

# Tree reconstruction from statistical perspectives

Lam Si Tung Ho

Department of Mathematics and Statistics



Contact: [Lam.Ho@dal.ca](mailto:Lam.Ho@dal.ca)

## Statistical inference

- **Data:**  $(Y_i)_{i=1}^n$
  - **Model:**  $(Y_i)_{i=1}^n$  follow a distribution  $\mathcal{P}_{\theta^*}$  where  $\theta^* \in \Theta \subset \mathbb{R}^d$
  - **Estimation method:** approximate  $\theta^*$
- 

## Tree reconstruction

- **Data:** sequences
- **Model:** a substitution model along a true tree  $\mathbb{T}$
- **Reconstruction method:** Maximum likelihood, Bayesian, ...

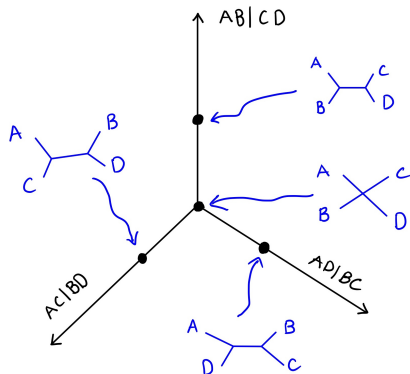
However,  $\mathbb{T} \notin \mathbb{R}^d$ , and the tree topology is a discrete object

Standard statistical theory:  $\hat{\theta}_{\text{MLE}} \rightarrow \theta^*$

- Model identification
- Parameter space  $\Theta$  is compact
- The log likelihood function  $\ell(\theta \mid Y)$  is continuous in  $\theta$  for almost all  $Y$
- $E [\sup_{\theta} |\ell(\theta \mid Y)|] < \infty$

*“Several workers . . . concerned that the discrete, unordered nature of a tree topology variable prevents it from being the sort of parameter required . . .”*

(Rogers, 2001)



## Embedding

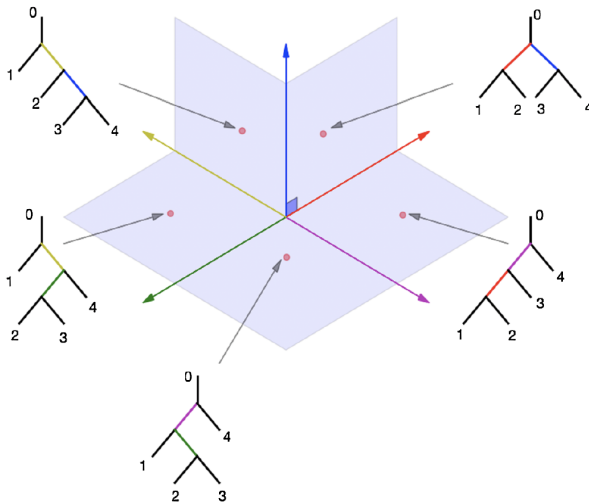
$$\mathbb{T} \hookrightarrow \sum_{s \in \mathcal{S}} e_s \zeta_s$$

- $\mathcal{S}$ : set of all tree splits
- $e_s$ : edge length
- $\zeta_s$ : basis vector

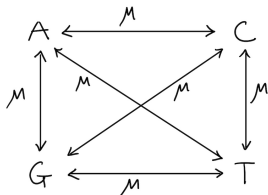
## Distance:

- Branch score distance
- Geodesic distance

# Continuous tree space (Billera-Holmes-Vogtmann)



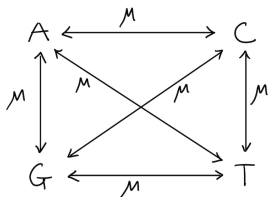
- Model identification
  - ▶ well-studied
- Parameter space  $\mathcal{T} \times \Theta$  is compact
  - ▶ bounded model parameters
  - ▶ bounded branch lengths
  - ▶ external branch lengths are bounded away from 0
- The log likelihood function  $\ell(\mathbb{T}, \theta \mid Y)$  is continuous in  $\mathbb{T}, \theta$ 
  - ▶ often true
- $E \left[ \sup_{\mathbb{T}, \theta} |\ell(\mathbb{T}, \theta \mid Y)| \right] < \infty$



$$P = \begin{pmatrix} \frac{1}{4} + \frac{3}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} \\ \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} + \frac{3}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} \\ \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} + \frac{3}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} \\ \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} + \frac{3}{4}e^{-t\mu} \end{pmatrix}$$

- Model identification
- Parameter space  $\mathcal{T} \times \Theta$  is compact
- The log likelihood function  $\ell(\mathbb{T}, \theta \mid Y)$  is continuous in  $\mathbb{T}, \theta$

$$P(Y \mid \mathbb{T}) = \frac{1}{4} \sum_{(x,y)} \prod_{(u,v) \in E} P[v = y \mid u = x, t = e_{(u,v)}]$$



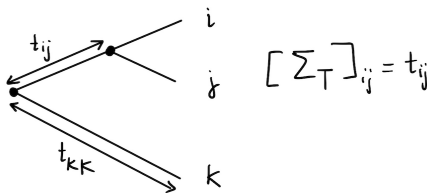
$$P = \begin{pmatrix} \frac{1}{4} + \frac{3}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} \\ \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} + \frac{3}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} \\ \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} + \frac{3}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} \\ \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} - \frac{1}{4}e^{-t\mu} & \frac{1}{4} + \frac{3}{4}e^{-t\mu} \end{pmatrix}$$

$$E [\sup_{\mathbb{T}, \theta} |\ell(\mathbb{T}, \theta \mid Y)|] < \infty$$

- Bound  $P(Y \mid \mathbb{T})$  away from 0 by setting all internal nodes to A
- Probability of transition  $A \rightarrow A$  is at least  $1/4$
- Done since all external edges are bounded away from 0

$$P(Y \mid \mathbb{T}) = \frac{1}{4} \sum_{(x,y)} \prod_{(u,v) \in E} P[v = y \mid u = x, t = e_{(u,v)}]$$

- Rooted trees
- Observe the frequency of alleles
- $Y_i \mid \mathbb{T} \sim_{iid} \mathcal{N}(\kappa \mathbf{1}, \Sigma_{\mathbb{T}})$  (Brownian motion model)



## MLE is a consistent tree reconstruction method

- Use the continuous representation of tree space
- Verify the conditions of Wald (1949) in the form given by Redner (1981)  
(RoyChoudhury et al., 2015)

Frequency model:

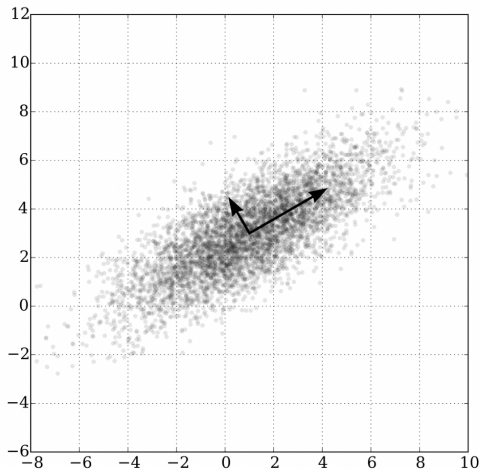
- Model identification
- Parameter space  $\mathcal{T} \times \Theta$  is compact
  - ▶ Without loss of generality, set  $\kappa = 0$ .
- The log likelihood function  $\ell(\mathbb{T}, \theta \mid Y)$  is continuous in  $\mathbb{T}, \theta$

$$\ell(\mathbb{T} \mid Y) = -\frac{1}{2} \sum_{i=1}^n Y_i^T \Sigma_{\mathbb{T}}^{-1} Y_i - \frac{n}{2} \log |\Sigma_{\mathbb{T}}|$$

- $E \left[ \sup_{\mathbb{T}, \theta} |\ell(\mathbb{T}, \theta \mid Y)| \right] < \infty$ 
  - ▶ upper bound  $Y_i^T \Sigma_{\mathbb{T}}^{-1} Y_i$
  - ▶ external edges are bounded away from 0 implies  $\Sigma_{\mathbb{T}} \geq cI$  for some  $c > 0$
  - ▶  $Y_i^T \Sigma_{\mathbb{T}}^{-1} Y_i \leq \frac{1}{c} Y_i^T Y_i$  and  $E(Y_i^T Y_i) = \Sigma_{\mathbb{T}^*}$

- Principal component analysis
- Hamiltonian Monte Carlo
- Regularized Estimation Methods

# Principal component analysis



Wikipedia, CC BY 4.0

- Given trees  $\{T_i\}_{i=1}^n$ , construct a central point  $T_0$ :

$$T_0 = \arg \min_T \sum_{i=1}^n d(x, T_i)^2$$

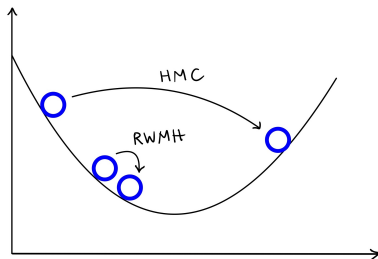
- For a geodesic line  $L$  through  $T_0$ , find the projection:

$$T_i^{(L)} = \arg \min_{T \in L} d(T, T_i)^2$$

- Find the line  $L_{\text{opt}}$  that optimizes an objective function:

$$L_{\text{opt}} = \arg \max_L \sum_{i=1}^n d(T_0, T_i^{(L)})^2$$

(Nye, 2011)

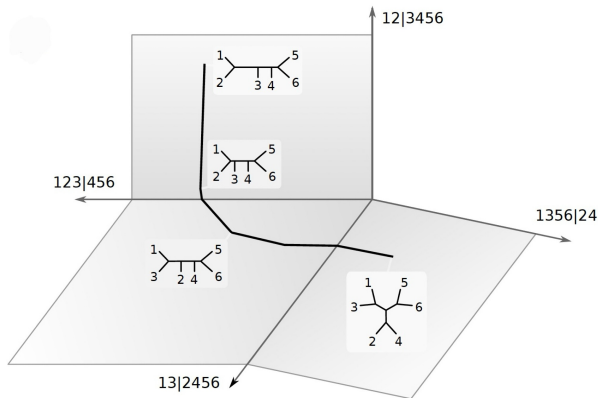


Hamiltonian's equations

$$\frac{dx_i}{dt} = \frac{\partial H}{\partial p_i}, \quad \frac{dp_i}{dt} = -\frac{\partial H}{\partial x_i},$$

where  $H(x, p) = U(x) + K(p)$ , with  $U(x) = -\log f(x)$  and  $K(p) = \|p\|_2^2/2$

# Hamiltonian Monte Carlo for sampling trees



(Dinh et al., 2017)

$$\hat{\theta} = \arg \max_{\theta \in \Theta} \underbrace{\ell(\theta | Y)}_{\text{log likelihood}} - \lambda \underbrace{R(\theta)}_{\text{penalty}} = \arg \min_{\theta \in \Theta} - \underbrace{\ell(\theta | Y)}_{\text{log likelihood}} + \lambda \underbrace{R(\theta)}_{\text{penalty}}$$

- Ridge regression (L2 regularization)

$$R(\theta) = \|\theta - \theta_0\|_2^2 = \sum_{i=1}^d (\theta_i - \theta_0)^2$$

- Lasso (L1 regularization)

$$R(\theta) = \|\theta\|_1 = \sum_{i=1}^d |\theta_i|$$

$$\hat{\mathbb{T}}_{\text{ridge}} = \arg \min_{\mathbb{T} \in \mathcal{T}} -\frac{1}{k} \ell(\mathbb{T} \mid Y) + \lambda_k [d_{\text{geodesic}}(\mathbb{T}, \mathbb{T}_0)]^2$$

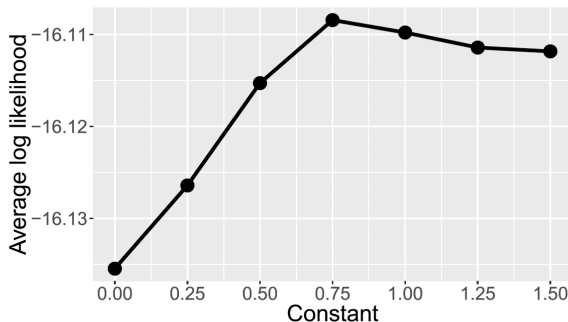
- insufficient signal in the gene sequences
- introduce extra information ( $\mathbb{T}_0$ )

## Convergence rate (Jukes-Cantor)

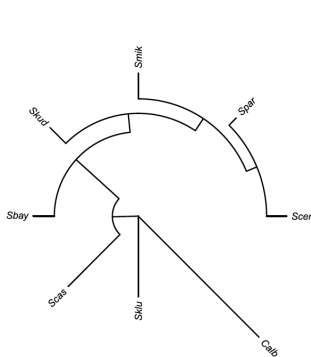
$$d_{\text{geodesic}}(\hat{\mathbb{T}}_{\text{ridge}}, \mathbb{T}^*) = \mathcal{O} \left( \frac{\log k}{\lambda_k \sqrt{k}} + \lambda_k \right)^{1/2}$$

$$\hat{\mathbb{T}}_{\text{ridge}} = \arg \min_{\mathbb{T} \in \mathcal{T}} -\frac{1}{k} \ell(\mathbb{T} \mid Y) + \frac{C}{k^{1/4}} [d_{\text{geodesic}}(\mathbb{T}, \mathbb{T}_0)]^2$$

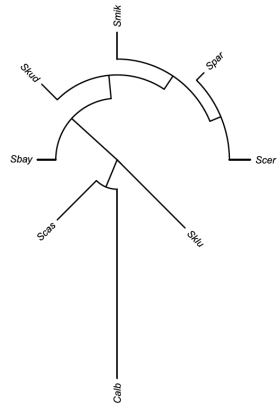
- $\mathbb{T}_0$ : concatenated gene tree
- $C = 0, 0.25, 0.5, 0.75, 1, 1.25, 1.5$



# Yeast gene-tree reconstruction (YKL120W)

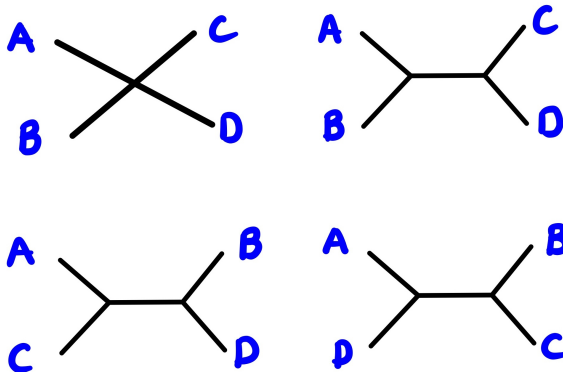


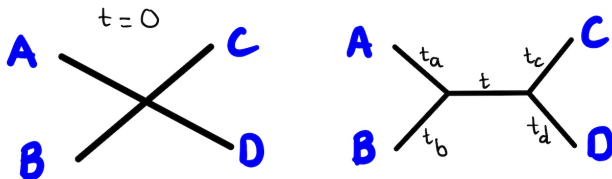
(a) Regularized method



(b) MLE method

# Nonbifurcating tree





Lasso

$$(\hat{t}_a, \hat{t}_b, \hat{t}_c, \hat{t}_d, \hat{t}) = \arg \min -\frac{1}{k} \ell(t_a, t_b, t_c, t_d, t) + \lambda_k (t_a + t_b + t_c + t_d + t)$$

Adaptive Lasso

$$(\tilde{t}_a, \tilde{t}_b, \tilde{t}_c, \tilde{t}_d, \tilde{t}) = \arg \min -\frac{1}{k} \ell(t_a, t_b, t_c, t_d, t) + \eta_k \left( \frac{t_a}{\hat{t}_a^\gamma} + \frac{t_b}{\hat{t}_b^\gamma} + \frac{t_c}{\hat{t}_c^\gamma} + \frac{t_d}{\hat{t}_d^\gamma} + \frac{t}{\hat{t}^\gamma} \right)$$

(Zhang et al., 2021)

## Embedding

$$\mathbb{T} \mapsto \sum_{s \in \mathcal{S}} e_{\mathbb{T},s} \zeta_s$$

## Adaptive Lasso

- Step 1: MLE

$$\hat{\mathbb{T}} = \arg \max_{\mathbb{T} \in \mathcal{T}} \ell_k(\mathbb{T})$$

where  $\ell_k(\mathbb{T})$  is the log likelihood function

- Step 2: regularization

$$\hat{\mathbb{T}}_{\text{AL}} = \arg \min_{\mathbb{T} \in \mathcal{T}} -\frac{1}{k} \ell_k(\mathbb{T}) + \lambda_k \left( \sum_{s \in \mathcal{S}} \frac{e_{\mathbb{T},s}}{e_{\hat{\mathbb{T}},s}^\gamma} \right)$$

## Consistency

- $e_{\hat{\mathbb{T}}_{\text{AL}},s} \rightarrow_p e_{\mathbb{T}^*,s}$
- If  $e_{\mathbb{T}^*,s} = 0$ , then  $e_{\hat{\mathbb{T}}_{\text{AL}},s} = 0$  with high probability

## Lemmas

- Convergence rate of MLE

$$d(\hat{\mathbb{T}}, \mathbb{T}^*) \leq \left( \frac{\log k}{\sqrt{k}} \right)^{1/\beta}$$

- Lojasiewicz inequality

$$\phi(\mathbb{T}) - \phi(\mathbb{T}^*) \geq c_{\mathcal{T}} d(\mathbb{T}, \mathbb{T}^*)^{\beta}, \quad \forall \mathbb{T} \in \mathcal{T}$$

- Concentration inequality

$$\left| \frac{1}{k} \ell_k(\mathbb{T}) - \phi(\mathbb{T}) \right| \leq c \frac{\log k}{\sqrt{k}}, \quad \forall \mathbb{T} \in \mathcal{T}$$

where  $\phi(\mathbb{T}) = E[\ell_1(\mathbb{T})]$

Define

$$M(\mathbb{T}) = \sum_{s \in \mathcal{S}} \frac{e_{\mathbb{T},s}}{e^{\hat{\gamma}_{\hat{\mathbb{T}},s}}}$$

$$\begin{aligned} c_{\mathcal{T}} d(\hat{\mathbb{T}}_{\text{AL}}, \mathbb{T}^*)^\beta &\leq \phi(\mathbb{T}^*) - \phi(\hat{\mathbb{T}}_{\text{AL}}) \\ &\leq c \frac{\log k}{\sqrt{k}} + \frac{1}{k} \ell_k(\mathbb{T}^*) - \frac{1}{k} \ell_k(\hat{\mathbb{T}}_{\text{AL}}) \\ &= c \frac{\log k}{\sqrt{k}} + \frac{1}{k} \ell_k(\mathbb{T}^*) - \lambda_k M(\mathbb{T}^*) \\ &\quad - \frac{1}{k} \ell_k(\hat{\mathbb{T}}_{\text{AL}}) + \lambda_k M(\hat{\mathbb{T}}_{\text{AL}}) + \lambda_k M(\mathbb{T}^*) - \lambda_k M(\hat{\mathbb{T}})_{\text{AL}} \\ &\leq c \frac{\log k}{\sqrt{k}} + \lambda_k M(\mathbb{T}^*) \rightarrow 0 \end{aligned}$$

- Assume that  $e_{\mathbb{T}^*,s} = 0$  and  $e_{\hat{\mathbb{T}}_{\text{AL}},s} > 0$  for some  $s$
- $\mathbb{T}'$  is the same as  $\hat{\mathbb{T}}_{\text{AL}}$ , except  $e_{\hat{\mathbb{T}}_{\text{AL}},s} = 0$

$$\lambda_k \frac{e_{\hat{\mathbb{T}}_{\text{AL}},s}}{e_{\hat{\mathbb{T}},s}} \leq \frac{1}{k} \ell_k(\hat{\mathbb{T}}_{\text{AL}}) - \frac{1}{k} \ell_k(\mathbb{T}') \leq c_{\mathcal{T}} d(\hat{\mathbb{T}}_{\text{AL}}, \mathbb{T}') = c_{\mathcal{T}} e_{\hat{\mathbb{T}}_{\text{AL}},s}$$

On the other hand,

$$e_{\hat{\mathbb{T}},s} \leq d(\hat{\mathbb{T}}, \mathbb{T}^*) \leq \left( \frac{\log k}{\sqrt{k}} \right)^{1/\beta}$$

Contradiction!

- Continuous tree space is helpful if you want to study tree reconstruction from a statistical viewpoint
- Consistency of MLE
- Regularized estimation methods can be good alternatives for MLE

- Stein's Paradox
- “Large  $p$ , small  $n$ ”
- Space of phylogenetic networks

**Contact:** [Lam.Ho@dal.ca](mailto:Lam.Ho@dal.ca)

