, \delta; \hat{\gamma}, \hat{\rho})$  at the optimal values  $\hat{\alpha}, \hat{\gamma}$  and  $\hat{\rho}$  obtained before. You should obtain one plot for each p.
- (3) For  $(\alpha, \beta, \delta) = (\hat{\alpha}, \hat{\beta}, \hat{\delta})$ , plot on the same graph the simulated  $I_{sim}$  and the preprocessed rate of detected infections  $I(t) = V(t + t_0 + \tau_0) V(t + t_0 \tau_0)$ . Plot on the same graph the predicted number of deaths  $Y_{sim} = \hat{\gamma}R_{sim}$  and the observed number of deaths Y(t) from your data set. You should obtain two plots for each value of p.
- (4) Repeat the previous questions for each pair  $(c_I, c_Y)$ .