### Serosurveillance

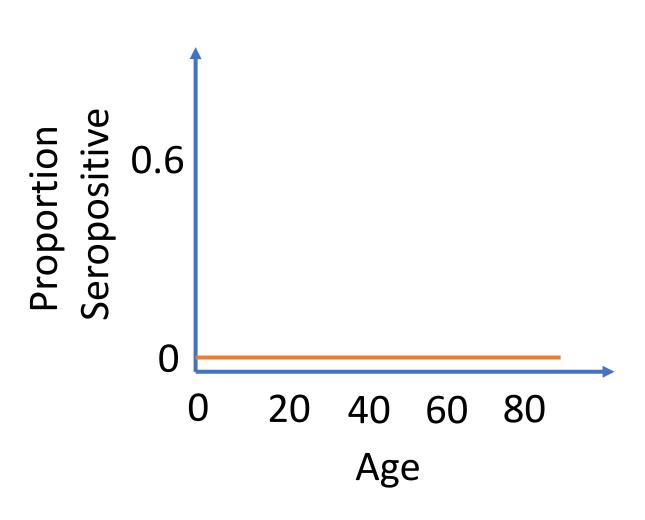
Henrik Salje
University of Cambridge
20 June 2022

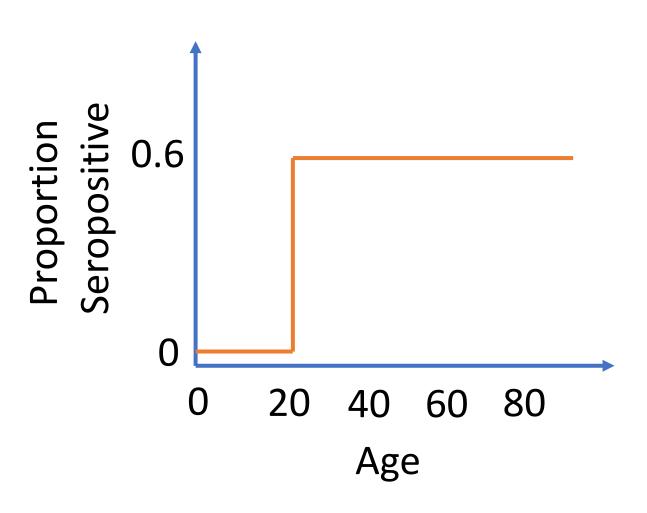
### What is serosurveillance?

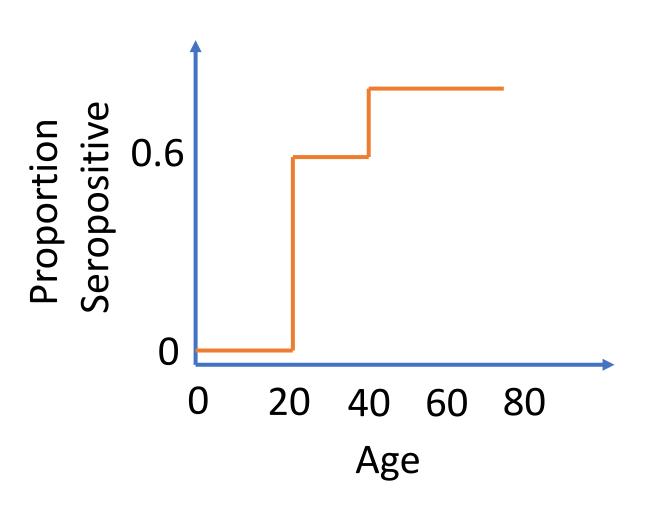
# Why is it useful as compared to case-based surveillance?

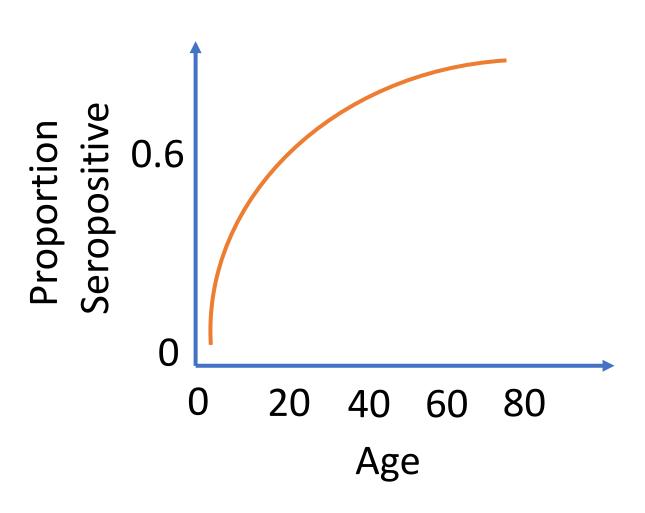
### Potential sources of sera

### Different assays

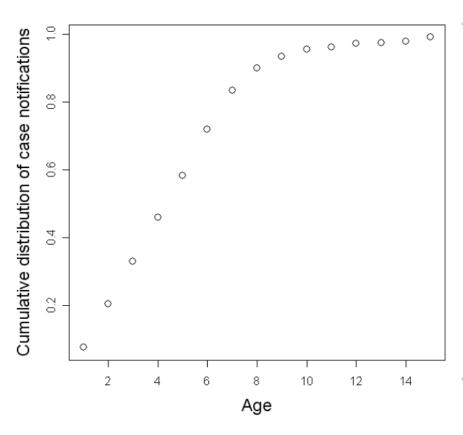








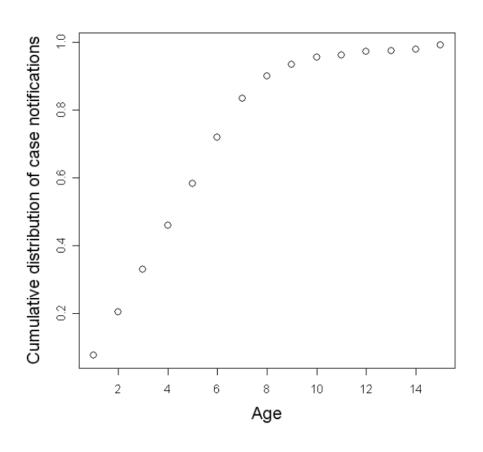
### Age-specific Seroprevalence



- Seroprevalence of measles in Aberdeen, UK
  - age-specific seroprevalence
  - this can be used to learn additional details about the disease dynamics
- What is the force of infection for this group?

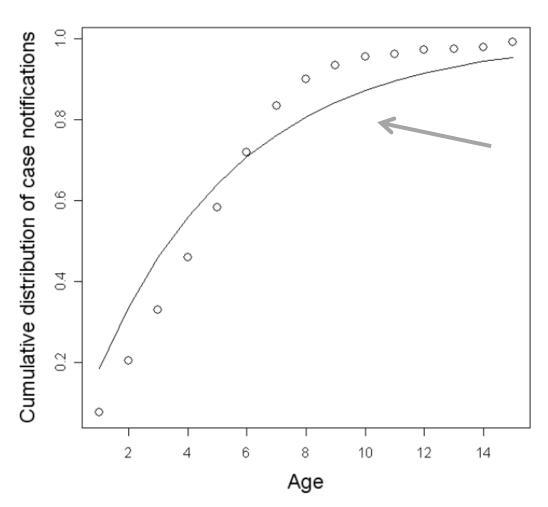
- Force of infection
  - FOI
  - Hazard of infection
  - the per capita rate at which people acquire infection
  - usually symbolized with  $\lambda$
  - in an SEIR model, equivalent to βI

### FOI for Measles in Aberdeen



What is the force of infection for this group?

### FOI for Measles in Aberdeen

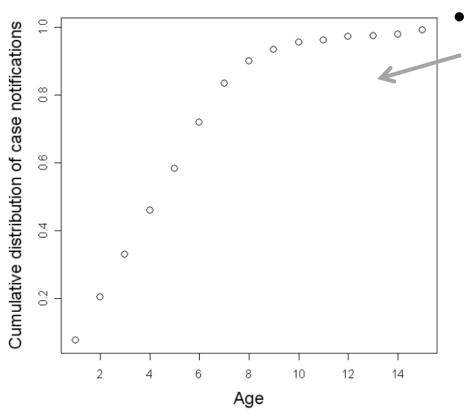


- What is the force of infection for this group?
- If we fit a constant force of infection, we see the fit is not very good
  - $\lambda = 0.21$
  - 21% chance of becoming infected each year

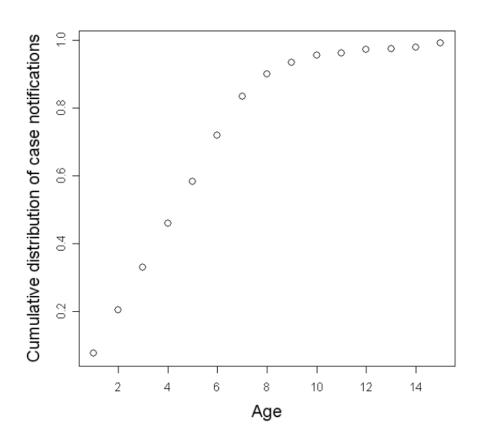
- Force of infection
  - the per capita rate at which people acquire infection
- FOI unlikely to be constant with age
  - infection risk changes with age
  - school entry, sexual activity, specific job exposures
- FOI likely to change over time
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- These factors indicate that constant FOI is a poor assumption
- Age-specific FOI would be much better and agespecific seroprevalence can help estimate this!



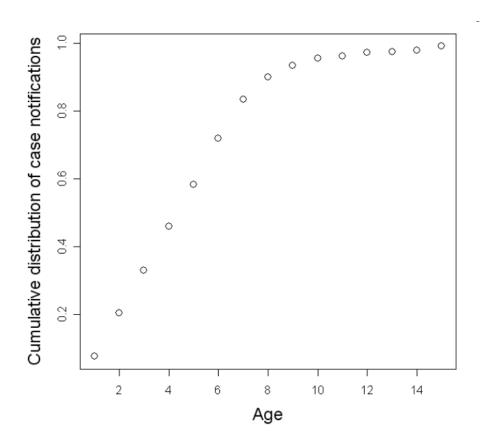
- As age increases, the proportion who are seronegative decreases
  - we expect more people to become seropositive over time
  - proportion of individuals who are seronegative at a given age (x(a)) is related to force of infection λ



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- proportion of individuals who are seronegative at a given age (x(a)) is related to force of infection λ
  - $x(a) = e^{-\lambda a}$
- proportion of individuals who are seropositive at a given age (y(a)) can then be calculated

• 
$$y(a) = 1 - x(a)$$

- Cumulative incidence by age (F(a)) is equivalent to seroprevalence
  - assumes permanent immunization
  - $F(a) = 1 e^{-\lambda a} \approx y(a)$

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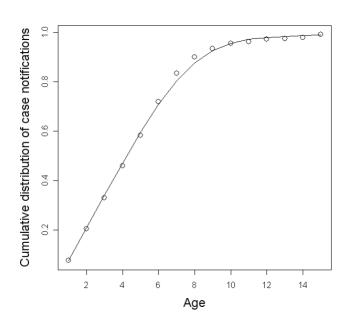
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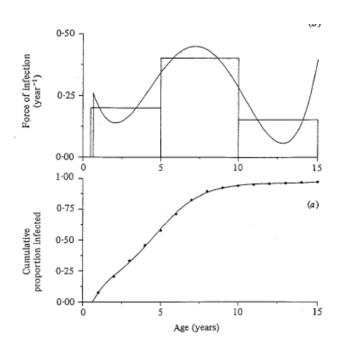
- We can use a binomial (statistical) framework to relate the seroprevalence to force of infection
- we are estimating a joint likelihood
- by using this statistical method, we can:
  - account for the different sample sizes among age groups
  - calculate confidence intervals for our estimate of FOI

### FOI for Measles in Aberdeen



- What is the force of infection for this group?
- If we fit age-specific force of infection, we see the fit is excellent
  - three separate FOIs for three age groups

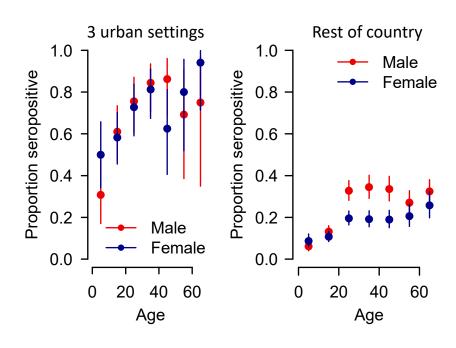
### FOI for Measles in Aberdeen



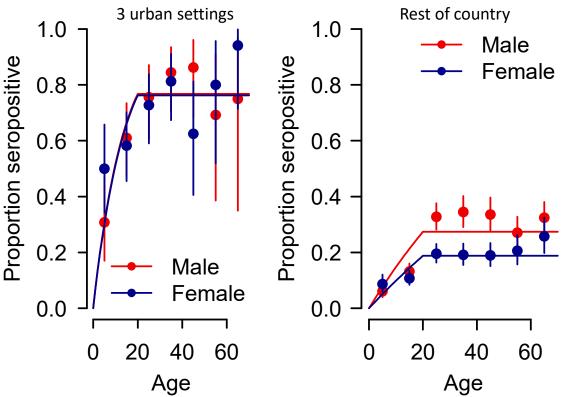
- What is the force of infection for this group?
- If we fit age-specific force of infection, we see the fit is excellent
  - three FOIs for three age groups
  - pre-school (<5 years old): 0.18
  - early school (5-10 years old): >0.3
  - late school (10-15 years old): <0.18</li>

#### Serosurveillance for dengue virus in Bangladesh



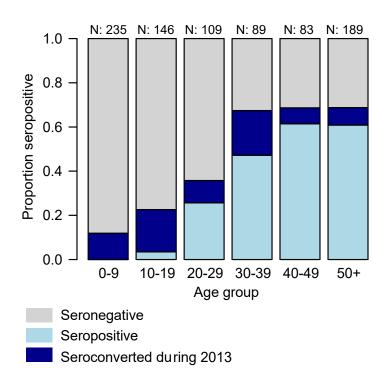


### What did we assume here to fit these FOIs for DENV in Bangladesh?



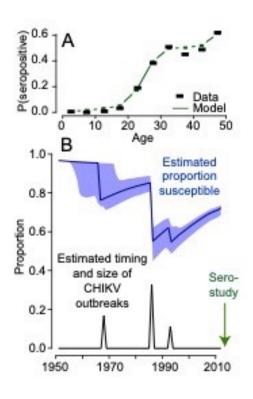
- Using serocatalytic models, we estimated that 40 million [34.3–47.2] people have been infected nationally (24% population), with 2.4 million ([1.3–4.5]) annual infections.
- 0.6% of participants reported having had dengue, of which half didn't actually have antibodies

# We can also identify more complex past infection histories — example with chikungunya



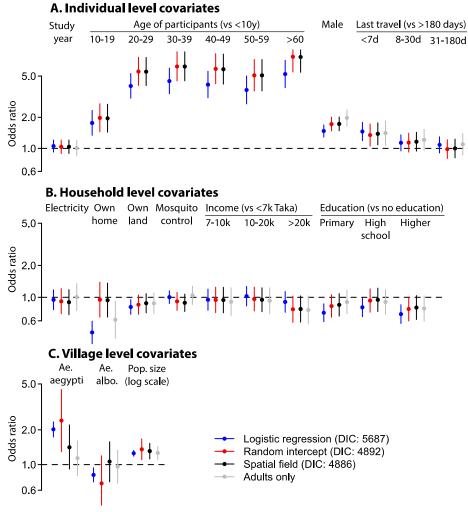
- In 2012 AFRIMS set up a cohort for chikungunya in Cebu, Philippines.
- No case of chikungunya had ever been reported in Cebu

# We can also identify more complex past infection histories

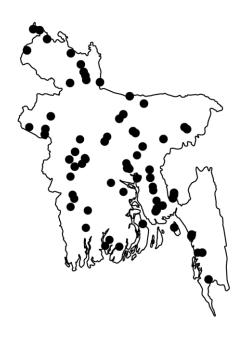


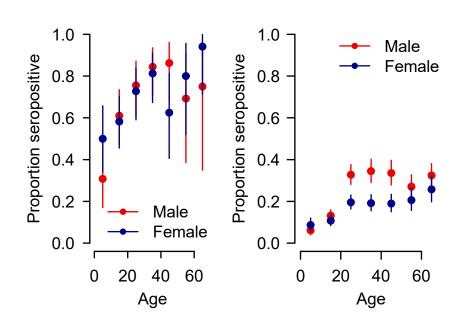
- This single seroprevalence study could identify that there had been three outbreaks in Cebu
- Using information on how the demography has changed in Cebu we could identify how many people had ever been infected (~350,000) and the changing level of immunity in the population.

We can identify risk factors for infection

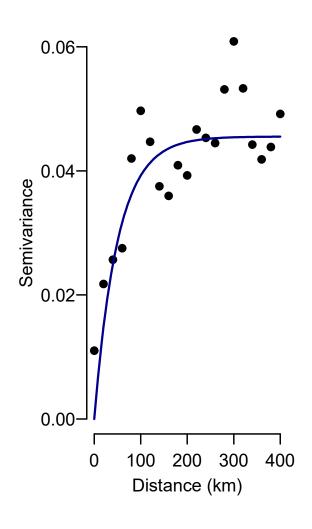


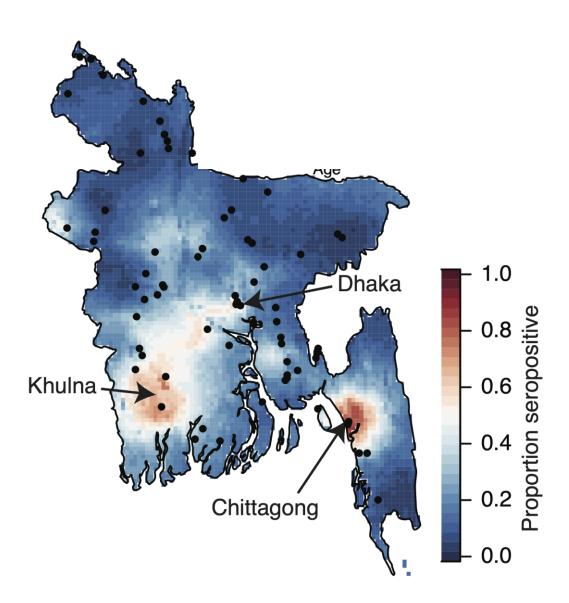
# What about places we didn't sample?



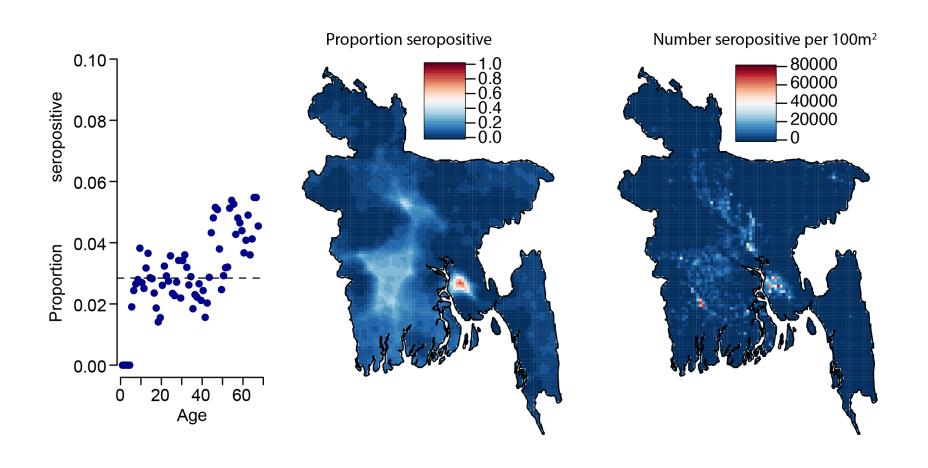


### We can use spatial correlation

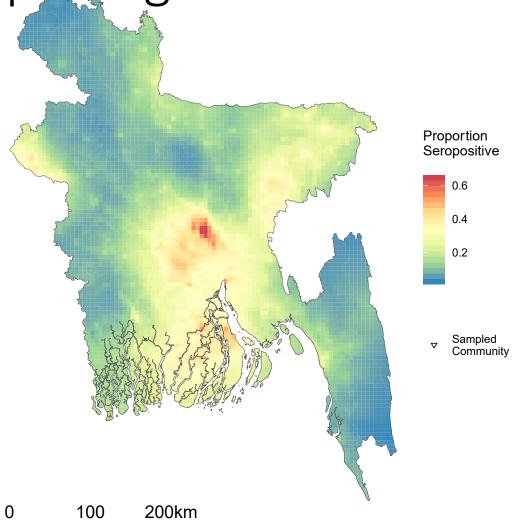




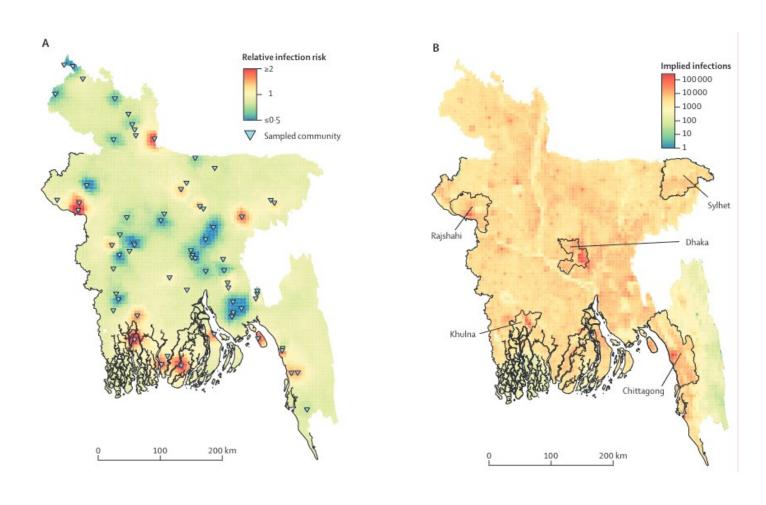
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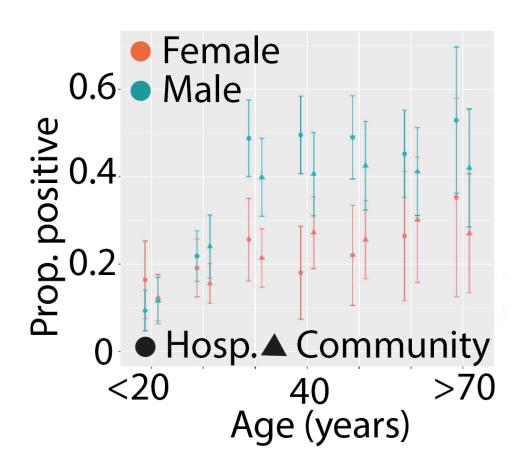


### What pathogen is this?



# Alternatives to community-based studies

# What about using sera anyway collected in hospitals?



### Conclusions

- Serosurveillance is an excellent way of understanding underlying level of infection
- Mathematical models can help maximise inferences, including the development of risk maps
- Need careful consideration of:
  - How antibodies behave post infection
  - Cross-reactivity
  - Biases in source of samples