

# *Epidemic models 1*

*Ben Bolker*

2021-01-11 ©BMB (except textbook material/other images)

*motivation*

- P & I data from Philadelphia 1918 flu:

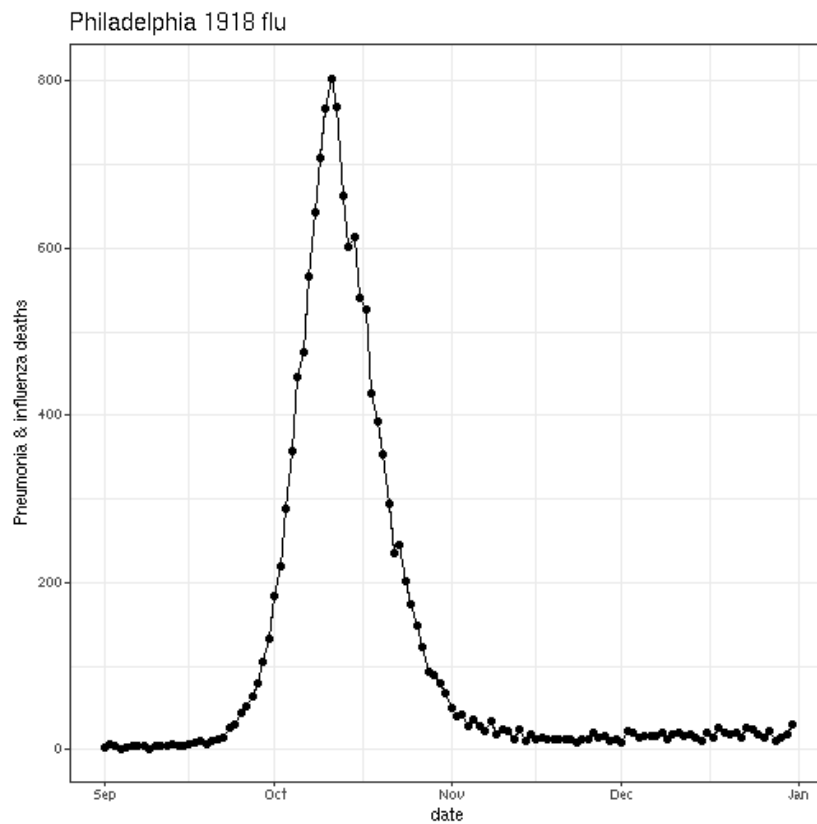


Figure 1: plot of chunk phila-data

some questions

*exponential growth*

- one variable (=1D model)
- how does disease spread? → equation

*what do we want to figure out?*

*what shall we assume?*

- classify individuals as  $S$ ,  $I$  (**compartmental** model; **microparasite** or **intensity-independent**)
- disease is transmitted from  $S$  to  $I$
- $S \rightarrow I$  instantaneously (zero latent period, no  $E$ )
- population is **homogeneous** (no heterogeneity in susceptibility, infectiousness, contact)
- fixed population size (birth = migration = 'natural' death = 0)
- transmission rate is time-invariant

- 
- assumption 2 is OK (Pasteur, [Koch's postulates](#) ...)
  - all the rest are approximations

start simple!

- parsimony
- robustness?
- applicability/estimation?

Levins (1966) (also Orzack et al. (1993), Levins (1993), Weisberg (2007))

---

*what variables should we use?*

- time ( $t$ )
- state variable: incidence, prevalence, death rate, death toll (= cumulative death?)
- deaths loosely connected to transmission

but deaths are observed!

---

when are deaths a good **proxy** for incidence?

- infection  $\rightarrow$  death time is fixed

- homogeneity? (might not matters?)
- mortality curve is shifted epidemic

(COVID context ... we observe case reports, number of tests, hospitalizations, and deaths)

- **incidence**: number of infections per unit time (rate or flow)
- **prevalence**: number of currently infected people (quantity or stock)

prevalence is closer to the **mechanism**

---

model components:

- $I(t)$  (state variable: prevalence)
- $I(0)$  (initial conditions)
- $\beta$  (parameter) = avg contacts **per susceptible per infective per unit time**

$$I(t + \Delta t) \approx I(t) + \beta I(t) \Delta t$$

Take  $\lim \Delta t \rightarrow 0$  (and solve):

$$\frac{dI}{dt} = \beta I \rightarrow I(t) = I(0) \exp(\beta t)$$

*model criticism*

- Ignored discrete nature of individuals
- Ignored time-varying  $\beta$  (e.g. **diurnal** fluctuations)
- Ignored finite infectious periods (recovery/death)

---

**Next:** What if we make infectious periods finite? (i.e., including recovery (**clearance**) or death)

$$dI/dt = \beta I - \gamma I$$

*mean infectious period*

$$I(t) = I(0) \exp(-\gamma t)$$

$$\text{proportion uninfected} = \exp(-\gamma t)$$

$$\text{proportion infected} = 1 - \exp(-\gamma t) (= \text{CDF} := C(t))$$

$$\text{PDF} := C'(t) = \gamma \exp(-\gamma t)$$

$$\text{substitute } x = \gamma t, dx = \gamma dt$$

$$\text{mean} = E[t] = \int t \exp(-\gamma t) dt = \int x \exp(-x) dx / \gamma = 1/\gamma$$

*dimensional analysis*

rates and characteristic times/scales

- is  $I$  a proportion or a density or a number ... ?
- what are the units of  $\beta$ ,  $\gamma$  ?

*nondimensionalization*

- standardize any values that can be eliminated **without loss of (mathematical) generality**
- what can we do here?
- $\gamma = 1$
- $I$  ? (depends on how we have defined it initially)  $\rightarrow$

*references*

- Levins, R. 1966.. *American Scientist* 54: 421–431. <https://www.jstor.org/stable/27836590>.
- Levins, R. 1993.. *Quarterly Review of Biology* 68 (4): 547–555.
- Orzack, SH et al. 1993.. *Quarterly Review of Biology* 68 (4): 533–546.
- Weisberg, M. 2007.. *Biology & Philosophy* 21 (5) (January): 623–645. doi:10.1007/s10539-006-9051-9. <http://www.springerlink.com/content/v141t07655057614/>.