

Bayesian inference:

Phylogenetic tree = Evolutionary tree
= tree showing evolutionary relationship among species.

Consider n species, with aligned sequences of length N .

For $D_i \in \{0, 1\}^N$, let $\vec{D} = (D_1, \dots, D_n)$.

(More generally, $D_i \in \{0, 1\}^N$ is a DNA sequence)
 $D_i \in \{A, C, G, T\}^N$

Let $\mathcal{T} =$ phylogenetic trees on n leaves
= trees with internal degree 3 & n leaves.

For $T \in \mathcal{T}$, let $V_{\text{ext}} =$ leaves (there are n leaves)
& $V_{\text{int}} =$ internal vertices.

Each edge in T has a mutation probability $p \in [0, \frac{1}{2}]$

for $e \in E(T)$, let $p(e)$ be its probability.

& $\vec{p} = (p(e))_{e \in E}$

(More generally, can replace $p(e)$ by a 4×4 matrix.)

Given a tree T & probabilities \vec{P} on edges,
 what's the probability of generating this data \vec{D} ?

Consider $D_i \in \vec{D}$ as assignment of $\{0, 1\}$ to each leaf.
 $D_i: V_{ext} \rightarrow \{0, 1\}$.

~~Each~~ An arbitrary internal node chooses a random spin $\{0, 1\}$ & then every edge e :
 flips with prob. $p(e)$
 & stays the same with $1 - p(e)$.

Hence,

$$Pr(D_i | T, \vec{P}) = \frac{1}{2} \sum_{\substack{\hat{D}_i \in \{0, 1\}^V \\ \hat{D}_i(V_{ext}) = D_i(V_{ext})}} \prod_{\substack{e=(y,z) \in E(T) \\ \hat{D}_i(y) = \hat{D}_i(z)}} (1 - p(e)) \prod_{\substack{e \\ \hat{D}_i(y) \neq \hat{D}_i(z)}} p(e).$$

Then,

$$Pr(\vec{D} | T, \vec{P}) = \prod_{i=1}^N Pr(D_i | T, \vec{P})$$

$$= \exp\left(\sum_{i=1}^N \log(Pr(D_i | T, \vec{P}))\right)$$

③

When $p(e)=0$ then no mutations

& $p(e)=\frac{1}{2}$ then endpoints are independent of each other.

Bayes rule implies:

Posterior Probability:

$$\Pr(T|\vec{D}) = \frac{\int_{\vec{p}} \Pr(\vec{D}|T, \vec{p}) \psi(T, \vec{p}) d\vec{p}}{\Pr(\vec{D})}$$

$$= \frac{\int_{\vec{p}} \Pr(\vec{D}|T, \vec{p}) \psi(T, \vec{p}) d\vec{p}}{\sum_{T'} \int_{\vec{p}} \Pr(\vec{D}|T', \vec{p}) \psi(T', \vec{p}) d\vec{p}}$$

For tree T , let $w(T) = \int_{\vec{p}} \Pr(\vec{D}|T, \vec{p}) \psi(T, \vec{p}) d\vec{p}$

Note, $\psi(T, \vec{p})$ is a prior, & we need that

$$\sum_{T'} \int_{\vec{p}} \psi(T', \vec{p}) d\vec{p} = 1$$

For uniform priors, $\psi(T, \vec{p}) = \psi(T', \vec{p})$ then:

$$\Pr(T|\vec{D}) = \frac{\int_{\vec{p}} \Pr(\vec{D}|T, \vec{p}) d\vec{p}}{\Pr(\vec{D})} = \frac{\Pr(\vec{D}|T)}{\Pr(\vec{D})}$$

So for uniform priors,

Posterior Distribution = likelihoods

$$\Pr(T | \vec{D}) = \Pr(\vec{D} | T)$$

Hence, sampling from posterior distribution is more general than sampling proportional to likelihoods

What priors? Mr Bayes allows uniform or exponential distributions for priors.

For ψ with a simple formula,

$$w(T) = \int_{\vec{p}} \Pr(\vec{D} | T, \vec{p}) \psi(T, \vec{p}) d\vec{p}$$

can be computed efficiently via Dynamic Programming.

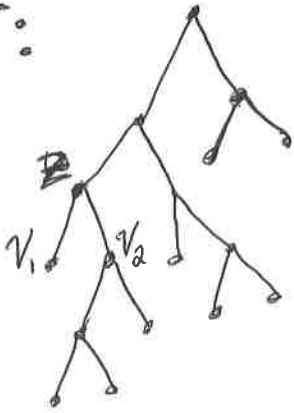
DP idea:

Consider a tree T :
(root it somewhere)

For $v \in V$,

let $T_v =$ tree

hanging below v .



For $v \in V, s \in \{0, 1\}$,

Let $w(T, v, s) = \Pr(\vec{D} | T_v, s \text{ at } v)$

Then $w(T, z, 0) = \prod_{i=1}^2 [p \times w(T_{v_i}, v_i, 1) + (1-p) \times w(T_{v_i}, v_i, 0)]$

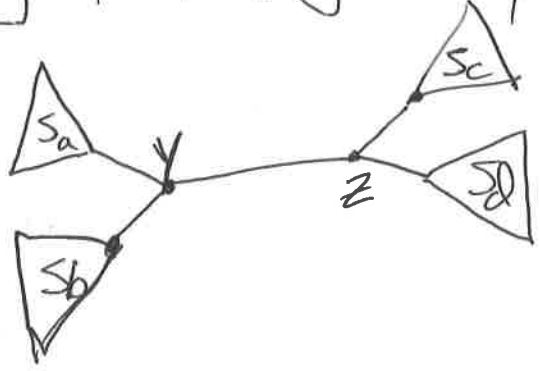
NNI MC: (nearest neighbor interchanges)

From $X_t \in \mathcal{Z}$,

1. Pick an internal edge (y, z) uniformly at random

2. Let ~~S_a~~ S_a & S_b be the subtrees hanging off y
& S_c & S_d " z

(Using that degree of y & z are 3)



3. There are 3 ways of hanging these

4 subtrees S_a, S_b, S_c, S_d off of $e=(y,z)$.

4. Choose one of these 3 at random.

Set T' as new tree

5. Set $X_{t+1} = \begin{cases} T' & \text{with prob. } \min\left\{1, \frac{w(T')}{w(X_t)}\right\} \\ X_t & \text{o/w.} \end{cases}$

SPR = subtree pruning & regrafting.

SPR moves:

1. Choose a random internal edge $e = (y, z)$.
2. Two subtrees hanging off it, T_y & T_z .
3. Choose one of these 2 (T_y or T_z) at random.
4. Say T_y , in $T \setminus T_y$, choose random edge e' & insert T_y onto e' .

Fix tree T^* & \vec{P}^*

Generate \vec{B} from $\mathcal{M}_{T^*, \vec{P}^*}$ = random spin at internal node, then flip with prob. $p(e)$, same with $1-p(e)$.

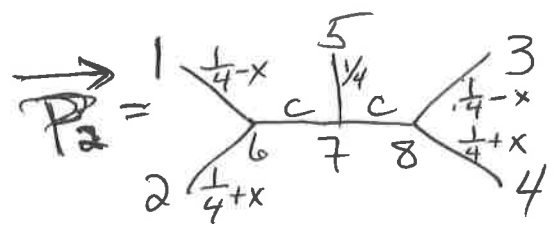
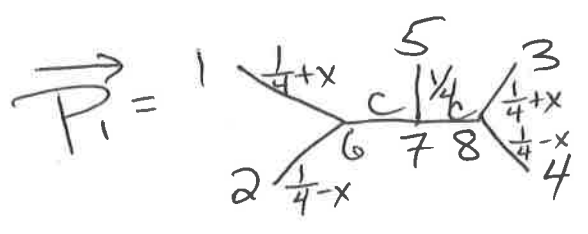
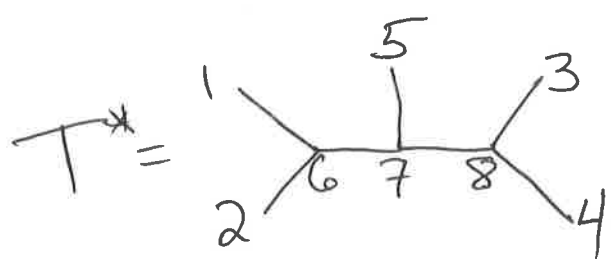
What is mixing time?

Tree T^* is max likelihood as $N \rightarrow \infty$
 i.e., $\max_T w(T) = w(T^*)$

Rapid mixing for all T^*, \vec{P}^* ?

Consensus tree = summary tree from multiple MCMC runs.

5-taxon tree:



$$c = \frac{1}{2} \left(1 - \sqrt{\frac{1-16x^2}{1+16x^2}} \right)$$

Let
$$\mu = \frac{1}{2} \left(\mu_{T^*, \vec{P}_1} + \mu_{T^*, \vec{P}_2} \right)$$

$\exists x_0 > 0, \forall x < x_0,$ with Prob. $\geq 1 - \exp(-\Omega(N))$,

$$T_{mix} = \exp(\Omega(N))$$

for NNI chain.

Open: What about MC³ = Metropolis-coupled MCMC,
 Can we construct a bad example?