# STATS 700-002 Class 3. The Volz Coalescent

Edward Ionides and Aaron King

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## The Volz Coalescent

Volz, E. M., Kosakovsky Pond, S. L., Ward, M. J., Leigh Brown, A. J., & Frost, S. D. (2009). Phylodynamics of infectious disease epidemics. Genetics, 183(4), 1421-1430. https://doi.org/10.1534/genetics.109.106021.

Overview: Kimgman's coalescent provides a link between the observed rate of coalescence, the *effective population size* and the generation interval, in a fixed-size population.

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- Overview: Kimgman's coalescent provides a link between the observed rate of coalescence, the *effective population size* and the generation interval, in a fixed-size population.
- ▶ A time-varying population can be approximated as piecewise constant. Volz et al. (volz09) noticed that this can be inappropriate for fast-changing populations and derived an alternative approach.

## Cluster size distributions

▶ Why are (Volz et al., 2009) so interested in cluster size distribution?

#### The ancestor function

igwedge A(t,T) is defined as the "fraction of the population at t with sampled progeny extant at T.

Does "population" here mean "host population, N" or "population of infected hosts,  $\mathcal{I}$ "?

#### Intuition

▶ From the introduction, "Coalescent rates are low near peak prevalence, but higher when there is a large ratio of incidence to prevalence."

How is this evident?

## Definition or derivation

Consider equation (2):

$$dA(t,T)/dt = f_{SI}p_c.$$

Is this a definition of A(t,T) for some model, or is it derived?

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- lacktriangle Either way, once a model gives us  $p_c(t,T)$ , this defines a coalescent process having coalescent rate  $f_{SI}p_c$ .
- Note, this depends on the expected number of lineages (normalized as the large sample fraction) not the actual number.

## Likelihood

lacktriangle Equation (12) is written in terms of A(t,T), which is a property of the model not of the data.

How can this be considered a likelihood?

## Comparison with the skyline plot

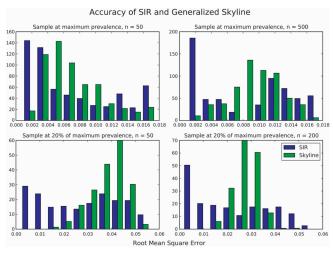


FIGURE 3.—Root mean square error of SIR and generalized skyline estimates of epidemic prevalence. Data are based on 300 simulated epidemics ( $R_0 = 2$ ). RMSE is averaged over 100 time points.

## References I

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