# Measuring the strength and speed of epidemics

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### **Outline**

#### Introduction

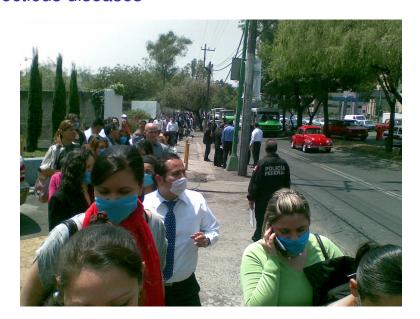
Speed of epidemics Strength of epidemics

Generation intervals through time

Estimating the effect of generation intervals Moment approximations

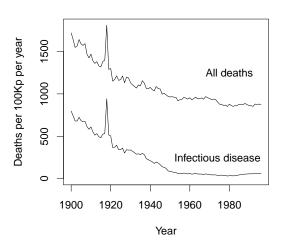
Strength and Speed of Epidemics HIV example

# Infectious diseases

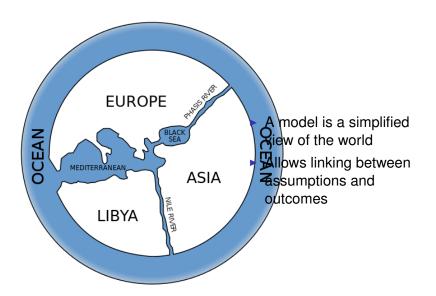




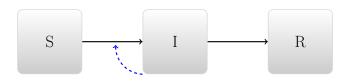
### US annual mortality rate (CDC)



### Models



# Dynamic models

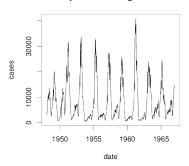


Small-scale events ⇔ Large-scale patterns and outcomes

# Dynamic models



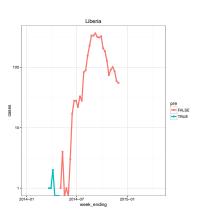
#### Measles reports from England and Wales



# Dynamic modeling

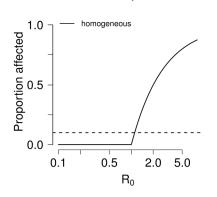
#### Connects scales





### Yellow fever in Panama

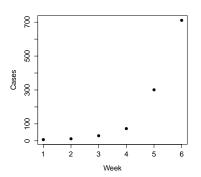
#### endemic equilibrium



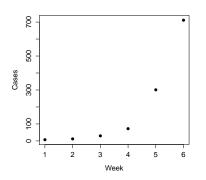


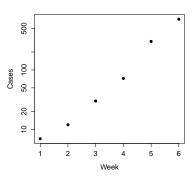
# Speed of epidemics

- Exponential growth:
  - Growth proportional to size

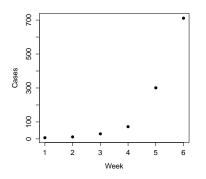


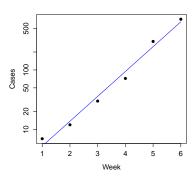
# Exponential growth





# Exponential growth



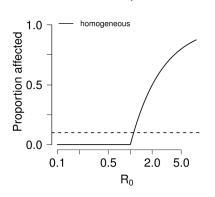


## **Speed**

- We measure epidemic speed using little r:
  - ► The ratio of the *change* in disease impact to the *amount* of disease impact
  - Units: [1/time]
  - Disease increases like e<sup>rt</sup>
- ▶ Time scale is C = 1/r
  - ▶ Ebola, C ≈ 1 month
  - ▶ HIV in SSA,  $C \approx 18$ *month*

# Strength of epidemics

#### endemic equilibrium

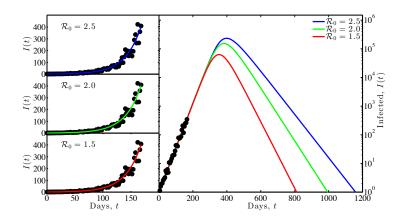




## Basic reproductive number

- We describe epidemic strength with big R
- Number of potential new cases per case
- To eliminate disease, we must:
  - ▶ Reduce transmission by a factor of R or
  - Reduce number of susceptible people by a factor of R or
- Examples:
  - ▶ Ebola,  $\mathcal{R} \approx 2$
  - ▶ HIV in SSA,  $\mathcal{R} \approx 5$

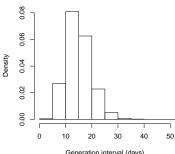
# Linking little r and big R



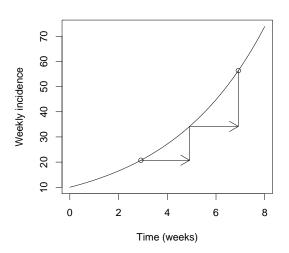
### Generation intervals

- The generation distribution measures generations of the disease
  - Interval between "index" infection and resulting infection
- What does G tell us about how dangerous the epidemic is?
  - It depends on what else we know!

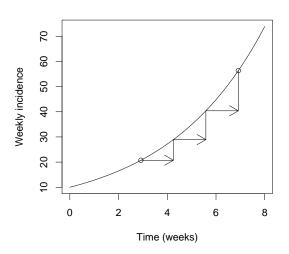
#### Approximate generation intervals



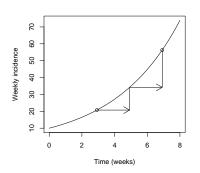
### Generations and $\mathcal{R}$

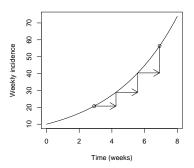


### Generations and $\mathcal{R}$



### Generations and $\mathcal{R}$





## Conditional effect of generation time

- ▶ Given the reproductive number R
  - faster generation time G means faster spread time C
  - More danger
- Given the spread time C
  - faster generation time G means  $smaller \mathcal{R}$
  - Less danger

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Speed of epidemics
Strength of epidemics

### Generation intervals through time

Estimating the effect of generation intervals Moment approximations

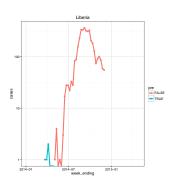
Strength and Speed of Epidemics HIV example

## Generation intervals through time

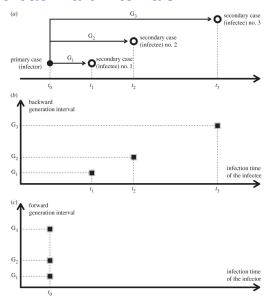
- Generation intervals can be estimated by:
  - Observing patients:
    - How long does it take to become infectious?
    - How long does it take to recover?
    - What is the time profile of infectiousness/activity?
  - Contact tracing
    - Who (probably) infected whom?
    - When did each become ill (serial interval)?

# Growing epidemics

- Generation intervals look faster at the beginning of an epidemic
  - A disproportionate number of people are infectious right now
  - They haven't finished all of their transmitting
  - We are biased towards observing faster events

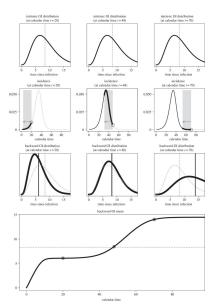


### Forward and backward intervals

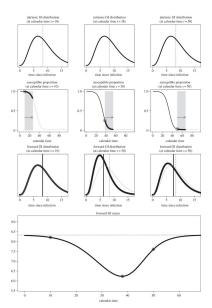


16 Champredon and Dushoff, 2015. DOI:10.1098/rspb.2015.2026

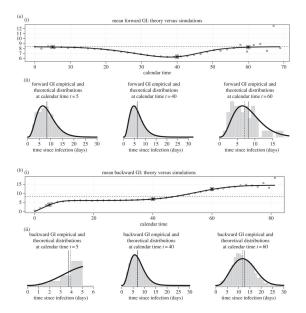
## What changes backward intervals?



## What changes forward intervals?



## Comparison to simulations



### Conclusion

- Backward intervals change if the number of infectious individuals is changing as you look back
- Forward intervals change if the number of susceptible individuals is changing as you look forward
- Lack of care in defining generation intervals can lead to bias
  - Results also tell us how to correct this bias

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Strength and Speed of Epidemics HIV example

## The "effective" generation time

▶ If the generation interval were absolutely fixed at a time interval of G, then

 $\mathcal{R} = \exp(G/C)$ 

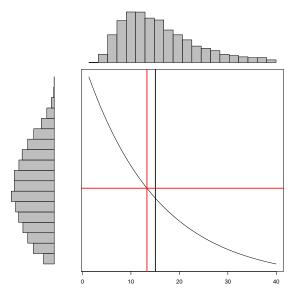
Define the effective generation time so that this remains true:

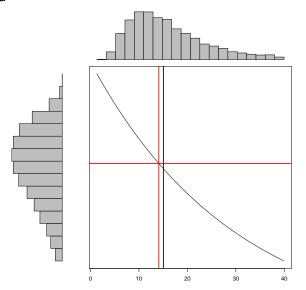
 $\mathcal{R} = \mathsf{exp}(\hat{ ilde{G}}/ ilde{C})$ 

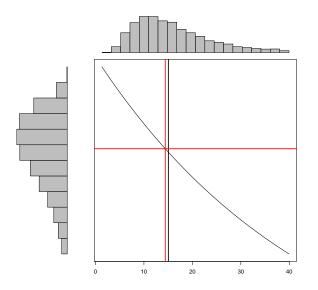
▶ We can show  $\hat{G}$  is a "filtered mean" of the distribution g:

 $\exp(-\hat{ extit{G}}/ extit{ extit{C}}) = \langle \exp(- au/ extit{ extit{C}}) 
angle_g.$ 

C = 15d





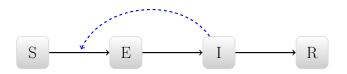


## Example: Post-death transmission and safe burial

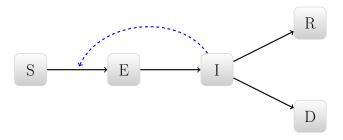
- How much Ebola spread occurs before vs. after death
- Highly context dependent
  - Funeral practices, disease knowledge
- Weitz and Dushoff Scientific Reports 5:8751.



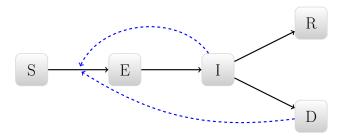
### Standard disease model



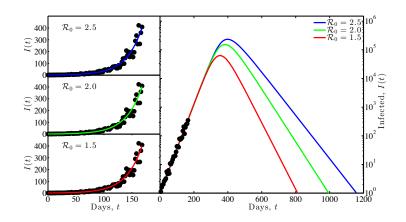
## Disease model including post-death transmission



## Disease model including post-death transmission



### **Scenarios**

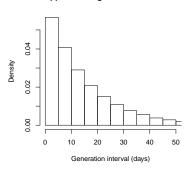


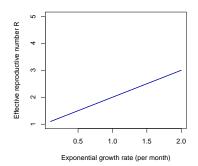
#### Conclusions

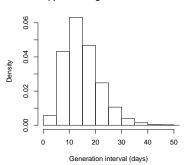
- Different parameters can produce indistinguishable early dynamics
- More after-death transmission implies
  - ▶ Higher R<sub>0</sub>
  - Larger epidemics
  - Larger importance of safe burials

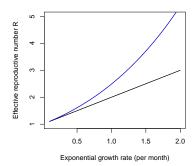
## An approximation

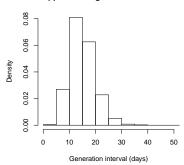
- We connect these quantities with a moment approximation
- ▶ Define  $\kappa = \sigma_G^2/\mu_G^2$  the squared coefficient of variation of the generation distribution
- $\mathcal{R} \approx (1 + \kappa \bar{G}/C)^{1/\kappa}$ 
  - Equal when G has a gamma distribution
  - Simple and straightforward
  - When is it a useful approximation?

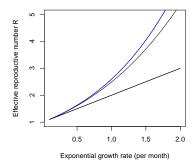


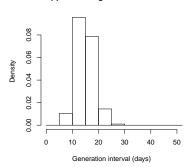


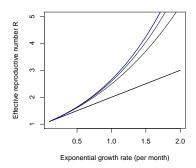








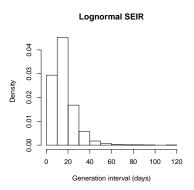




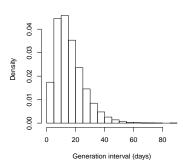
### Fitting to Ebola

- Simulate generation intervals based on data and approach from WHO report
- Use both lognormals and gammas
  - WHO used gammas
  - Lognormals should be more challenging

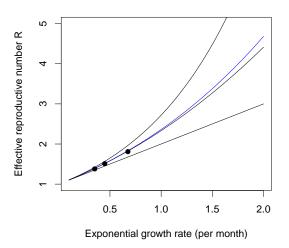
## Approximating the distribution



#### Single-gamma approximation



## Approximating the curve



## Summary

- ▶ Generation intervals are the missing link between r and R
- We need better methods for estimating them, and propagating uncertainty to other parts of the model
- For many practical applications:
  - Knowing the mean generation interval is not enough
  - But knowing the mean and CV may be enough

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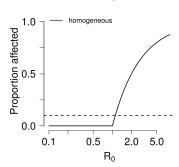
Estimating the effect of generation intervals Moment approximations

Strength and Speed of Epidemics HIV example

## Strength: R – the reproductive number

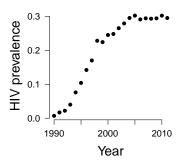
- Expected number of new cases per cases
- $\triangleright \mathcal{R} = \beta DS/N$ 
  - Disease increases iff R > 1

#### endemic equilibrium



## Speed: r – the growth rate

- $\rightarrow$   $i(t) \approx i(0) \exp(rt)$
- ightharpoonup C = 1/r
- ►  $T_2 = \ln(2)/r$

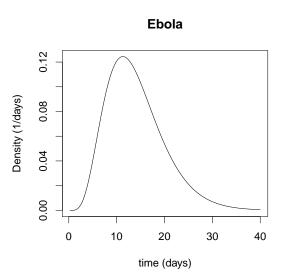


## $\ensuremath{\mathcal{R}}$ and the generation interval

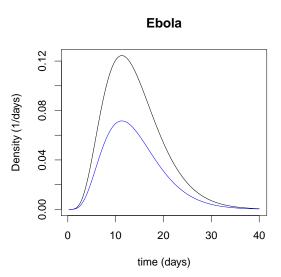
$$i(t) = \int k(\tau)i(t-\tau)\,d\tau$$

- $ightharpoonup \mathcal{R} = \int k(\tau) d\tau$
- ▶ Define the intrinsic generation interval distribution:  $k(\tau) = \mathcal{R}g(\tau)$

## ${\cal R}$ and the generation interval



## ${\cal R}$ and the generation interval

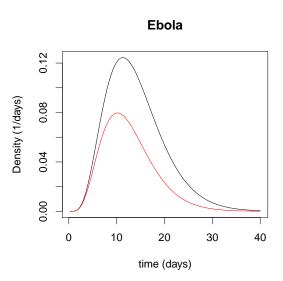


## r and the (other) generation interval

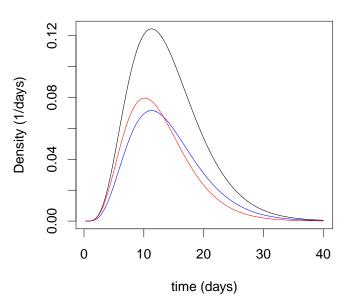
$$i(t) = \int k(\tau)i(t-\tau)\,d\tau$$

- if i(t) grows like  $\exp(rt)$ , then
- ▶ 1 =  $\int k(\tau) \exp(-r\tau) d\tau$
- ▶  $b_0(\tau) = k(\tau) \exp(-r\tau)$  is the initial *backwards* generation interval

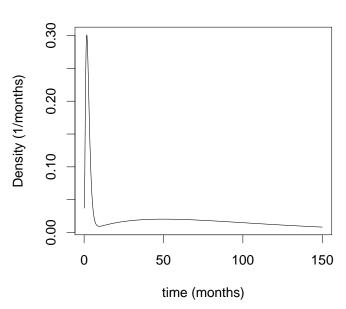
## r and the (other) generation interval



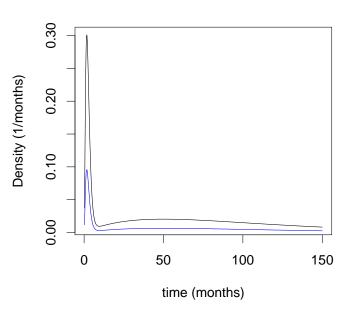
#### **Ebola**



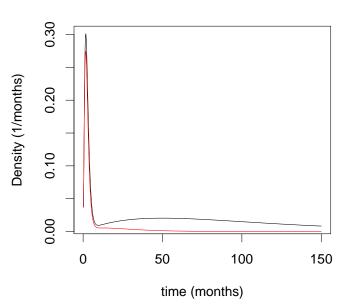
### HIV



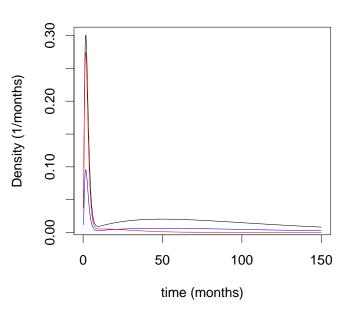
### HIV







### HIV



## Strength of intervention

- Imagine we have an intervention that reduces transmission
  - $k(\tau) \rightarrow k(\tau)/L(\tau)$
  - ▶ Define *strength*  $\theta = \mathcal{R}/\hat{\mathcal{R}}$  the proportional amount by which the intervention reduces transmission.
- We then have:
  - $\bullet \ \theta = 1/\langle 1/L(\tau)\rangle_{g(\tau)}$
  - θ is the harmonic mean of L, weighted by the generation distribution g.
- ▶ Outbreak can be controlled if  $\theta > \mathcal{R}$

## Speed of intervention

▶ Define the *speed* of an intervention be  $\phi = r - \hat{r}$  – the amount by which the intervention slows down spread.

$$1 = \left\langle \frac{\exp(\phi \tau)}{L(\tau)} \right\rangle_{b(\tau)}$$

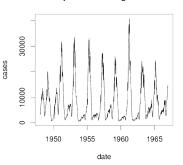
- \$\phi\$ is sort of a mean of the hazard associated with L
  - Averaged over the initial backwards generation interval
- ▶ Outbreak can be controlled if  $\phi > r$ .

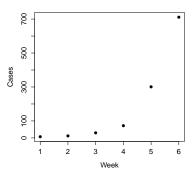
# A new way of looking



## Measuring the epidemic

#### Measles reports from England and Wales





# Measuring the intervention



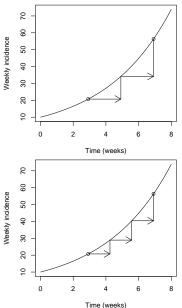


## Can treatment stop the HIV epidemic?



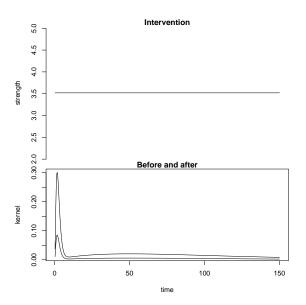
## Are HIV generations fast or slow?

- Fast generations mean:
  - Testing and treating will help less
  - but lower epidemic strength

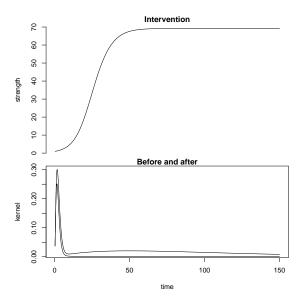




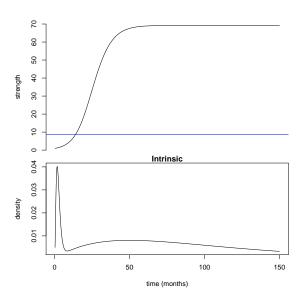
#### Condom intervention



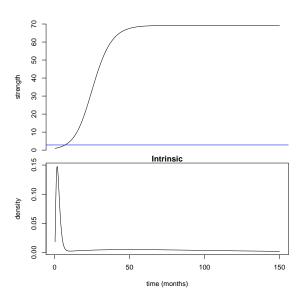
#### Find and treat



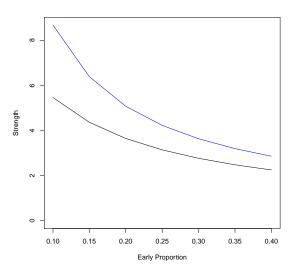
## Low early transmission



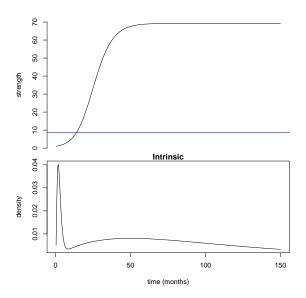
## High early transmission



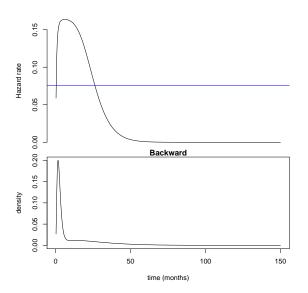
## Range of estimates



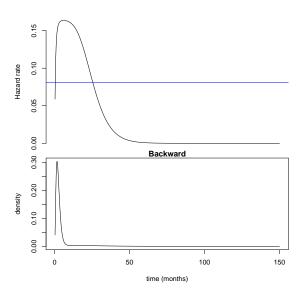
#### Find and treat



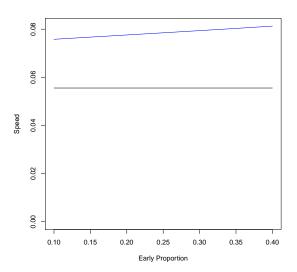
## Low early transmission



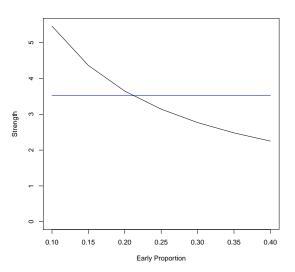
## High early transmission



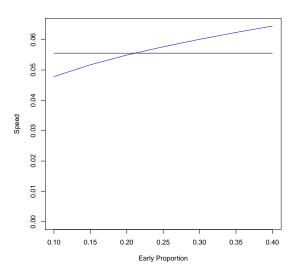
## Range of estimates



### Condom intervention



### Condom intervention



#### Conclusion

- This perspective helps us understand why find and treat predictions are robust to assumptions about transmission
- Sometimes "strength" and sometimes "speed" can help us see epidemic control questions more clearly

#### **Thanks**

- Organizers
- Audience
- Collaborators
- ► Funders: NSERC, CIHR