

# STATS 700-002 Class 3.

## The Volz Coalescent

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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: Kingman'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: Kingman'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

- ▶  $A(t, T)$  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?

# Likelihood

- ▶ 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

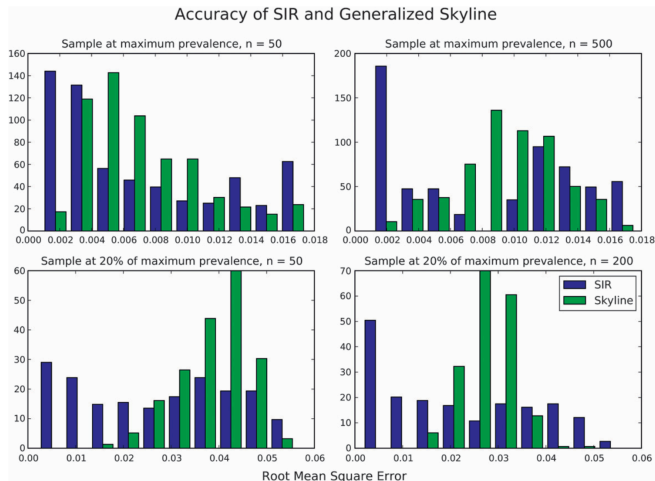


FIGURE 3.—Root mean square error of SIR and generalized skyline estimates of epidemic prevalence. Data are based on 300 simulated epidemics ( $R_0 = 2$ ). RMSE is averaged over 100 time points.



## References I

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