(see Felsenstein, J. (1973). Maximum likelihood and minimum-steps methods for estimating evolutionary trees from data on discrete characters. *Systematic Biology*, *22*(3), 240-249.)

Rooted tree T=(V,E) with root  $r\in V$ . A leaf of T is a node with no children and we call the set of all leaves L(T). Consider a discrete, finite alphabet  $\Sigma$  and let  $d\coloneqq |\Sigma|$ .

Let  $\mathcal{D}$  be a data matrix  $\in \mathbb{N}^{|L(T)| \times d}$  that encodes the symbol that has been observed at each leaf.

An *evolutionary tree* is a tuple  $(T, \theta)$  with a rooted tree T = (V, E) where each node  $V_i \in V$  is a random variable with values in  $\Sigma$  and parameters  $\theta = (\tau, Q, \pi)$ . For each  $e \in E$   $\tau_e$  is the evolutionary time along the tree edge,  $Q \in \mathbb{R}^{d \times d}$  is a rate matrix and  $\pi$  is the equilibrium distribution at the root (also see: <u>probabilistic model of evolution</u>).

**Goal:** Estimate  $P(\mathcal{D}|T,\theta)$ .

Let  $\mathcal{D}_{|u}$  for any  $u \in V \setminus L(T)$  denote the data restricted to leaves below u.

## **Algorithm (dynamic programming)**

Input:  $T_{\sigma}$ ,  $\mathcal{D}$ 

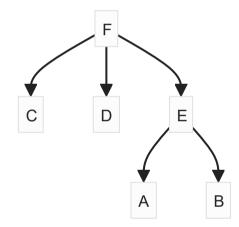
**Output:**  $lpha(u,v)=P(\mathcal{D}_{|u}|V_u=v,T, heta)$  for all  $u\in V\setminus L(T)$ 

The  $\alpha(u, v)$  are computed dynamically starting with leaf edges. This depends on the  $P_{a,b}$  of a substitution model (see: <u>probabilistic model of evolution</u>).

Then we have  $P(\mathcal{D}|T,\theta) = \Sigma_v \alpha(r,v) * \pi_v$ 

## **Example**

T



 $\mathcal{D}$ 

|   | Α | В | С | D |
|---|---|---|---|---|
| 1 | 1 | 0 | 0 | 0 |
| 2 | 0 | 1 | 0 | 0 |
| 3 | 0 | 0 | 0 | 0 |
| 4 | 0 | 0 | 1 | 1 |

$$d=4$$

Let  $\tau = 1$  for all edges.

Model: Jukes-Cantor

$$P_{i,i}^ au = rac{1}{4} + rac{3}{4} ext{exp}(-rac{4}{3} au)$$

$$P_{i,j}^{ au} = rac{1}{4} - rac{1}{4} \exp(-rac{4}{3} au)$$

Compute P(D|T) = P(A = 1, B = 2, C = D = 4|T) with the following steps:

1. 
$$P(A=1|E)=P^{ au_{E,A}}D_{,A}pprox(rac{7}{16},rac{3}{16},rac{3}{16},rac{3}{16})$$

2. 
$$P(B=2|E)=P^{ au_{E,B}}D_{,B}pprox (rac{3}{16},rac{7}{16},rac{3}{16},rac{3}{16})^T$$

3. 
$$P(C=4|F)=P^{ au_{F,C}}D_{;C}pprox (rac{3}{16},rac{3}{16},rac{7}{16})^T$$

4. 
$$P(D=4|F)=P^{ au_{F,D}}D_{;D}pprox (rac{3}{16},rac{3}{16},rac{7}{16})^T$$

5. 
$$P(A=1,B=2|E) = P(A|E)P(B|E) \approx (\frac{21}{256},\frac{21}{256},\frac{9}{256},\frac{9}{256})^T$$

- 6.  $P(A=1,B=2|F)=P^{ au_{E,D}}P(A=1,B=2|E)pprox (0.064,0.064,0.053,0.053)^T$
- 7.  $P(A=1,B=2,C=4,D=4|F)=P(A=1,B=2|F)P(C=4|F)=P(D=4|F)\approx (0.002,0.002,0.0018,0.01)^T$  Symbol 4 is more likely at the root than 1,2,3, since we observed it 2 times at C and D and require 2 substitutions for A and B, whereas for any other symbol, we require at least 3 mutations.