April 10, 2018 Heroin Model

S=susceptibles, P=prescribed opioid users,

A=addicted to opioids, H=heroin users/addicted, R=treatment/rehabilitation  $S(0) = S_0$ ,  $P(0) = P_0$ ,  $A(0) = A_0$ ,  $H(0) = H_0$ ,  $R(0) = R_0$ 

Assume  $H_0 > 0$ ,  $\mu_H > \mu_A$  and  $\theta_3 > \theta_1, \theta_2$ 

$$\begin{split} \frac{\mathrm{d}S}{\mathrm{d}t} &= -\alpha S - \beta (1-\xi)SA - \beta \xi SP - \theta_1 SH + \epsilon P + \delta R + \mu (P+R) + (\mu + \mu_A)A + (\mu + \mu_H)H \\ \frac{\mathrm{d}P}{\mathrm{d}t} &= \alpha S - \epsilon P - \gamma P - \theta_2 PH - \mu P \\ \frac{\mathrm{d}A}{\mathrm{d}t} &= \gamma P + \sigma_A R + \beta (1-\xi)SA + \beta \xi SP - \zeta A - \theta_3 AH - (\mu + \mu_A)A \\ \frac{\mathrm{d}H}{\mathrm{d}t} &= \theta_1 SH + \theta_2 PH + \theta_3 AH + \sigma_H R - \nu H - (\mu + \mu_H)H \end{split}$$

 $\frac{\mathrm{d}R}{\mathrm{d}t} = \zeta A + \nu H - \delta R - \sigma_A R - \sigma_H R - \mu R$ 

The following is a brief description of each parameter in the system:

 $\alpha$ : the rate at which people are prescribed opioids

 $\beta$ : total probability of becoming addicted to opioids other than by prescription

 $\beta(1-\xi)$ : proportion of which the non-prescribed, susceptible population becomes addicted to opioids by black market drugs and other addicts

 $\beta \xi$ : proportion of which the non-prescribed, susceptible population obtains extra prescription opioids and becomes addicted

 $\theta_1$ : rate at which the non-prescribed, susceptible population becomes addicted to heroin by black market drugs and other addicts

 $\epsilon$ : rate at which people come back to the susceptible class after being prescribed opioids (i.e. not addicted)

 $\delta$ : rate at which people come back to the susceptible class after successfully finishing treatment

 $\mu$ : natural death rate

 $\mu_A$ : enhanced death rate for opioid addicts (overdose rate which results in death)

 $\mu_H$ : enhanced death rate for heroin addicts (overdose rate which results in death)

 $\gamma$ : rate at which prescribed opioid users become addicted

 $\theta_2$ : rate at which opioid prescribed user population becomes addicted to heroin

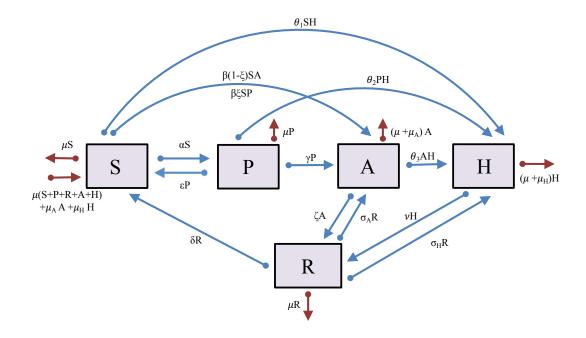
 $\sigma_A$ : rate at which people relapse from treatment into the opioid addicted class

 $\zeta$ : rate at which addicted opioid users enter treatment/rehabilitation

 $\theta_3$ : rate at which the opioid addicted population becomes addicted to heroin

 $\sigma_H$ : rate at which people relapse from treatment into the heroin addicted class

 $\nu$ : rate at which heroin users enter treatment/rehabilitation



To find the addiction-free equilibrium (AFE), we set Eqs. (FILL IN) equal to zero and require that A = H = R = 0. We are left with the system

$$0 = -\alpha S^* - \beta \xi S^* P^* + \epsilon P^* + \mu P^*$$
$$0 = \alpha S^* - \epsilon P^* - \gamma P^* - \mu P^*$$
$$0 = \gamma P^* + \beta \xi S^* P^*$$

If P=0, then the only solution is  $S^*=P^*=H^*=R^*=0$ . Thus, will assume  $P\neq 0$ . This forces  $\gamma+\beta\xi S^*=0$  and since all of our parameters and variables are non-negative, then it must be  $\gamma=0$  and either  $\beta=0$  or  $\xi=0$ . Under the assumption that  $\gamma=0=\xi$  to ensure the existence of our AFE and that 1=S+P+A+H+R, we calculate the AFE to be

$$S^* = \frac{\epsilon + \mu}{\alpha + \epsilon + \mu}$$

$$P^* = \frac{\alpha}{\alpha + \epsilon + \mu}$$

$$A^* = 0$$

$$H^* = 0$$

$$R^* = 0$$

Calculating the Basic Reproduction Number,  $R_0$ 

From this point on, we will assume  $\gamma = 0$  and  $\xi = 0$  (thus,  $\beta \neq 0$ ) in order to ensure the existence of the AFE. This results in the infected compartment Eqns. (FILL IN) reducing to:

$$\frac{dA}{dt} = \sigma_A R + \beta S A - \zeta A - \theta_3 A H - (\mu + \mu_A) A$$

$$\frac{dH}{dt} = \theta_1 S H + \theta_2 P H + \theta_3 A H + \sigma_H R - \nu H - (\mu + \mu_H) H$$

$$\frac{dR}{dt} = \zeta A + \nu H - \delta R - \sigma_A R - \sigma_H R - \mu R$$

Thus, under the assumption of A, H and R as the infected compartments and parameter restrictions stated above, the assumptions of the Next Generation Method are satisfied for matrices  $\mathscr{F}$  and  $\mathscr{V}$  shown below. Note that  $\mathscr{F}_i$  represents the rate that secondary infections enter infected compartment i and  $\mathscr{V}_i$  represents the difference between the rate of transfer out of compartment i and the rate of transfer into compartment i by means different than a secondary infection. Using this method results in the following matrices:

$$\mathscr{F} = \begin{pmatrix} 0 \\ 0 \\ \beta SA \\ \theta_1 SH + \theta_2 PH \\ 0 \end{pmatrix}$$

$$\mathscr{V} = \begin{pmatrix} \alpha S + \beta SA + \theta_1 SH - \epsilon P - \delta R - \mu(P + R + A + H) - \mu_A A - \mu_H H \\ -\alpha S + \epsilon P + \theta_2 PH + \mu P \\ -\sigma_A R + \zeta A + \theta_3 AH + (\mu + \mu_A) A \\ -\theta_3 AH - \sigma_H R + \nu H + (\mu + \mu_H) H \\ -\zeta A - \nu H + \delta R + \sigma_A R + \sigma_H R + \mu R \end{pmatrix}$$

$$F = \begin{pmatrix} \beta S^* & 0 & 0 \\ 0 & \theta_1 S^* + \theta_2 P^* & 0 \\ 0 & 0 & 0 \end{pmatrix}$$

$$V = \begin{pmatrix} \zeta + \mu + \mu_A & 0 & -\sigma_A \\ 0 & \nu + \mu + \mu_H & -\sigma_H \\ -\zeta & -\nu & \delta + \sigma_A + \sigma_H + \mu \end{pmatrix}$$

The eigenvalues of  $FV^{-1}$  are calculated to be:

$$\sigma(FV^{-1}) = \big\{0, \tfrac{(r+s) - \sqrt{(r-s)^2 + 4\beta S^* z \sigma_A \zeta \sigma_H \nu}}{2 det(V)}, \tfrac{(r+s) + \sqrt{(r-s)^2 + 4\beta S^* z \sigma_A \zeta \sigma_H \nu}}{2 det(V)}\big\}$$

 $\mathcal{R}_0$  may then be determined as the spectral radius of  $FV^{-1}$ :

$$\mathcal{R}_0 = \frac{(r+s) + \sqrt{(r-s)^2 + 4\beta S^* z \sigma_A \zeta \sigma_H \nu}}{2 det(V)}$$

where 
$$a = \zeta + \mu + \mu_A$$
,  $b = \nu + \mu + \mu_H$ ,  $c = \delta + \sigma_A + \sigma_H + \mu$ ,  $z = \theta_1 S^* + \theta_2 P^*$ ,  $r = \beta S^*(bc - \sigma_H \nu)$ ,  $s = z(ac - \sigma_A \zeta)$ , and  $det(V) = a(bc - \sigma_H \nu) - \sigma_A \zeta b$ .

We note that the radicand  $(r-s)^2 + 4\beta S^* z \sigma_A \zeta \sigma_H \nu$  is positive, since all parameters are positive. In addition, r is positive since bc contains the term that cancels with  $-\sigma_H \nu$ , s is positive since ac contains the term that cancels with  $-\sigma_A \zeta$  and finally, det(V) is positive since abc contains terms that cancel with  $-\sigma_A \zeta (\nu + \mu + \mu_H) - \sigma_H \nu$ .