

# Defining and Measuring 'Recent infection'

## Application to Incidence Estimation

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# What is 'Recent Infection'?

- Introduction of a time scale into a biological state definition poses some fundamental quandaries:
  - Is recency of fixed duration?
  - Is it linked to the presence of a biomarker?
  - Is there a 'gold standard' for evaluating a test?
- Why would we bother with knowing about recent infection?
  - **Incidence is related to 'prevalence' of recent infection**
  - Clinical/sociological response to diagnosis

# Crucial Epidemiological Indicators

- Reliable estimation of **prevalence** and **incidence** (even more so) are central to the determination of epidemiological trends.
  - **Prevalence** is the number of infected individuals.
  - **Incidence** is the 'rate' of occurrence of new infections.
- These estimates are important for:
  - Assessing **outbreaks**
  - Allocating **resources**
  - Evaluating **interventions**
  - **Planning** cohort and community based studies

# 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

# (Almost Consistent) Colour Coding

- **Terms on first use / at time of definition**
- **Jargon which is hopefully clear**
- **General Emphasis**

# Outline

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# Longitudinal Study Design

- 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

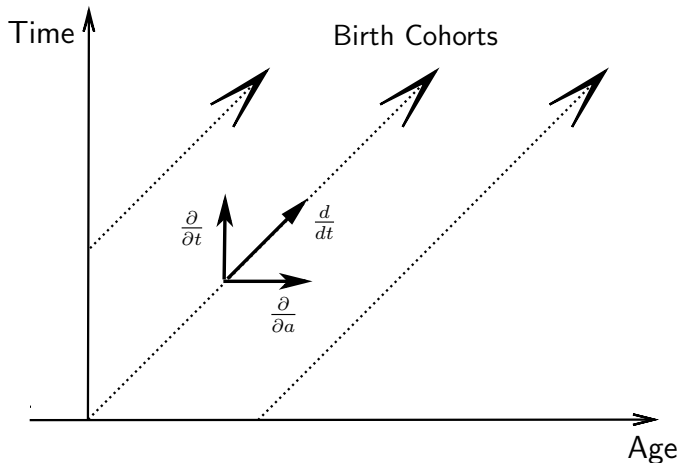
$$I_{\text{est}} = \frac{N_{IE}}{T} = \frac{25}{992} \approx \frac{25}{1000} = 2.5\% \text{ p.a.}$$



# '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



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

# '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 ( $\approx 0.02$  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

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

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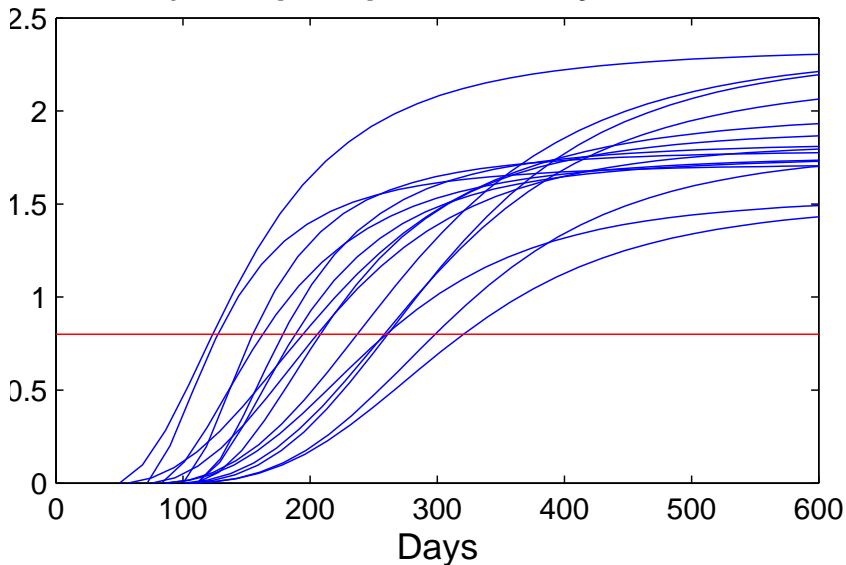
# Can Recent Infection be Defined by a Fixed Time?

- In principle, Yes - **BUT**
- As a purely time based definition, it requires an omniscient investigator
- No conceivable study design uses this level of information
- If you had a perfectly discriminating biomarker - how would you know?
  - (Think seroconverter cohort sampling intervals)

# 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

## A) Multiple Optical Density Curves



# Talking Points

- There is **always inter-subject variability**
- There is always a dynamic range of some kind - and hence the need for a **threshold**
- 'Raising the threshold' eventually, **always breaks** the concept of 'recent':
- Just **measurement noise** alone means there is always some FRR
- Choosing assay conditions and **threshold is everything**

# What Matters for Incidence Surveillance?

- Only two things **really** matter:
  - Minimization of bias (maximization of **accuracy**)
  - Minimisation of variance / Coefficient of Variation // (maximization of **precision**)
- In cross sectional surveillance of biomarkers, **accuracy** is linked to:
  - Good knowledge of the test properties
  - Unbiased sample frame
  - Use of correct analytical/statistical methodology

i.e. **NOT** the actual test properties themselves
- **Precision**, in turn is crucially linked to:
  - Long **M**ean **D**uration of **R**ecent **I**nfection
  - Low **F**alse **R**ecent **R**ate

**i.e. the actual test properties themselves!**

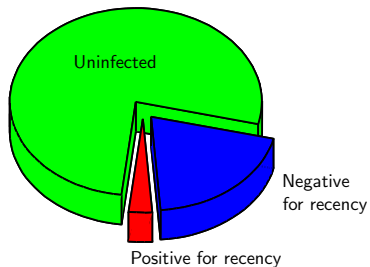
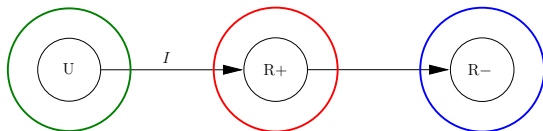
# Recent Infection Defined **Directly** by Biomarkers

- Now there is, *a priori*, no fixed time in the definition
- Lack of a diagnostic gold standard - **the biomarker is everything!**
- *Something like* Survival Analysis gives **mean recency** duration
- Recover the naive estimator
- Explicit formula for implicit temporal **averaging/weighting**

$$\hat{I} = \frac{\int IW}{\int W}$$

- (No need for **equilibrium** assumptions!)

## Simple Estimator based on knowledge of Idealised Recency

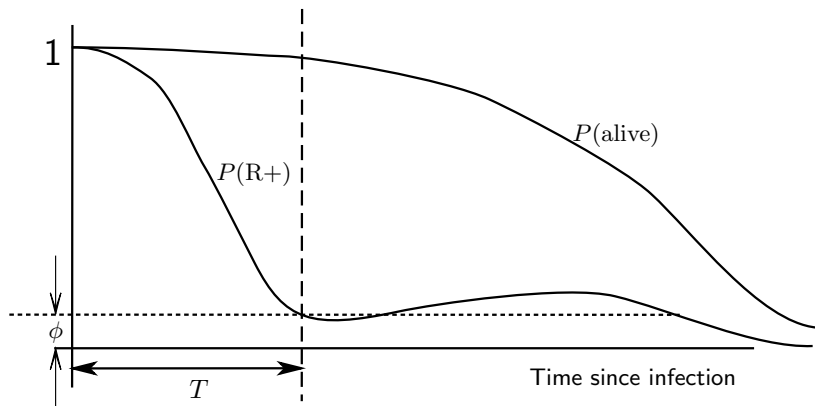


Simple **incidence estimator**:

$$I_{\text{est}} = \frac{N_{R+}}{N_U \tau_R}$$

# Summarising Recency Test Performance into **Two** parameters

(MDRI  $\Omega$  and False Recent Rate  $\epsilon$ )





# Talking Points

- There is some persistent disagreement about use of T
- (but actually a storm in a T cup)
- 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

# Estimating a Mean Duration of Recent Infection

- Recruit newly infected individuals
- Follow them up frequently
- Apply full recent infection test repeatedly
- Regression of various kinds can estimate mean recency
- Once data is reasonable - rough estimates are easy
- Uncertain infection/seroconversion time is a computational bottleneck

# Estimating a False Recent Rate

- Recruit non-newly infected individuals
- If possible, stage by time of infection
- Apply full recent infection test (including VL, etc)
- Estimate FRR 'globally' and by 'risk factors'
- Provide context for 'risk factor' *weighting* (tricky)
- Is there prior information/FRR estimate or this is all we have?

# Evaluating Recent Infection Biomarkers - I

- Context matters (stage of epidemic)
- Is new candidate biomarker supplemental or stand-alone?
- How ambitious is the funding programme?
- Can devise a variety of estimates and hypothesis tests on performance

# Evaluating Recent Infection Biomarkers - II

- Does MDRI Confidence Interval **include/exclude** critical values?
- Does FRR Confidence Interval include/exclude critical values?
- Is MDRI/FRR **superior/non-inferior** to other products (accounting for uncertainty in those)
- Given context, is CoV of Incidence Estimate, superior, non inferior to **other products**, or **targets**
- How **orthogonal** are biomarkers