

Modern Concepts in Incidence Estimation

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The Problem of Estimating 'Rates'

- The “gold standard” for incidence measurement was long said to be a **prospective cohort follow-up** (a.k.a. “direct observation”).
- It's not all gold:
 - **Expensive** and logistically **complex** to run
 - **Time** consuming
 - **Biased** due to selection, intervention and follow-up
- Various approaches:
 - **Direct observation** such as in ongoing cohort studies
 - **Mathematical modeling** to interpret trends in **prevalence** data
 - **Cross-sectional surveys** using ‘recent infection tests’

Outline of the Talk

- 1 Introduction
- 2 Incidence Surveillance Basics
- 3 Recent Infection

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Longitudinal Study Design ('Real' Cohorts)

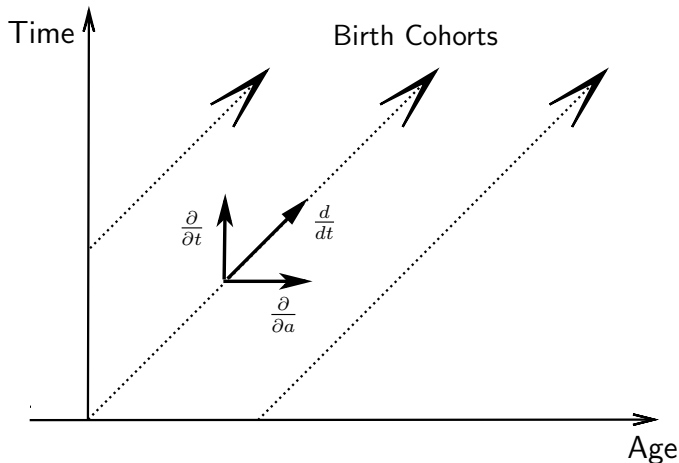
- Follow up (for 8 months) a cohort of E (1500) exposed individuals.
- Record N_{IE} (25) infection events.
- Estimate an exposure time T - a little (about 8) less than 1000 person years

$$I_{\text{est}} = \frac{N_{IE}}{T} = \frac{25}{992} \approx \frac{25}{1000} = 2.5\% \text{ p.a.}$$

'Synthetic' Cohort

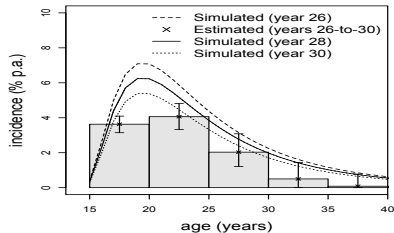
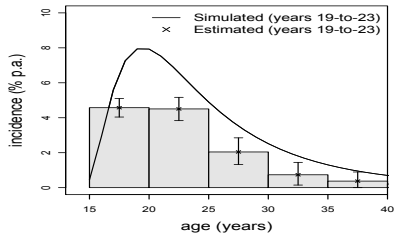
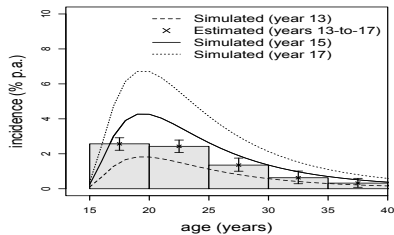
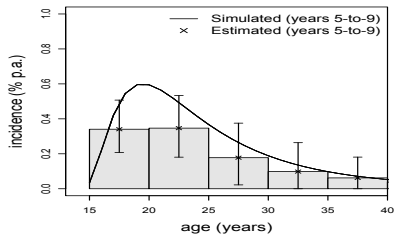
- Given **age** (and preferably also **time**) structured prevalence data
 - Need to get **external excess mortality** estimate
 - Combine with prevalence *trend* estimate
 - Obtain Incidence estimate
 - (Brunet and Struchiner, Williams et al, Hallett et al, Brookmeyer and Konikoff)
 - **Mahiane et al Plos ONE 2012**
- Note: **Considerable Complexity** in optimization, power/uncertainty analysis

Birth Cohorts

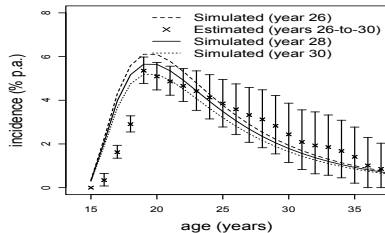
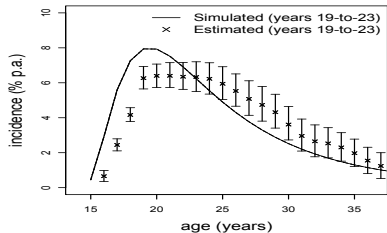
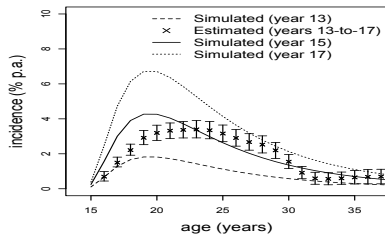
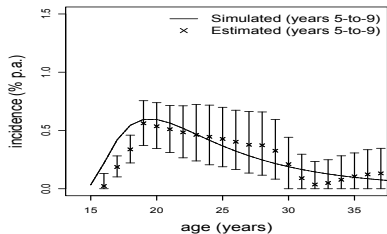


$$\frac{d}{dt} (\text{Birth Cohort}) = \left(\frac{\partial}{\partial a} + \frac{\partial}{\partial t} \right) (\text{Population Density})$$

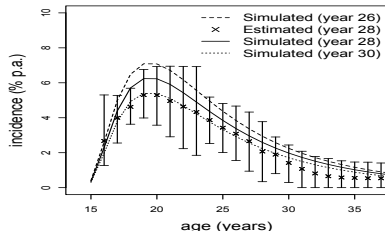
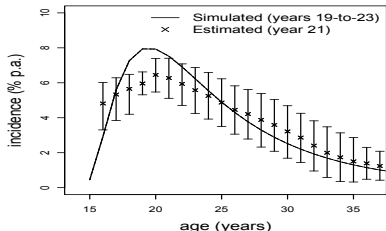
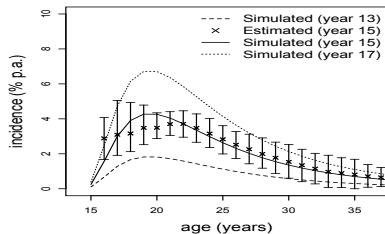
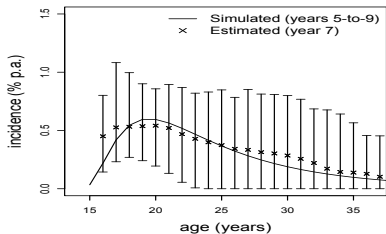
5 year Bins



Age-Smoothing Prevalence, but averaging over time



Fully Instantaneous Formulation



'Fitting' Models

- Given a general mix of country/region level data
 - Antenatal Clinic Data
 - Household Survey
 - AIDS Indicator Survey
 - Demographic and Health Survey
 - Burden of disease estimates
 - Life Tables
- Fit your best/favourite model and read off the fitted incidence
- Note:
 - Parametric assumptions unavoidable
 - Difficulty in weighting diverse sources of data
 - No formal uncertainty estimation
 - No validation

Instantaneous Population State

- Let persistence in the infected state have a mean duration D .
- For example

$$\dot{N}_I = IE - \frac{1}{D}N_I$$

- At **equilibrium**, the number of **currently** infected individuals is

$$N_I = EID$$

- Rearrange:

$$I_{\text{est}} \approx \frac{N_I}{ED}$$

- Note: **Same basic structure** as longitudinal estimate!

Cross Sectional Study Design

- Let mean persistence in the infected state $D = 1$ week (≈ 0.02 years).
- 'Survey' 100 000 civil servants
- 4000 have flu on a given date

$$I_{\text{est}} \approx \frac{4000}{96,000 \times 0.02} = 208\% \text{ p.a.}$$

- (which may mean a 10% Risk of infection in a month long flu outbreak)

When Survival in the Infected State is Long.

- If D is long \implies an uninformative, widely smoothed, 'weighting'.
- Survival post infection is not well known, and it is changing (ARV's).
- So - can we use a transient Recent Infection state?

Cross Sectional HIV estimate

- Survey 3,000 Individuals
- Prevalence 10%
- Mean duration of recent infection of test is half a year
- 15 'recent infections' detected

$$I_{\text{est}} = 1.1\% \text{ p.a.} \quad (0.6 - 1.7)$$

Outline

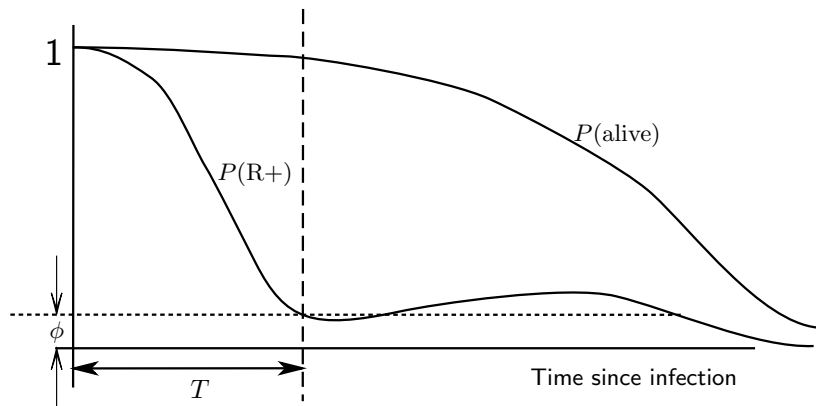
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Can Recent Infection be Usefully Defined?

- Need a **biomarker** which **correlates well** to a **meaningful** time scale
- **Many** biomarkers evolve post infection to provide **heuristic** correlates of 'recent infection'
- Need to understand what makes a biomarker useful for surveillance

Summarising Recency Test Performance into **Two** parameters

(MDRI Ω and False Recent Rate ϵ)



Talking Points

- Choosing T requires some care but is not a conventional 'optimization'
- (but choosing a threshold is a conventional, **context specific**, optimization)
- **MDRI**, importantly, contains (or should contain) mainly **biology**
- **FRR** will vary, and will inevitably be imperfectly known
- FRR should be small enough to:
 - Not imply a huge effect
 - Be acceptably approximated with low precision