

Chapter 13: More estimators for R_0

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Overview

Estimators based on ...

- ... final size data.
- ... age-structured data.
- ... a transmission experiment.
- ... intrinsic growth rate.

Estimators based on age-structured data.

- + age-dependent estimation of force of infection and R_0
- indeterminacy of quantities, such as contact frequency matrix between age classes

Next-generation operator

$$(K\phi)(a) = \int_0^{\infty} k(a, \alpha) \phi(\alpha) d\alpha \quad (1)$$

$$k(a, \alpha) = \int_0^{\infty} h(\tau, \alpha) c(a, \alpha + \tau) N(a) \frac{\mathcal{F}_d(\alpha + \tau)}{\mathcal{F}_d(\alpha)} d\tau \quad (2)$$

- ▶ $h(\tau, \alpha)$: measure of infectiousness
- ▶ $c(a, \alpha + \tau)$: quantification of age-dependent contact structure
- ▶ $N(a)$: steady-state population size
- ▶ $\frac{\mathcal{F}_d(\alpha + \tau)}{\mathcal{F}_d(\alpha)}$: ratio of survival functions

Short-disease approximation

- ▶ combination of assumptions
- ▶ length of infectious period \ll average individual lifespan

$$\frac{\mathcal{F}_d(\alpha + \tau)}{\mathcal{F}_d(\alpha)} \approx 1$$

$$c(a, \alpha + \tau) \approx c(a, \alpha)$$

$$H(\alpha) = \int h(\tau, \alpha) d\tau$$

Age-specific force of infection

$$\Lambda(a) = \int_0^\infty \int_0^\infty h(\tau, \alpha) c(a, \alpha + \tau) N(a) \Lambda(\alpha) \mathcal{F}_i(\alpha) \frac{\mathcal{F}_d(\alpha + \tau)}{\mathcal{F}_d(\alpha)} d\tau d\alpha \quad (3)$$

- ▶ $N(\alpha)$: steady-state population
- ▶ $\Lambda(\alpha)$: force of infection at age of infection α
- ▶ $\mathcal{F}_i(\alpha)$: susceptibility survival function

Separable mixing

$$c(a, \alpha) = f(a)g(a)$$

- ▶ assumption
- ▶ eigenvalue problem for K becomes one-dimensional
- ▶ explicit expression for R_0 possible

Approximation of R_0

- ▶ assumptions:
- ▶ 1) separable mixing
- ▶ 2) only one age class
- ▶ 3) $\mathcal{F}_i(a) = e^{-Qa}$
- ▶ 4) g is constant
- ▶ 5) h is independent of α
- ▶ 6) $\mathcal{F}_d(a) = e^{-\mu a}$
- ▶ 7) $r \approx 0$

$$\hat{R}_0 = 1 + \frac{Q}{\mu + r} = \frac{L}{\bar{a}}$$

- ▶ Q : force of infection
- ▶ \bar{a} : mean age of infection
- ▶ L : life expectancy

Estimator for age-specific R_0 : Part I

$$R_0 = \int_0^{\infty} \psi(\alpha) f(\alpha) N(\alpha) d\alpha$$

$$\psi(\alpha) = \int_0^{\infty} h(\tau, \alpha) g(\alpha + \tau) \frac{\mathcal{F}_d(\alpha + \tau)}{\mathcal{F}_d(\alpha)} d\tau$$

Estimator for age-specific R_0 : Part II

- ▶ assumptions:
- ▶ 1) all individuals are susceptible at birth
- ▶ 2) natural mortality is independent of the force of infection
- ▶ $G(a)$: fraction of individuals of age a who will have antibodies to infection

$$G(a) = 1 - \mathcal{F}_i(a) = 1 - e^{-\int_0^a \Lambda(\alpha) d\alpha} \quad (4)$$

Estimator for age-specific R_0 : Part III

- ▶ another estimator for age-specific R_0 :

$$\hat{R}_0 = \frac{\int_0^\infty l(a)\Lambda(a)\mathcal{F}_d(a)da}{\int_0^\infty l(a)\Lambda(a)\mathcal{F}_i(a)\mathcal{F}_d(a)da} \quad (5)$$

Derivation of age-specific R_0

$$R_0 I(\alpha) = \int_0^\infty I(a) k(a, \alpha) da \quad (6)$$

$$R_0 \int_0^\infty \Lambda(\alpha) \mathcal{F}_i(\alpha) \mathcal{F}_d(\alpha) I(\alpha) d\alpha = \int_0^\infty \Lambda(\alpha) \mathcal{F}_i(\alpha) \mathcal{F}_d(\alpha) \quad (7)$$

$$\int_0^\infty I(a) k(a, \alpha) da d\alpha \quad (8)$$

$$(9)$$

- ▶ switch the order of integration on the RHS
- ▶ insert $\frac{N(\alpha)N(a)}{N(\alpha)N(a)}$
- ▶ use $\Lambda(a) = \int \Lambda(\alpha) \mathcal{F}_i(\alpha) \frac{N(\alpha)}{N(a)} k(a, \alpha) d\alpha$
- ▶ population of constant size: $\frac{N(\alpha)}{N(a)} = \frac{\mathcal{F}_d(\alpha)}{\mathcal{F}_d(a)}$
- ▶ simplify RHS & substitute a for α on LHS

Estimator for age-specific R_0 : Part IV

- ▶ impose additional assumption: separable mixing
- ▶ approximation for R_0

$$\hat{R}_0 = \frac{\int_0^\infty \Lambda^2(a) e^{-\mu a} da}{\int_0^\infty \Lambda^2(a) \mathcal{F}_i(a) e^{-\mu a} da} \quad (10)$$

- ▶ estimator of R_0 expressed by age-dependent force of infection $\Lambda(a)$
- ▶ estimate $\Lambda(a)$ for example via maximum-likelihood estimation

Maximum likelihood estimation

- ▶ assumption: $\Lambda(a) = \Lambda_i$
- ▶ n_i : # of seronegative individuals at age interval i
- ▶ m_i : # of seropositive individuals at age interval i
- ▶ binomial likelihood: $\binom{n_i}{m_i} G_i^{m_i} (1 - G_i)^{n_i - m_i}$
- ▶ $G_i = 1 - \exp(-\int_{a_i}^{a_{i+1}} \Lambda(a) da) = 1 - \exp(-\Lambda_i(a_{i+1} - a_i))$
- ▶ $L = \prod_{i=0}^{n-1} \binom{n_i}{m_i} G_i^{m_i} (1 - G_i)^{n_i - m_i}$
- ▶ estimate of force of infection: $\hat{\Lambda}_i = \frac{-\ln(\frac{n_i - m_i}{n_i})}{a_{i+1} - a_i}$

Back to Michael.