Modeling Epidemics: Introduction, Simple Model, and Linear Least Squares

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First Models

- Preliminary goal: Model the spread of an contagious illness through a population.
- Simplifying assumptions:
 - The total population *N* is constant in time.
 - A newly infected person becomes infectious the next day and remains infectious forever.
 - Each infectious person is equally likely (probability *p*) to infect each noninfectious person on a given day.

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• Let $\mathcal{I}(t)$ be the number of infectious people at the start of day *t*.

Stochastic Model

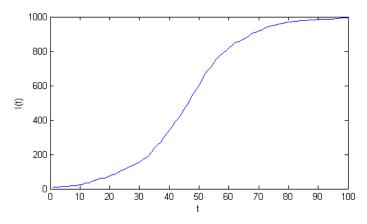
- Number the people from 1 to *N*.
- Let x_n(t) be the infectious status (1 if infectious, 0 if not) of person n at the start of day t.
- We can simulate a possible spread of the illness with the following program ("rand"= random number):

```
for t=1:T-1 for n=1:N let x(n,t+1)=x(n,t) for m=1:N if x(m,t)=1 and rand<p, then let x(n,t+1)=1 end end end
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Simulation Results

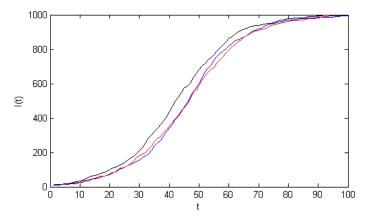
- Notice that $\mathcal{I}(t) = \sum_{n=1}^{N} x_n(t)$.
- Here are the results of a simulation with $p = 10^{-4}$, N = 1000, and $\mathcal{I}(1) = 10$:



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Simulation Results

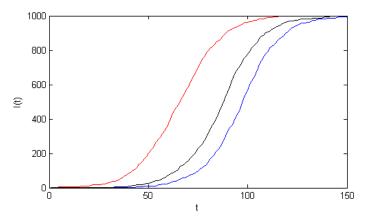
 And here are the results of three different simulations with p = 10⁻⁴, N = 1000, and I(1) = 10:



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Simulation Results

 Finally, here are the results of three different simulations with p = 10⁻⁴, N = 1000, and I(1) = 1:



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Expected (Average) Daily Outcome

- Let's determine the expected number of people infected on a day t that starts with I(t) infectious people and N – I(t) who are susceptible to infection.
- A susceptible person *n* has probability 1 − *p* of NOT being infected on day *t* by a given infectious person *m*. Therefore, person *n* has probability (1 − *p*)^{*I*(*t*)} of NOT being infected on day *t*.
- The expected number of people who are infected on day *t* is then [1 − (1 − *ρ*)^{*I*(*t*)}][*N* − *I*(*t*)], so

 $\boldsymbol{E}[\mathcal{I}(t+1)] = \mathcal{I}(t) + [1 - (1 - \boldsymbol{p})^{\mathcal{I}(t)}][\boldsymbol{N} - \mathcal{I}(t)]$

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Deterministic Models

 If both I(t) and N – I(t) are large enough, it may be reasonable to approximate I(t + 1) by its expected value, resulting in a deterministic model:

 $\mathcal{I}(t+1) = \mathcal{I}(t) + [1 - (1 - p)^{\mathcal{I}(t)}][N - \mathcal{I}(t)]$ (1)

• If $p\mathcal{I}(t)$ is small, we can approximate $(1 - p)^{\mathcal{I}(t)}$ by $1 - p\mathcal{I}(t)$, yielding a simpler model:

 $\mathcal{I}(t+1) = \mathcal{I}(t) + \rho \mathcal{I}(t)[N - \mathcal{I}(t)]$ (2)

• For these models, given $\mathcal{I}(1)$ we can compute $\mathcal{I}(2)$, $\mathcal{I}(3)$,

Deterministic versus Stochastic

- These deterministic models are much more efficient to compute (1 calculation versus N^2 for the stochastic model). Their predictions may be just as reasonable as any particular simulation of the stochastic model.
- The stochastic model can give some idea of the uncertainty of its predictions via multiple simulations; the deterministic models we've written down say nothing about their uncertainty.

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Continuous-Time Model

- The models discussed so far are called discrete-time models; time *t* takes on only integer values.
- When the quantities being modeled change slowly enough, we can approximate these models by continuous-time processes. Approximating model (2) by replacing Δ*I* = *I*(*t* + 1) *I*(*t*) by *dI*/*dt*, we get

$$d\mathcal{I}/dt = p\mathcal{I}(t)[N - \mathcal{I}(t)]. \tag{3}$$

This differential equation is commonly called the Logistic Growth Model.

• We can write down an exact solution to this differential equation:

$$\mathcal{I}(t) = \frac{N\mathcal{I}(0)}{\mathcal{I}(0) + [N - \mathcal{I}(0)]e^{-pNt}}$$

Fitting the Model to Data

- The solution I(t) of model (3) has three parameters:
 N, *p*, and I(0). Suppose we know *N* but not the other two parameters. Given a set of data points [t_j, I_j], we can ask which values of *p* and I(0) best fit the data.
- [A more fundamental (but more difficult) question is whether the model can adequately fit the data at all; are there ANY parameters of the model that fit the data reasonably well?]
- We could try to minimize the sum of the squares of the residuals I_j – I(t_j). However, this would be a nonlinear least squares problem, because I(t) does not depend linearly on p or I(0).

Method 1 to use Linear Least Squares

 If the data is given at consecutive values of *t*, say t_j = j, then one approach is to use model (2) and write

 $\mathcal{I}(t+1) - \mathcal{I}(t) = p\mathcal{I}(t)[N - \mathcal{I}(t)].$

The right-hand side is a linear function of the parameter p, and linear least squares yields the value of p that minimizes the sum of the squares of the residuals $\mathcal{I}_{j+1} - \mathcal{I}_j - p\mathcal{I}_j(N - \mathcal{I}_j)$.

• This doesn't resolve the question of which value of $\mathcal{I}(0)$ to use. If we let $t_0 = 0$ for the first data point, then we could let $\mathcal{I}(0) = \mathcal{I}_0$. However, this might not be the best choice of $\mathcal{I}(0)$ in order to make the residuals $\mathcal{I}_i - \mathcal{I}(t_i)$ small.

Method 2 to use Linear Least Squares

 Going back to the solution of model (3), we can make a transformation of variables so that the transformed solution does depend linearly on its parameters. First we divide both sides into N and simplify:

 $N/\mathcal{I}(t) = 1 + [N/\mathcal{I}(0) - 1]e^{-pNt}$

• Next subtract 1 and take the logarithm:

 $\log[N/\mathcal{I}(t) - 1] = \log[N/\mathcal{I}(0) - 1] - \rho Nt$

• Let $Z(t) = \log[N/\mathcal{I}(t) - 1]$; then the model becomes Z(t) = Z(0) - pNt. This is a linear function of the parameters pN and Z(0). One can transform the data to pairs (t_j, Z_j) , use linear least squares to determine values for pN and Z(0), and then solve for p and $\mathcal{I}(0)$.

Caveat

- Both methods of using linear least squares transform the model or its solution into a linear relationship between two quantities that can be computed from the data points (*t_j*, *I_j*); in the second method, the model predicts that *Z_j* is a linear function of *t_j*.
- Rather than simply accept the result of the least squares fit, one should graph the predicted relationship (e.g., *Z_j* versus *t_j*) and see if it actually looks linear. This gives some idea of how valid the model is.
- Regardless of how one determines values for *p* and *I*(0), one should also check directly how well the resulting *I*(*t*) fits the data.