to position

1. Consider again R/e << 1 so we let E R/P ~ 1- R/p only and substitute into the Regulation:

 $\frac{dR}{dt} = \mu \{ N - R - g_0 (1 - R/p) \} = \mu (1 - g_0/p) \{ \frac{N - g_0}{1 - g_0/p} - R \}$ With $R(0) = 0 , \text{ then } R(t) = \frac{N - g_0}{1 - g_0/p} (1 - e^{-M(1 - g_0/p)t})$ Thus, $\frac{dR}{dt} = \mu (N - g_0) e^{\mu (g_0/p - 1)t}$

So, for the apidemic case (Sorp), It > 0 for all to and, hence, there is no way to fit the data (figure) shown in class.

2. If you first look at the case Y=0, then we obtain Z=0 and are led to the (healthy population) equilibrium state (X,Y,Z)=(N,0,0). If $Y\neq 0$, then $X=\frac{b+v}{p}$, and working through some elementary algebra, we obtain another equilibrium state (X,Y,Z)=(b+v

state $(X,Y,Z) = (\frac{b+r}{p}) \frac{b(b+r)}{p(b+r)} \frac{b(b+r)}{p(b+r)}$ But we can only have the states exist mouningfully, i.e. have positive components, is Nois large enough. That is, we need $N > \frac{b+r}{p} = N_c$. Otherwise we only have the stable healthy rest state. (Actually, its not strickly stable since the eigenvalues are $\lambda = 0$, -b, BN-r-b, the latter being negative $(-b) = N < N_c$; but we have a zero eigenvalue.)

3. $v(+) = a + \rightarrow \frac{dx}{dt} = -a + x$, $x(0) = 1 - b \times (+) = e^{-at^2/2} \rightarrow dy(t) = a + e^{-a+^2/2}$, y(0) = 0, $y(+) = 1 - e^{-at^2/2}$.



