

STATS 700-002 Class 4.
Sampling through time in birth-death trees

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Stadler (2010)

Stadler, T. (2010). Sampling-through-time in birth–death trees. *Journal of Theoretical Biology*, 267(3), 396–404.
<https://doi.org/10.1016/j.jtbi.2010.09.010>.

- ▶ Here, we start developing models for heterochronous genealogical trees (not all leaves are at the same time).

Birth-death trees vs the coalescent

Stadler, T. (2009). On incomplete sampling under birth–death models and connections to the sampling-based coalescent. *Journal of Theoretical Biology*, 261(1), 58–66.

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- ▶ Deals with homochronous sampling (i.e., all samples at the end time, T)
- ▶ Similar to Kingman's coalescent, but not identical
- ▶ What are the advantages and disadvantages?

Definitions

- Notice the deletion of ancestors of unsampled lineages. We will call this *pruning*.

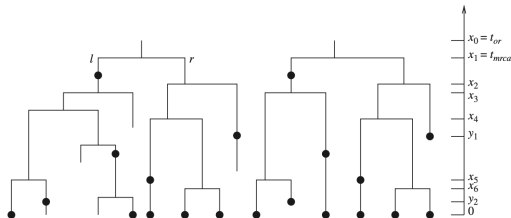


Fig. 1. An example of an oriented tree with root edge induced by the birth-death process displaying all sampled and non-sampled individuals is shown on the left. The left descendant of a bifurcation is l , the right descendant is r . Note that we only label one pair of descendants with l and r for easier readability. The right tree is the corresponding sampled tree with $n=5$ extant individuals, $m=2$ extinct individuals without sampled descendants, and $k=3$ extinct individuals with sampled descendants. In the sampled tree, the bifurcation times are at $x=(x_1, \dots, x_{n+m-1})$ and the sampling times of extinct individuals without sampled descendants are at $y=(y_1, y_2)$. The time of origin is $t_{or}=x_0$ and the time of the most recent common ancestor of the extant species is $t_{mrcu}=x_1$.

Lemma 2.3

► How do you prove the first identity?

$$\sum_{N=n}^{\infty} \binom{N}{n} a^N = \frac{a^n}{(1-a)^{n+1}},$$

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$$\sum_{N=n}^{\infty} \binom{N}{n} a^N = \frac{a^n}{(1-a)^{n+1}},$$

- ▶ You can always ask AI
- ▶ How do you know if AI got it right?

Master equations

$$\begin{aligned}\dot{p}_0(t) &= \mu - (\lambda + \mu + \psi)p_0(t) + \lambda p_0(t)^2, & p_0(0) &= 1 - \rho, \\ \dot{p}_1(t) &= -(\lambda + \mu + \psi)p_1(t) + 2\lambda p_0(t)p_1(t), & p_1(0) &= \rho.\end{aligned}$$

► What are these, and where do they come from?

An informal derivation

let $A_0(t)$, $A_1(t)$ be the events corresponding to $p_0(t)$, $p_1(t)$.

$$\begin{aligned} P[A_0(t+h)] &= E\left[P[A_0(t+h) \mid \text{outcomes in } [t, t+h]]\right] = \\ &\quad (1 - [\lambda + \mu + \phi]h)P[A_0(t)] & (1) \\ &\quad + \lambda h(P[A_0])^2 & (2) \\ &\quad + \mu h & (3) \\ &\quad + o(h) & (4) \end{aligned}$$

Now subtract $P[A_0(t)]$ from both sides, divide by h , and take a limit.

Note: we've passed a limit through expectation without justification.

Theorem 3.1.

How did Stadler guess this solution?

Theorem 3.5.

- ▶ This is a key result, since it gives the likelihood of a given genealogy under the linear birth-death-sampling model.

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- ▶ This is a key result, since it gives the likelihood of a given genealogy under the linear birth-death-sampling model.
- ▶ How is this derived?

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