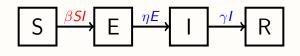
## Parameterizing Epidemic Models

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## SEIR epidemic model

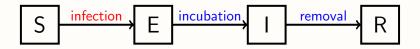


$$S' = -\beta SI$$
  
 $E' = \beta SI - \eta E$   
 $I' = \eta E - \gamma I$   
 $R' = \gamma I$ 

- ▶ Let N = S + E + I + R. Then N' = 0, so N is constant.
  - The R equation is not needed because R = N S E I.

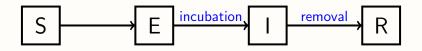
#### **Processes**

▶ Processes are either transmissions or transitions.



- Transmissions require interaction with another class.
  - Susceptibles are infected by Infectives.
- Transitions happen without any interaction.
  - Incubation of Latent individuals and removal of Infectious individuals happen spontaneously.

#### Processes – Transitions



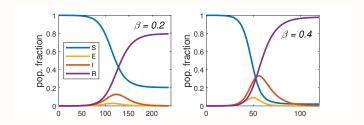
- Transition rates are (assumed to be) proportional to the leaving class
  - incubation rate =  $\eta E$
  - removal rate =  $\gamma I$
- Transition times are then exponentially distributed.
- ▶ Rate constants are reciprocals of average time in class.
  - Average removal time 10 days  $ightarrow \gamma = 0.1$

### Processes – Transmissions



- Transmission rates are proportional to the leaving class size
  - infection rate = force of infection  $*S = \lambda S$
- The force of infection is proportional to the sum of the transmitting classes (just I for SEIR)
  - force of infection =  $\beta I$
- ightharpoonup eta is the product of the encounter rate and transmission probability, neither easily measured.

### How Does $\beta$ Affect Outcomes?



- ▶ It changes the peak / value.
- ▶ It changes the final *S* value.

### SEIR Final Size Relation

 $\blacktriangleright$  We can relate S to R, independent of time.

$$\frac{dR}{dS} = \frac{R'}{S'} = \frac{\gamma I}{-\beta SI} = -\frac{1}{\mathcal{R}_0} \frac{1}{S}.$$

1. Integrate from time 0 to infinity and nondimensionalize:

• Let 
$$s = S/N$$
,  $r = R/N$ ,  $s_0 = S(0)/N$ ,  $r_0 = R(0)/N$ .  

$$\ln \frac{s_0}{s_\infty} = \mathcal{R}_0(r_\infty - r_0). \tag{1}$$

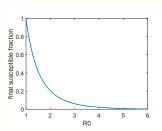
2. We can show that  $e_{\infty}=i_{\infty}=0$ , so  $s_{\infty}+r_{\infty}=1$ .

$$\ln \frac{s_0}{s_\infty} = \mathcal{R}_0(1 - r_0 - s_\infty). \tag{2}$$

#### SEIR Model - Final size relation

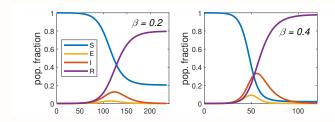
▶ Assume nearly everyone is susceptible, so  $s_0 \approx 1$  and  $r_0 = 0$ .

$$\ln \frac{1}{s_{\infty}} = \mathcal{R}_0(1 - s_{\infty}). \tag{3}$$



► This works great if  $\mathcal{R}_0 < 4$  and the epidemic ended without intervention. Then  $\beta = \gamma \mathcal{R}_0$ .

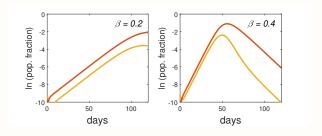
## How Does $\beta$ Affect Outcomes?



- ▶ It changes the peak / value.
- ▶ It changes the final *S* value.
- ► Maybe it does something we can't see in these graphs.

### How Else Does $\beta$ Affect Outcomes?

▶ Plot the logarithms of the infected populations *E* and *I*.



There is an extended period of exponential growth.

$$ln I = ln I_0 + \lambda t$$
,  $ln E = ln I + ln \rho$ 

for some  $I_0$ ,  $\rho$ .

$$I=I_0e^{\lambda t}, \quad E=
ho I.$$

## SEIR epidemic model – exponential phase

$$E' = \beta S_0 I - \eta E.$$
$$I' = \eta E - \gamma I.$$

With

$$E = \rho I, \quad I' = \lambda I, \quad E' = \rho \lambda I,$$

we get

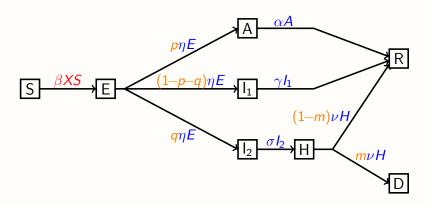
$$\beta = \frac{(\lambda + \eta)(\lambda + \gamma)}{S_0 \eta}.$$

The growth rate  $\lambda$  comes from the doubling time of class I:

$$\lambda = \frac{\ln 2}{t_d}.$$

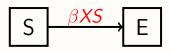
So the doubling time gives us  $\beta$ .

### SEAIHRD Epidemic Model - March 2020 Scenario



transition rates, probabilities, transmission rates

## SEAIHRD Transmission Details (March 2020)



$$X = f_c c_i I + \delta[(1 - c_i)I + f_a A]$$
 (4)

- c<sub>i</sub> is the fraction of confirmed cases for symptomatic infectives.<sup>1</sup>
- $ightharpoonup f_c$  and  $f_a$  are the infectivities of confirmed and asymptomatic cases, relative to an unconfirmed symptomatic infective.
- $\blacktriangleright$   $\delta$  is a 'contact factor' that incorporates physical distancing and mask use for unconfirmed cases.

<sup>&</sup>lt;sup>1</sup>There was no testing of asymptomatic people in spring 2020, ≥ → ⋅ ≥ → ∞ ⋅ ∞

### Transition Rate and Infectivity Parameters

- $\triangleright$  η, α, γ, σ,  $\nu$  are the reciprocals of the mean durations for classes E, A, I<sub>1</sub>, I<sub>2</sub>, and H.
- Best estimates from data for these times are
  - 5 days for incubation
  - 8 days for asymptomatic infectiousness
  - 10 days for symptomatic infectiousness
  - 6 days for transition to hospitalization
  - 8 days for hospitalization
- ▶ Best estimate for infectivity of asymptomatics is  $f_a = 0.75$ .

## Probability Parameters

- An antibody study of a random sample of people estimated about 40% of asymptomatic cases, so p = 0.4.
- ▶ About 25% of hospitalized patients died, so m = 0.25.
- ► There is no direct data for the fraction of patients who required hospitalization.
  - One study estimated that roughly 1 of 11 total cases was confirmed by testing (counting both asymptomatic and untested symptomatics), with 12% of confirmed cases leading to hospitalization.
  - From this data, we can estimate that about 1.8% of symptomatic cases required hospitalization, so q = 0.018.
- ▶ The combination q = 0.018 and m = 0.25 results in a fatality rate of just under 0.5%, which would be 1.6M Americans, assuming everyone gets infected and there is no treatment.

# $\beta$ and Initial Conditions (and $\mathcal{R}_0$ )

Assume exponential phase at start with hospitalization fraction  $H_0$ .

$$\frac{dE}{dt} = \beta S_0 X S - \eta E, \quad E(0) = e_0 H_0;$$

$$\frac{dA}{dt} = p \eta E - \alpha A, \quad A(0) = a_0 H_0;$$
(6)

$$\frac{dA}{dt} = p\eta E - \alpha A, \quad A(0) = {}_{0}H_{0}; \tag{6}$$

$$\frac{dI_1}{dt} = (1 - p - q)\eta E - \gamma I_1, \quad I_1(0) = i_{10}H_0; \tag{7}$$

$$\frac{dI_2}{dt} = q\eta E - \sigma I_2, \quad I_2(0) = i_{20}H_0; \tag{8}$$

$$\frac{dH}{dt} = \sigma I_2 - \nu H, \quad H(0) = H_0; \tag{9}$$

$$X = I + f_a A \tag{10}$$

# eta and Initial Conditions (and $\mathcal{R}_0$ )

- ▶ Unknowns  $\beta$ ,  $e_0$ ,  $a_0$ ,  $i_{10}$ ,  $i_{20}$  can be given in terms of  $\lambda$  and the other parameters.
- ▶ A history of estimates for R<sub>0</sub>:
  - December 2019: Published value of  $\mathcal{R}_0 = 2.6$  based on statistical analysis of data from China.
  - January 2020 through March 2020: Numerous published values, most between 2.5 and 3.5, but one over 5.
  - o April 8, 2020: Value  $\mathcal{R}_0 = 5.0$  published on UNL COVID modeling web page and used in teaching modules for Math 203 and elsewhere. Based on hospitalization data from New York City.
  - July 2020: 'Definitive' result  $\mathcal{R}_0 = 5.7$  published by Los Alamos research group (Sanche et al similar method to mine, but better data).

### Public Health Parameters

- ► The public health parameters are difficult to estimate and highly variable.
- ▶ The testing fraction for symptomatic patients  $c_i$  was probably in the range 0.1 to 0.6. This parameter is a good choice for a parameter study showing the impact of testing.
- ▶ The infectivity of isolation can only be guessed. I use  $f_c = 0.1$ .
- ▶ The contact factor  $\delta$  is another good choice for a parameter study. Cell phone data suggests values from 0.2 to 0.6.

### Some Conclusions

- Looking at data in different ways can yield insights that you might have missed.
- 'Nobody believes a model except the person who created it; everyone believes data except the person who collected it.'2
   We should be equally sceptical of models and data.
- ▶ It is often easier to measure a parameter by its effect than by direct measurement.
- ▶ If you get bad results from a good model, then at least one parameter value is wrong.

<sup>&</sup>lt;sup>2</sup>Paraphrase of a quotation from Albert Einstein.