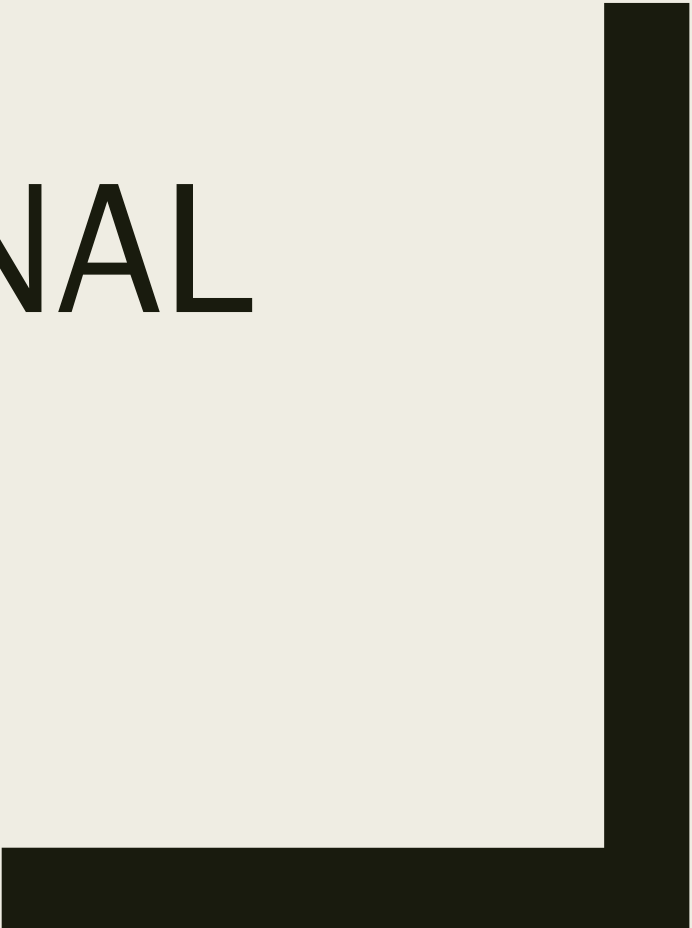


CS 364
COMPUTATIONAL
BIOLOGY

Sara Mathieson
Haverford College



Outline

- Parsimony
- Fitch's Algorithm
- Sankoff's Algorithm





Parsimony Problem

Ancestral State Reconstruction





- Given:
 - States at the leaves
 - Tree topology (and sometimes branch lengths)
- Find the ancestral states at each internal vertex



Characters: Heritable traits

	Wings	Flying	Swimming	Cute
	Present	Absent	Absent	Present
	Present	Present	Absent	Absent
	Absent	Absent	Present	Absent
	Absent	Absent	Absent	Present

Characters: Heritable traits

	Wings	Flying	Swimming	Cute
	1	0	0	1
	1	1	0	0
	0	0	1	0
	0	0	0	1

Images: BBC

Characters: DNA sequence



ACTGAGATAGGGAGGGCGGA



ACTCAGATAGGGAGAGGAGA



ACTGAGAGAGGGAGAGCAGA



ACTGAGAGAGGGAGAGGAGA

Characters: DNA sequence



00010010111110010010



00011010111111010010



00011010111010010010



000100101001-0010010

RNA, Protein sequence etc...

Evolution in Great Apes

List characteristics or traits

- Orange beard
- Miniaturized hair
- Way of movement
- ...



Orangutan
(Bornean)



Orangutan
(Sumatran)



Gorilla

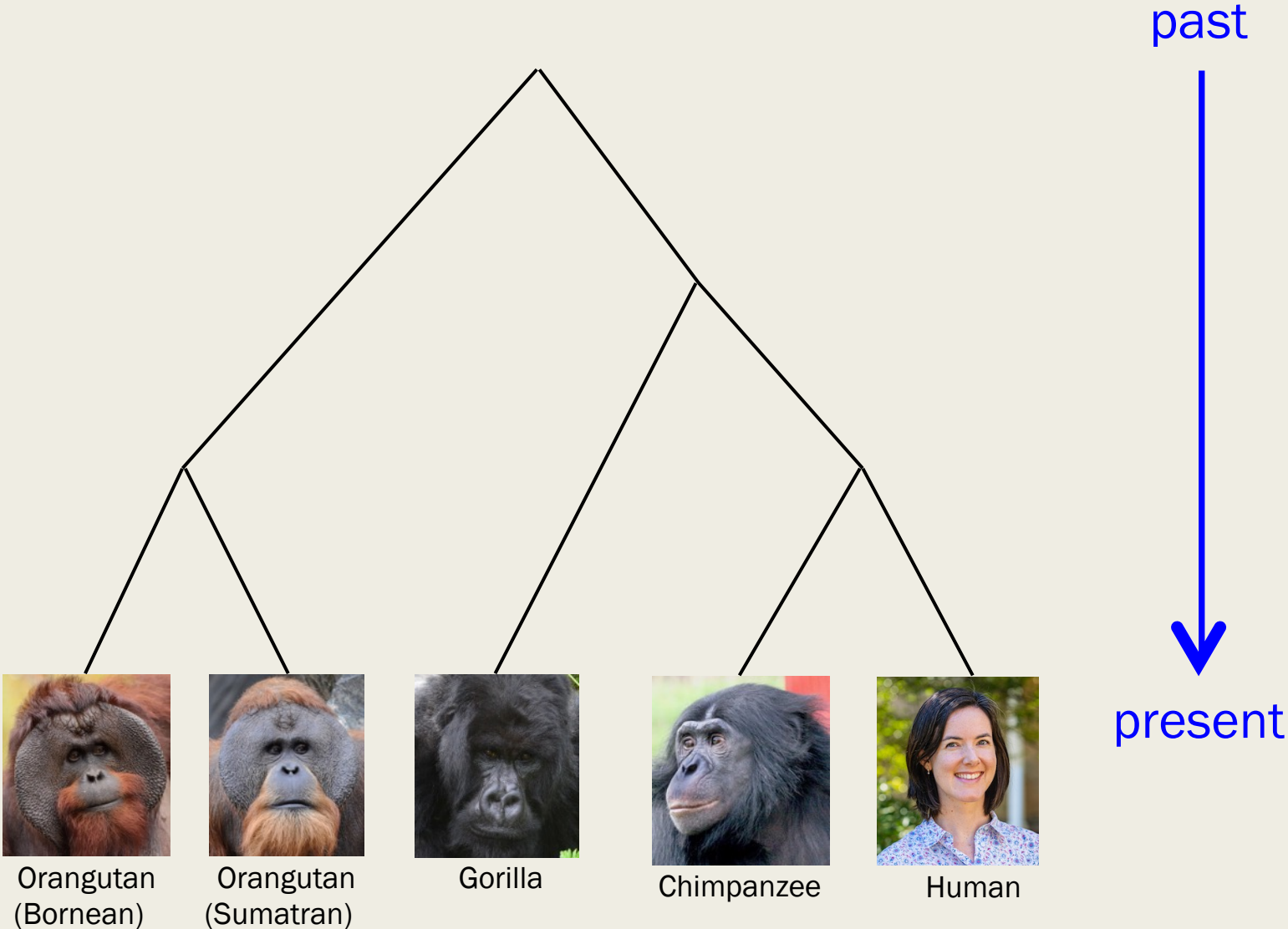


Chimpanzee

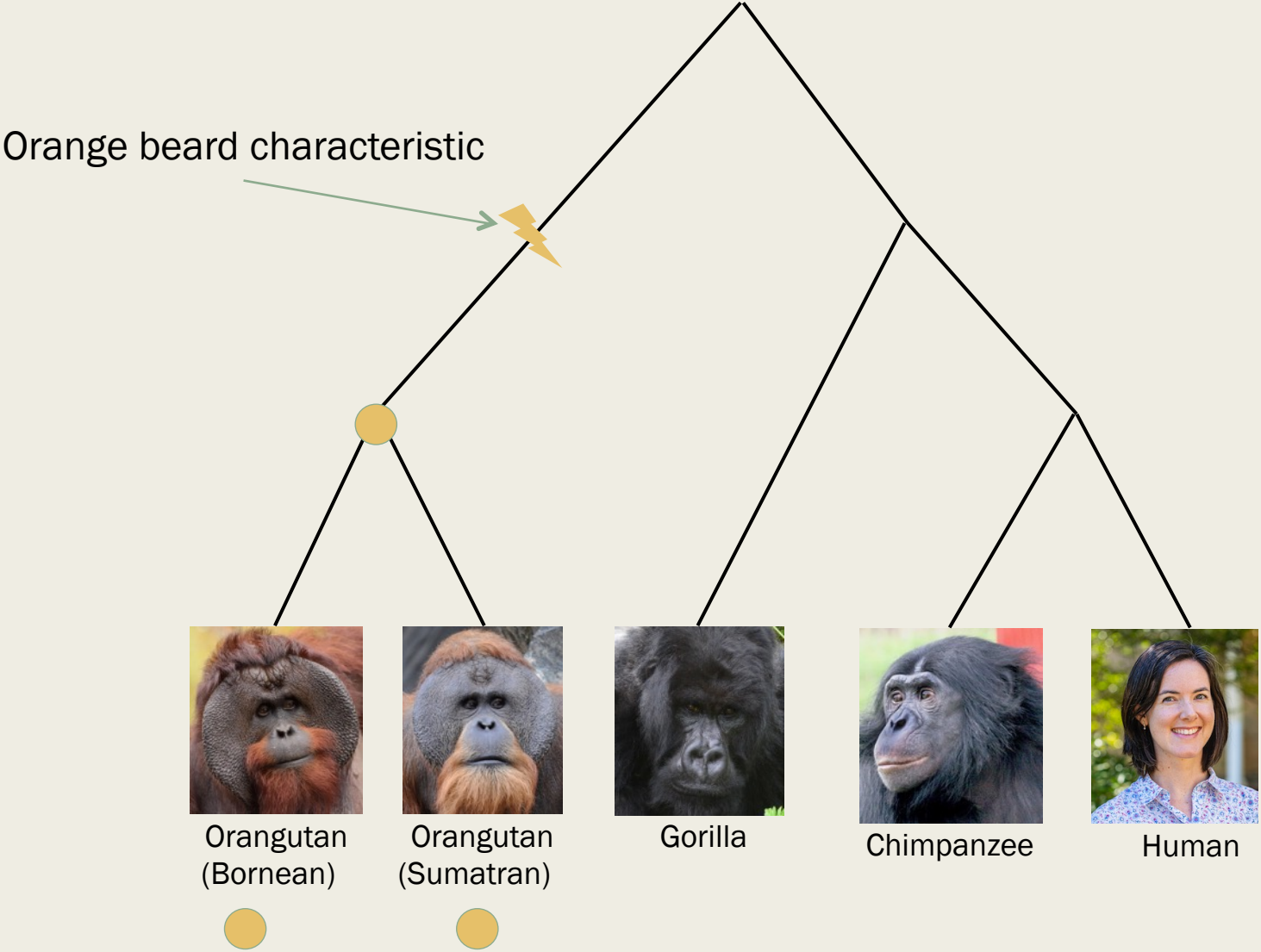


Human

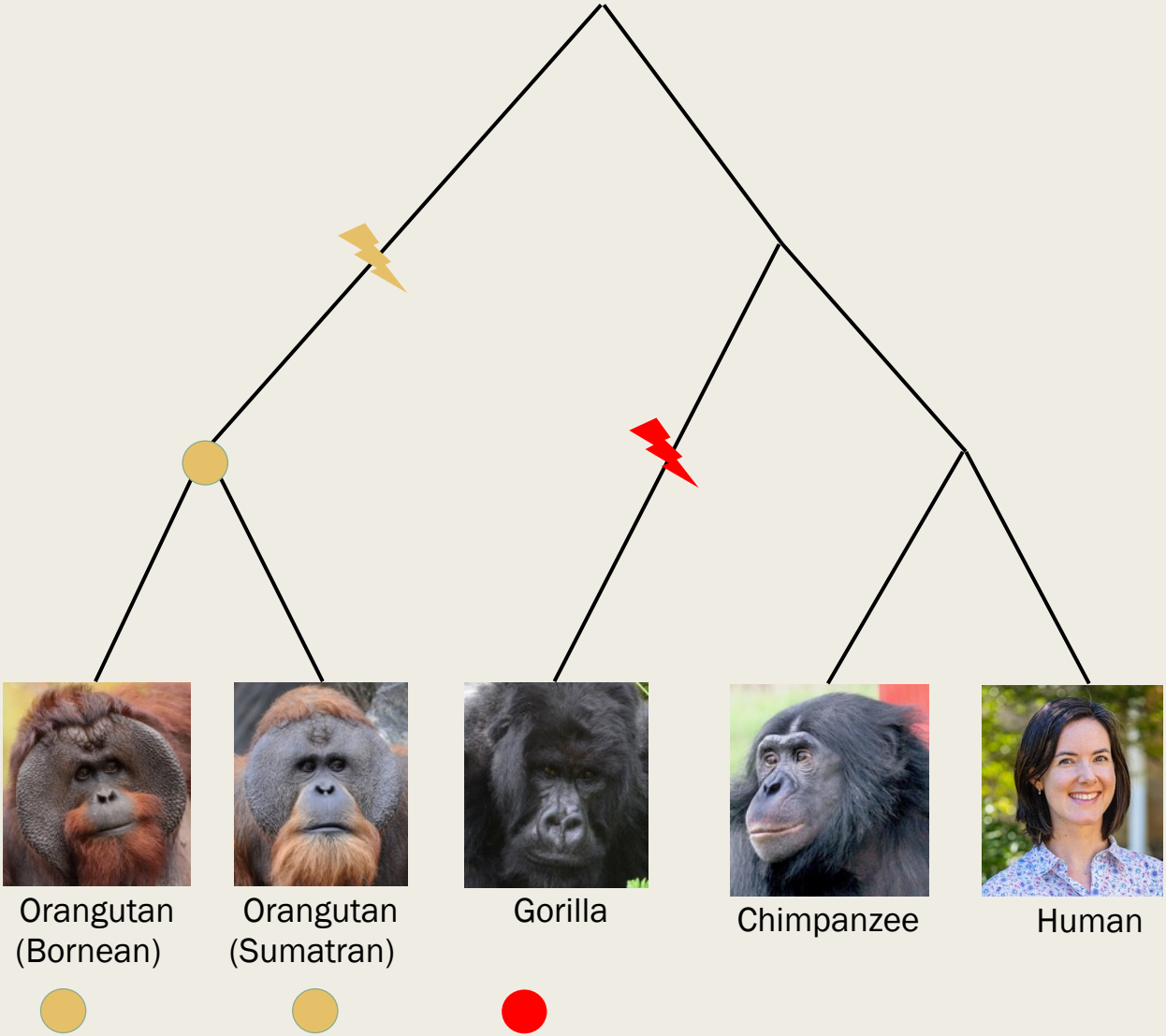
Evolution in Great Apes



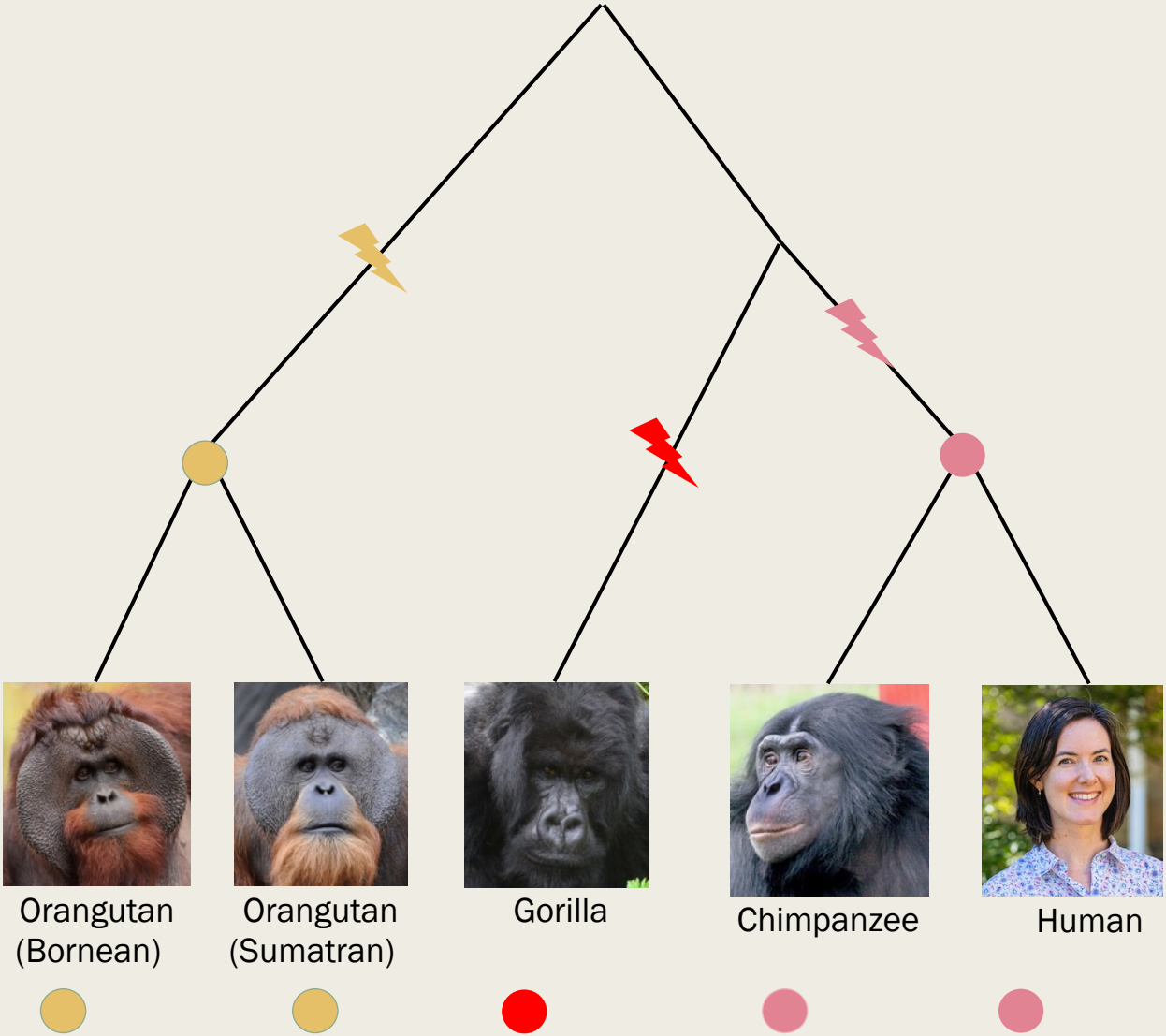
Evolution in Great Apes



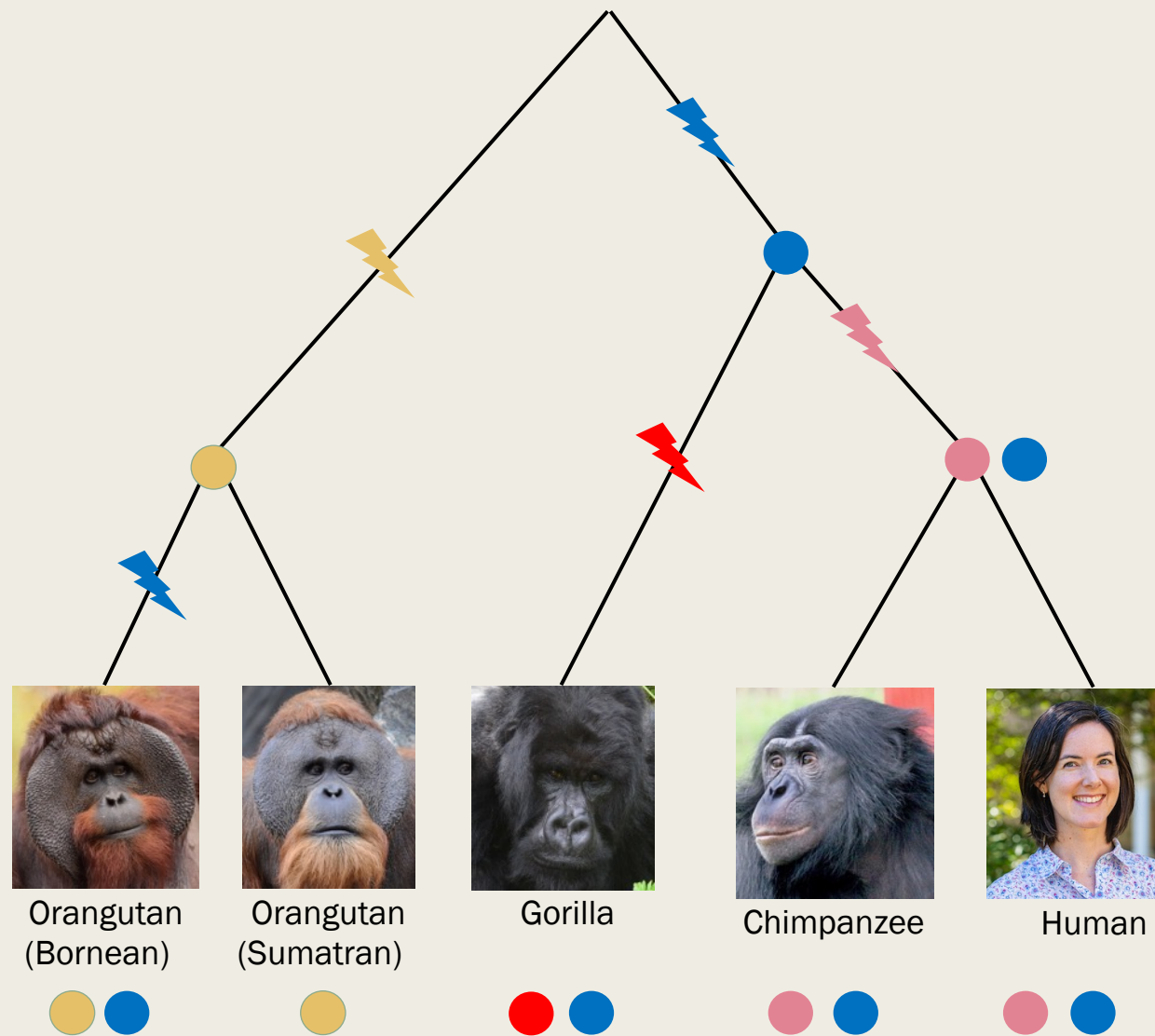
Evolution in Great Apes



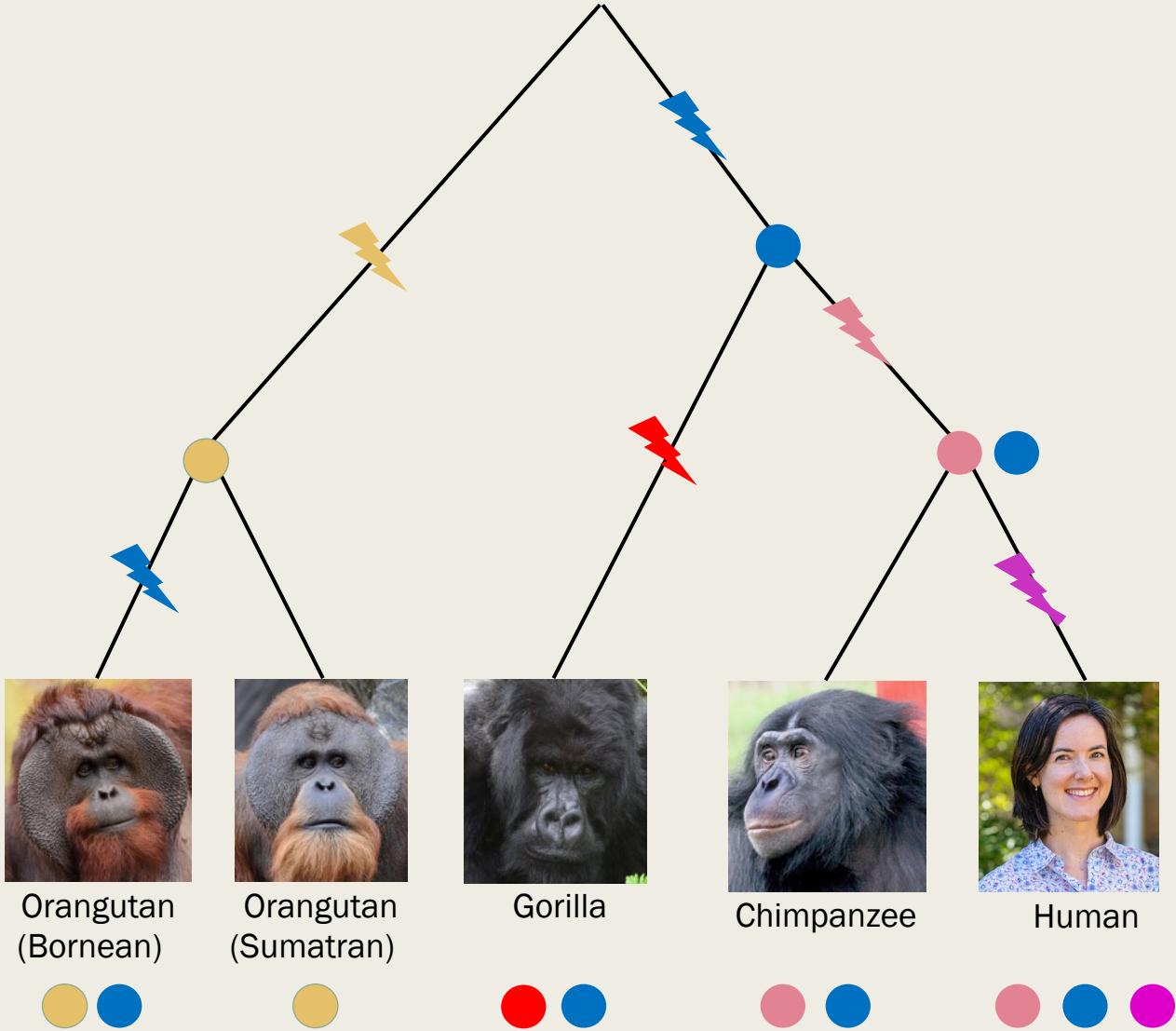
Evolution in Great Apes



Evolution in Great Apes



Evolution in Great Apes



Evolution in Great Apes

We do not observe the evolutionary (phylogenetic) tree.

Only the mutations/characters at the leaves

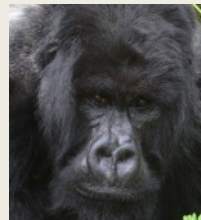
If we *knew* the shape of the tree, could we reconstruct the characters of the ancestors?



Orangutan
(Bornean)



Orangutan
(Sumatran)



Gorilla



Chimpanzee



Human

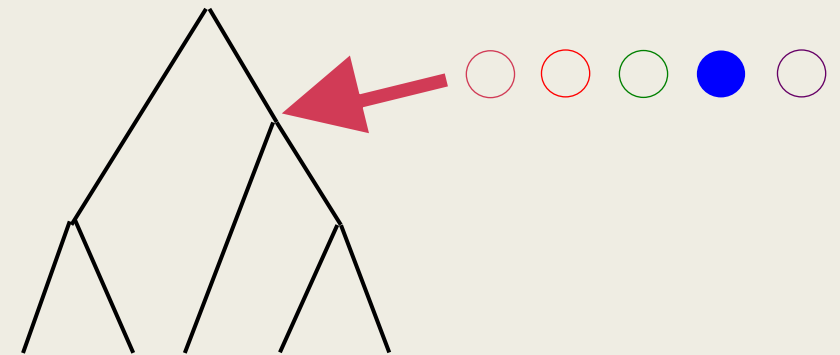


GOAL

■ Input:

- *Presence/absence of characteristics (states) at leaves*
- *Tree topology*

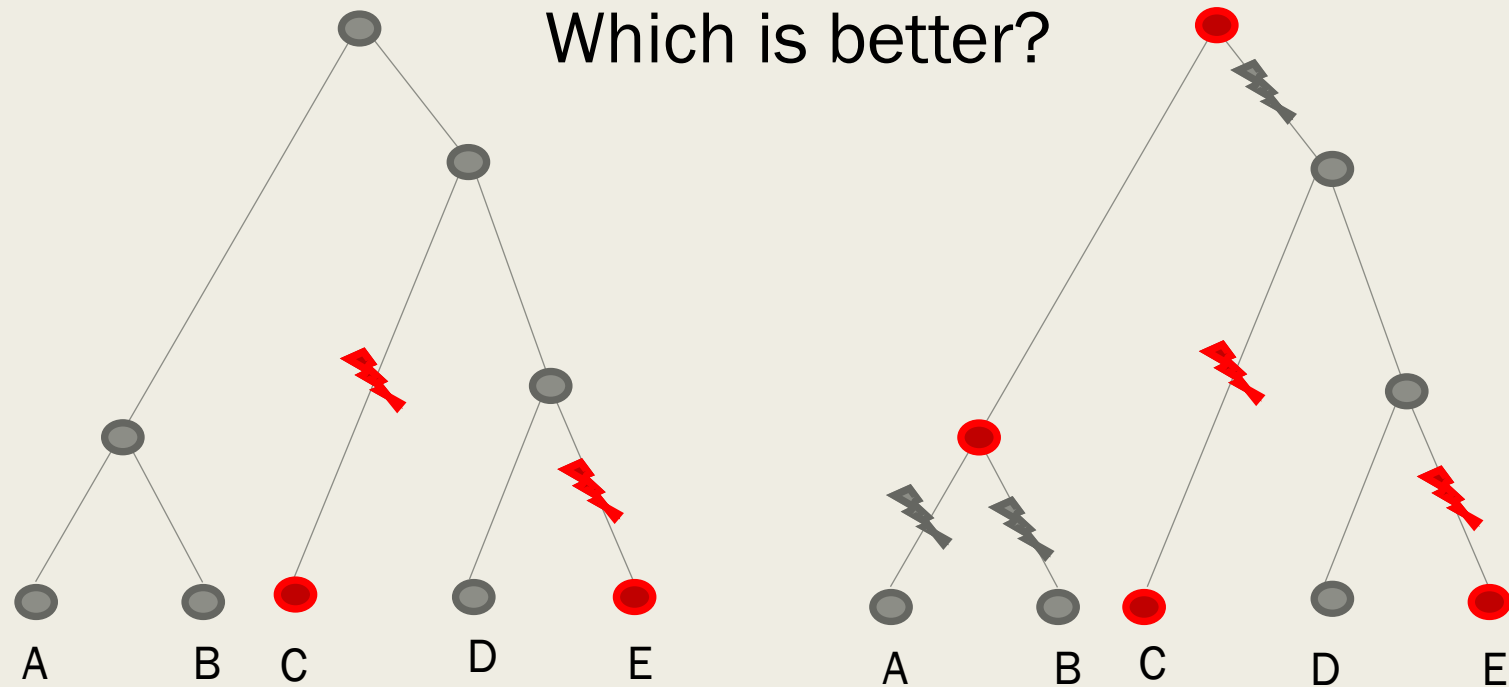
Orangutan (Bornean)	●	○	○	●	○
Orangutan (Sumatran)	●	○	○	○	○
Gorilla	○	●	○	●	○
Chimpanzee	○	○	●	●	○
Human	○	○	●	●	●



Output:

Ancestral states at each internal node (and root)

Parsimony



The parsimony principle: “Everything else being equal, the best explanation is the one that requires the fewest evolutionary changes”

Two parsimony problems

- The “Small” parsimony problem – given the character states at the leaves, and the topology of the tree, reconstruct the states at the ancestral nodes.
- The “Big” parsimony problem – given the character states at the leaves, reconstruct the tree, and the states at the ancestral nodes.

Two parsimony problems

- The “Small” parsimony problem – given the character states at the leaves, and the topology of the tree, reconstruct the states at the ancestral nodes.

Fitch’s algorithm (Fitch, 1971)

Sankoff’s algorithm (Sankoff, 1975)

The parsimony principle: “Everything else being equal, the best explanation is the one that requires the fewest evolutionary changes”

Fitch's algorithm (unweighted parsimony)

Example of Fitch's algorithm in the literature

BIOINFORMATICS ORIGINAL PAPER

Vol. 29 no. 23 2013, pages 2987–2994
doi:10.1093/bioinformatics/btt527

Genome analysis

Advance Access publication September 24, 2013

FPSAC: fast phylogenetic scaffolding of ancient contigs

Ashok Rajaraman^{1,2}, Eric Tannier^{3,4} and Cedric Chauve^{1,5,*}

¹Department of Mathematics, Simon Fraser University, Burnaby (BC) V5A1S6, Canada, ²International Graduate Training Center in Mathematical Biology, Pacific Institute for the Mathematical Sciences, Vancouver (BC), Canada, ³INRIA Grenoble Rhône-Alpes, Montbonnot 38334, France, ⁴Université de Lyon 1, Laboratoire de Biométrie et Biologie Évolutive, CNRS UMR5558 F-69622 Villeurbanne, France and ⁵LaBRI, Université Bordeaux I, 33405 Talence, France

Associate Editor: Michael Brudno

Black death pandemic in Europe: 1347-1351
Killed 30-60% of Europe's population

In this paper, they used Fitch's algorithm to help reconstruct the genome of the ancient Black death pathogen

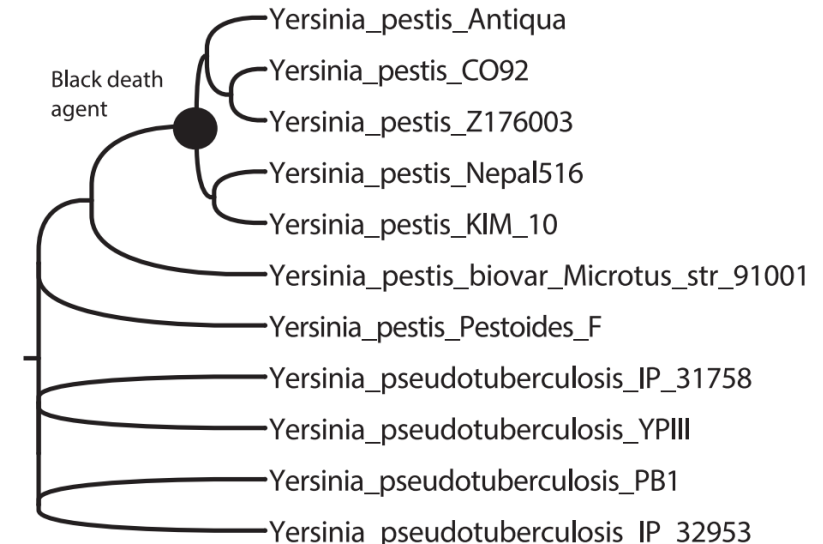
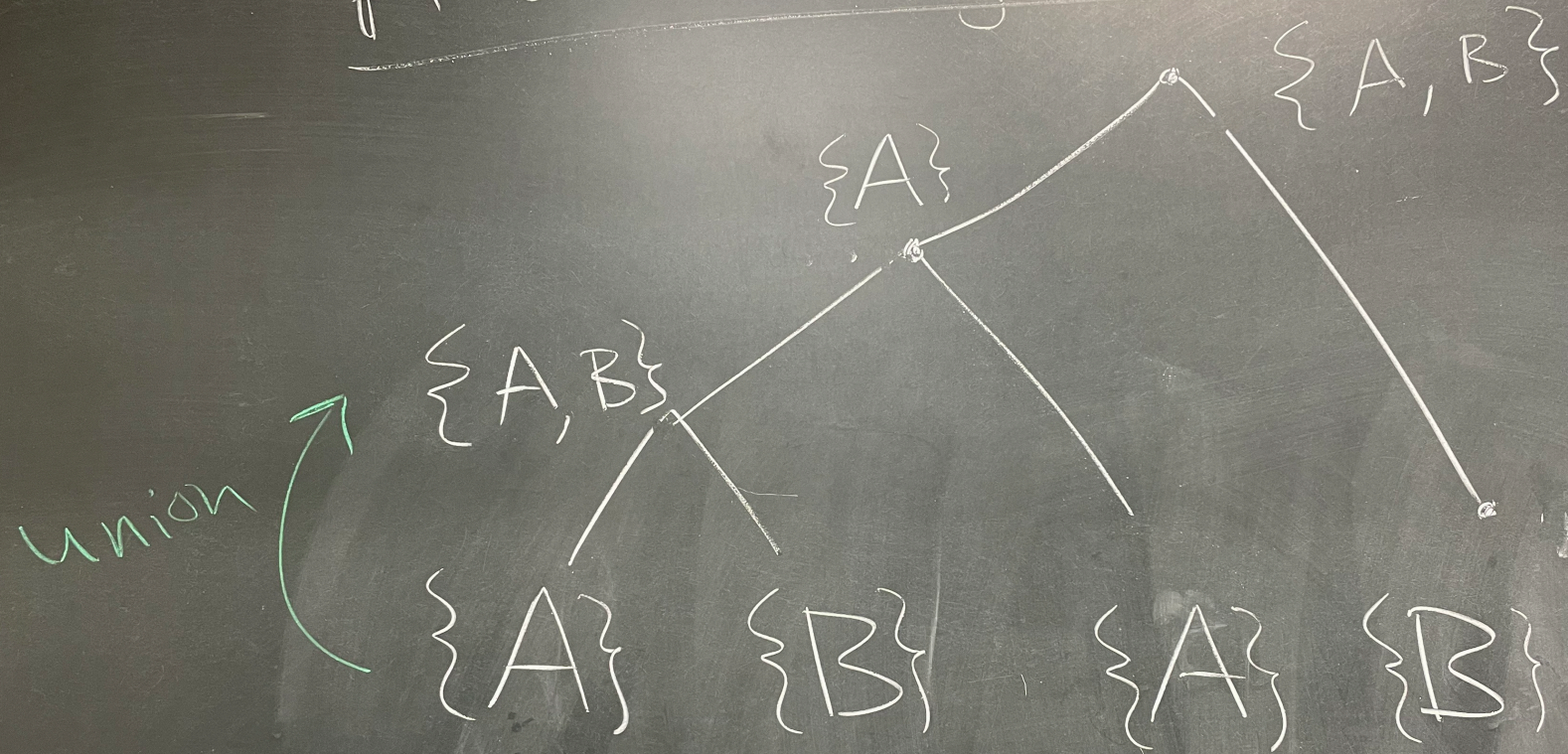


Fig. 3. Phylogeny of the considered genomes from Bos *et al.* (2011)

Fitch's Algorithm



bottom-up phase

- S_v = state set at node v
- $S_v = \{x\}$ if v is a leaf assigned base x

• internal vertices:

let c_1 & c_2 = children of v

$$S_v = \begin{cases} S_{c_1} \cap S_{c_2} & \text{if } S_{c_1} \cap S_{c_2} \neq \emptyset \\ S_{c_1} \cup S_{c_2} & \text{otherwise (o.w.)} \end{cases}$$

intersection

union

empty set

top-down phase

- assign root arbitrarily

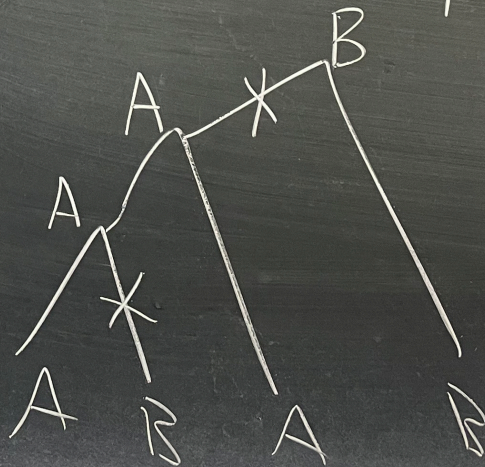
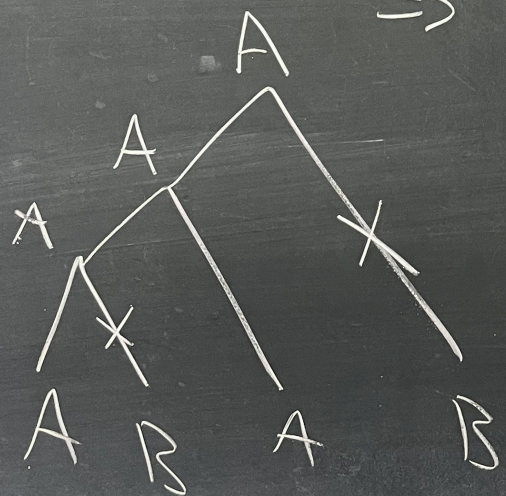
- assign node c :

if parent of c is $x \neq x \in S_c$

\Rightarrow then assign x to c

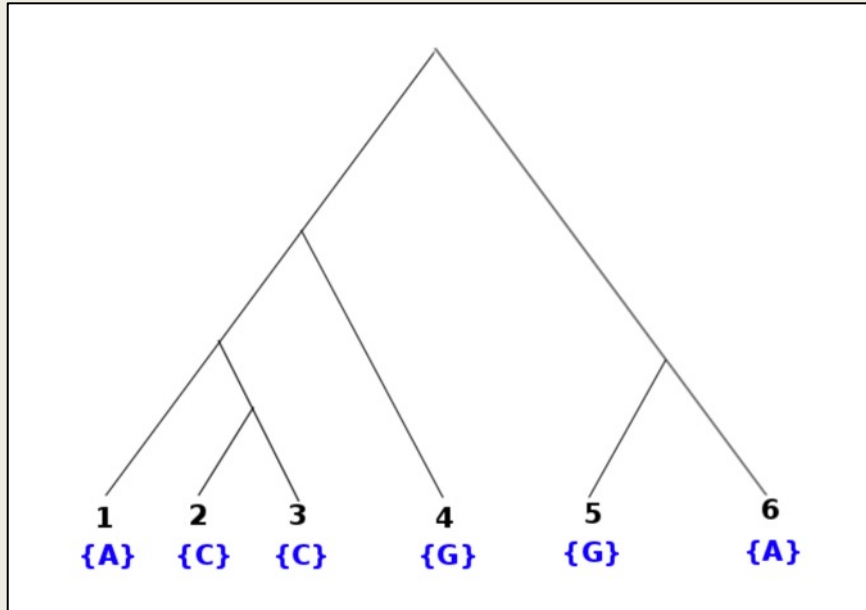
otherwise

\Rightarrow choose arbitrarily.



parsimony
score
= 2

Fitch's algorithm

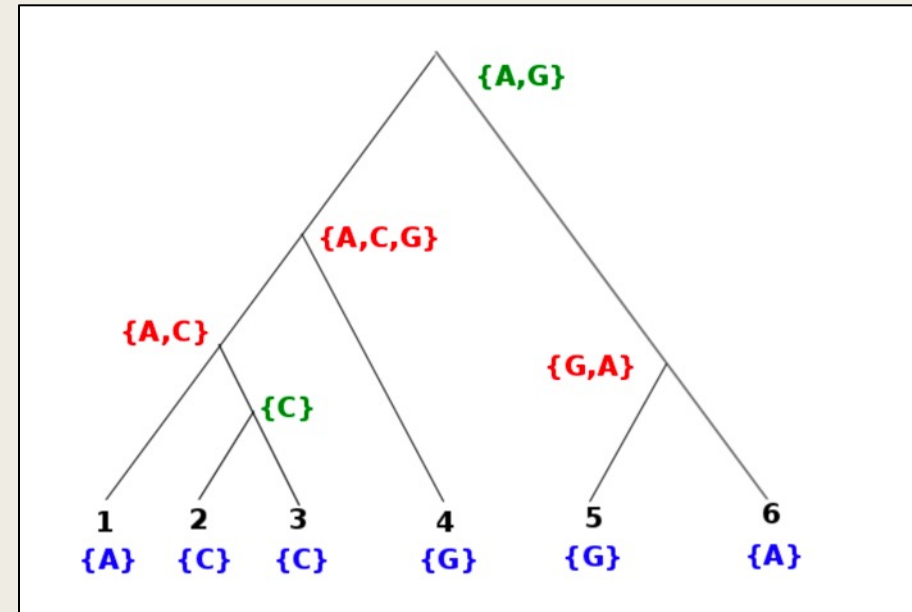
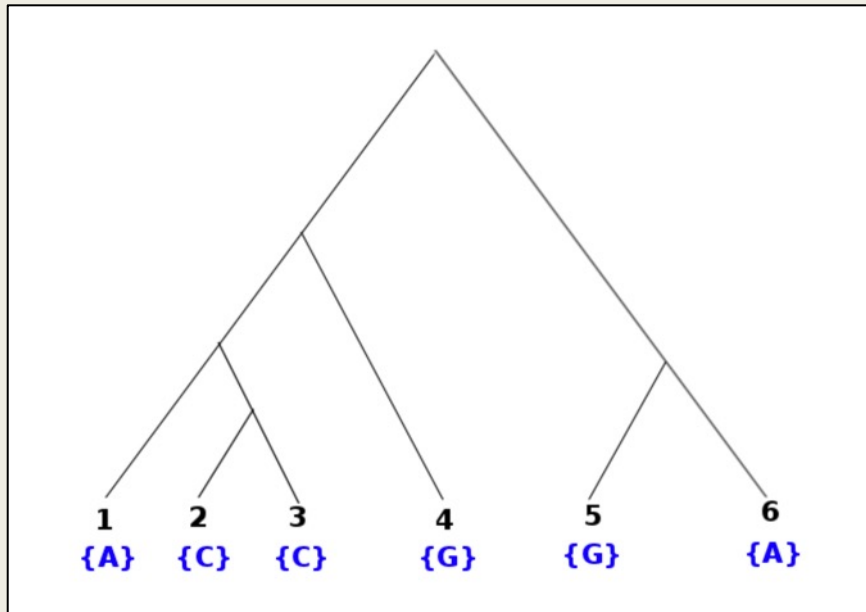


1. Traverse the tree from leaves to root, at each step assign to the parent node the set that is the intersection of the set of characters in the child nodes if it's non empty, but the union if the intersection is empty.

2. Traverse the tree from root to leaves. Assign state of the root randomly from its set. At each node, if the parent state is in that node's set, assign that. Otherwise, assign randomly.

Fitch's algorithm

1. Traverse the tree from leaves to root, at each step assign to the parent node the intersection of the set of characters in the child nodes if it's non empty, but the union if the intersection is empty.

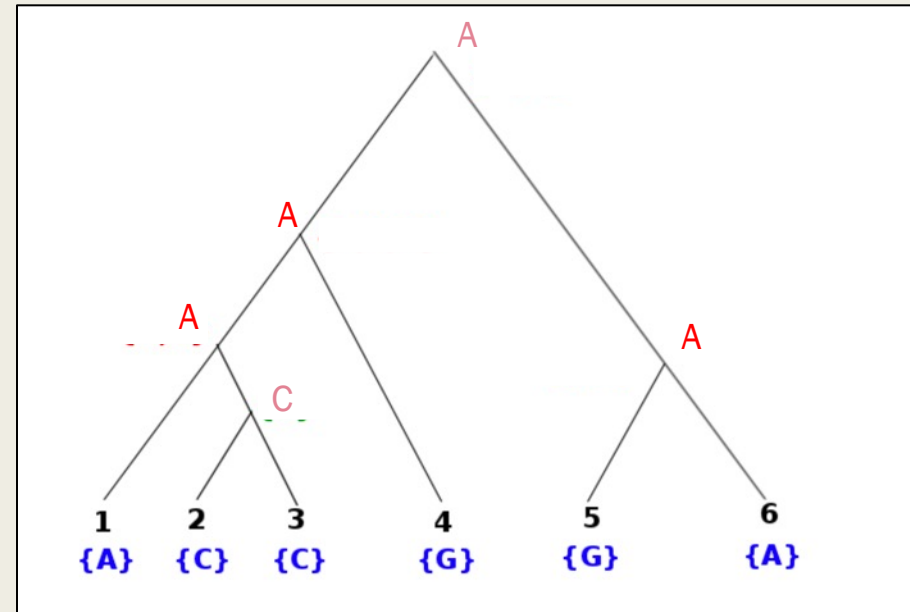
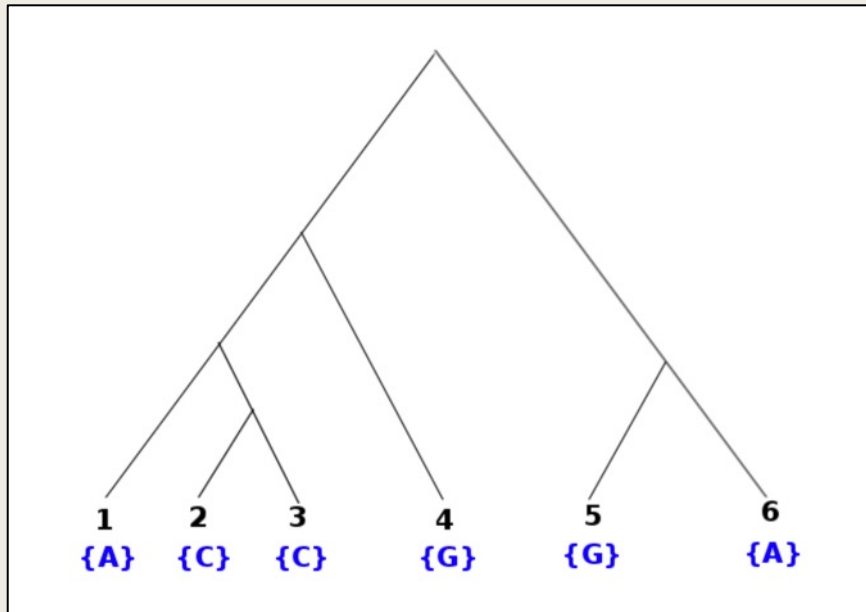


Intersection: no mutation

Union: mutation no matter what we choose

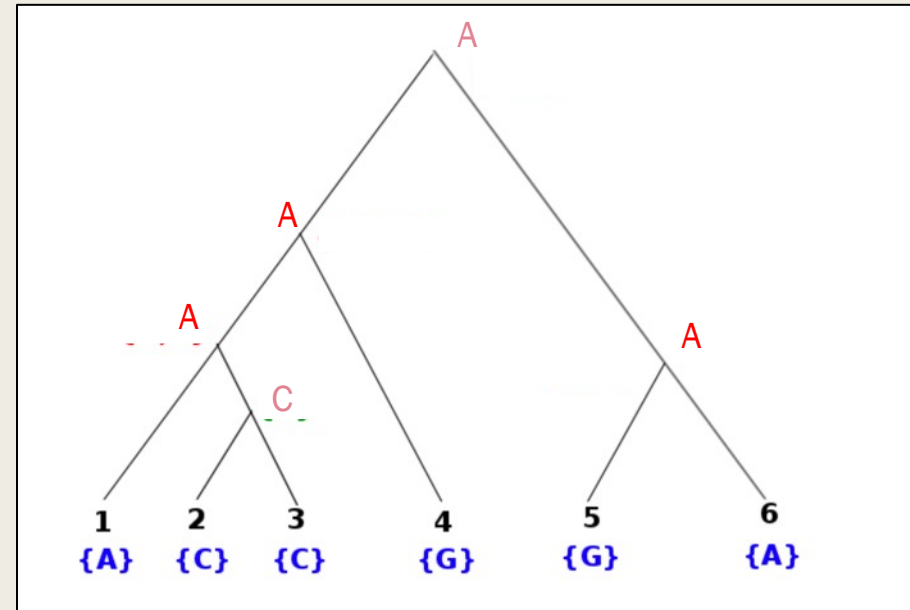
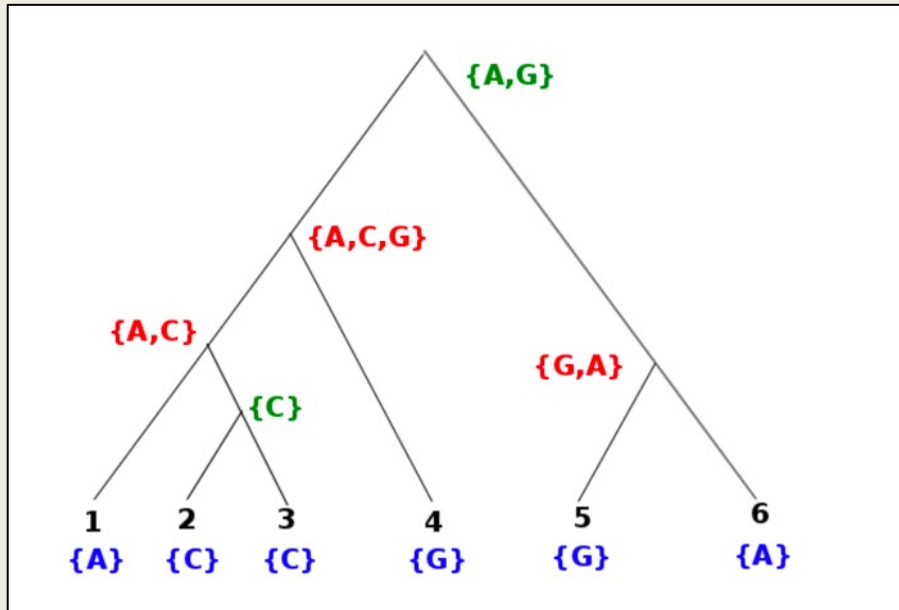
Fitch's algorithm

2. Traverse the tree from root to leaves. Assign state of the root randomly from its set. At each node, if the parent state is in that node's set, assign that. Otherwise, assign randomly.



Fitch's algorithm

The parsimony score is equal to the minimum number of mutations required

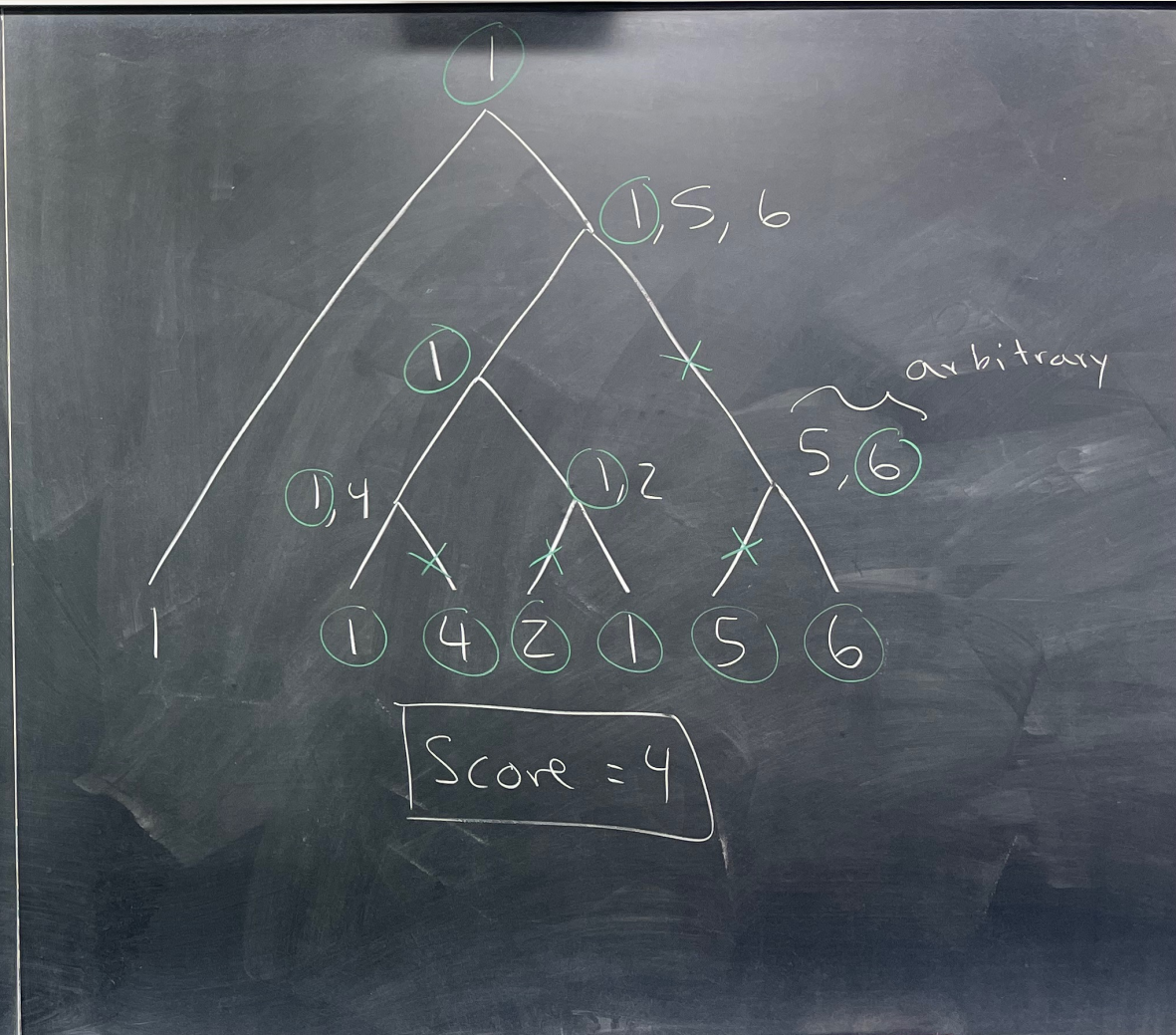
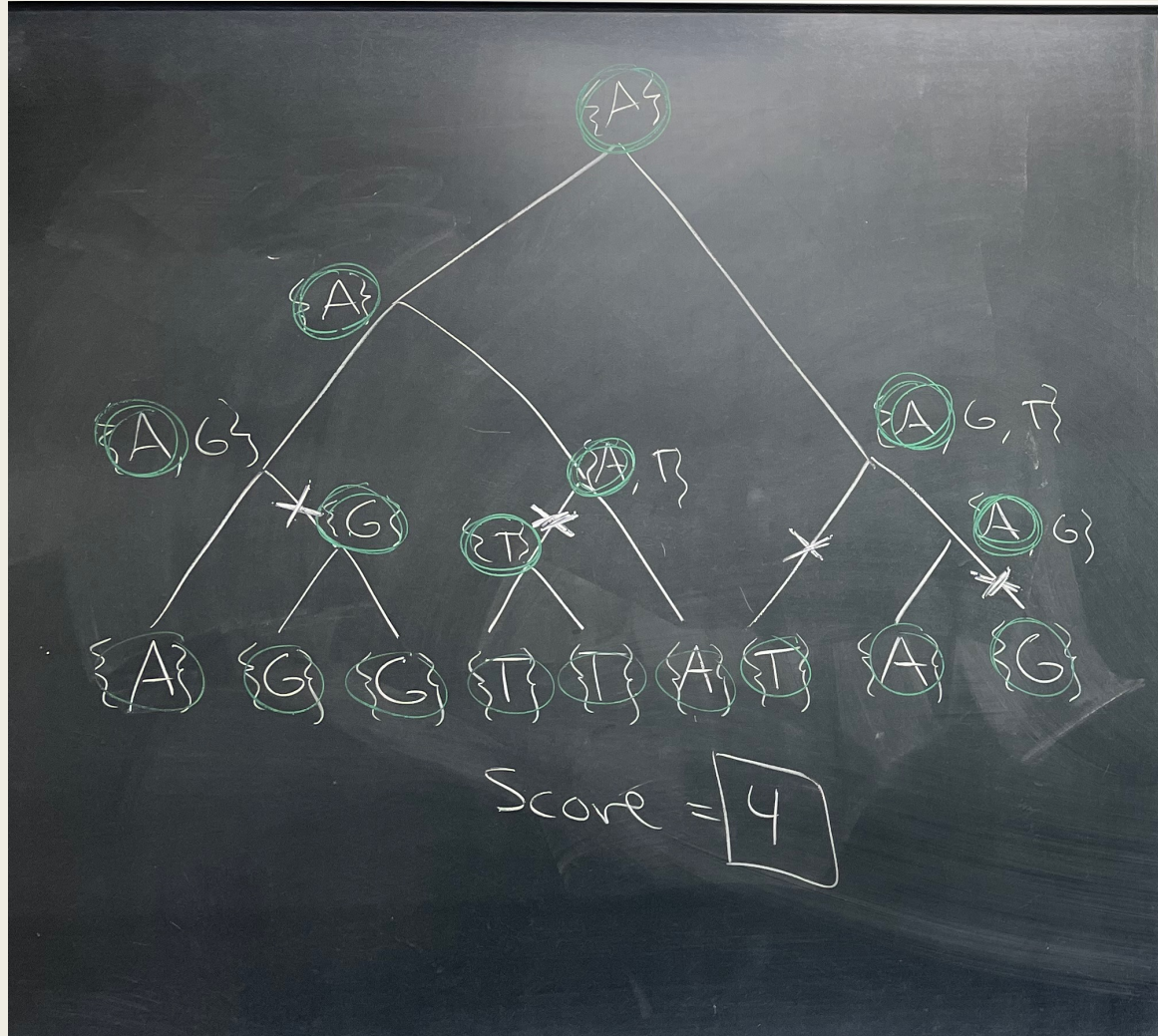


Fitch's algorithm handout (Handout 14, page 1)

1. Traverse the tree from leaves to root, at each step assign to the parent node the set that is the intersection of the set of characters in the child nodes if the intersection is non-empty, but the union if the intersection is empty.

2. Traverse the tree from root to leaves. Assign state of the root randomly from its set. At each node, if the parent state is in that node's set, assign that. Otherwise, assign randomly from the child node character set.

Fitch's algorithm handout (Handout 14, page 1)



Sankoff's algorithm (weighted parsimony)

Sankoff's algorithm

Fitch's algorithm assumes that every possible change is equally [un]likely

Sankoff's (1975) is a generalization that allows different changes to have different costs. Keeps track of the cost of being in each state at each node.

E.g.

	To	
σ	a	b
From a	0	2
b	1	0

a->b cost 2
b->a cost 1

Sankoff : weighted parsimony

input : rooted binary tree with
labeled leaves & cost
function Δ

i.e. $\Delta(a, b)$ is cost of
mutation $a \rightarrow b$

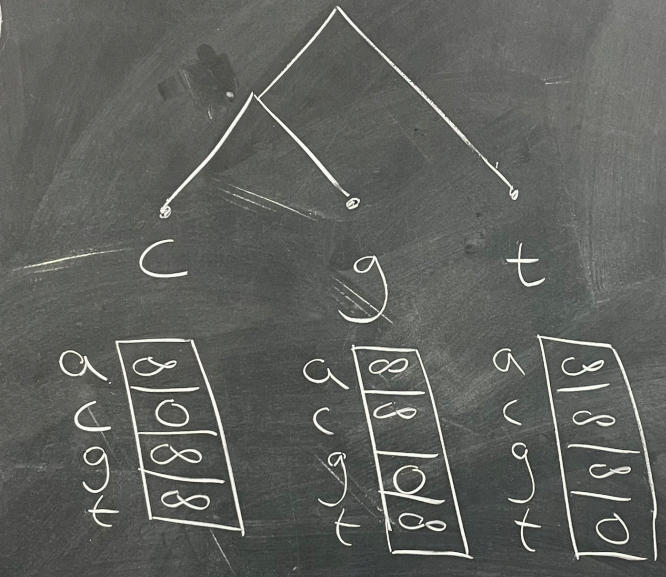
output : internal vertex labels
that minimize weighted
parsimony score

cost

parsimony
 tree with
 S & cost
 cost of
 b
 labels
 weighted
 one

initialization

leaf label characters assigned score
 of 0, other characters ∞



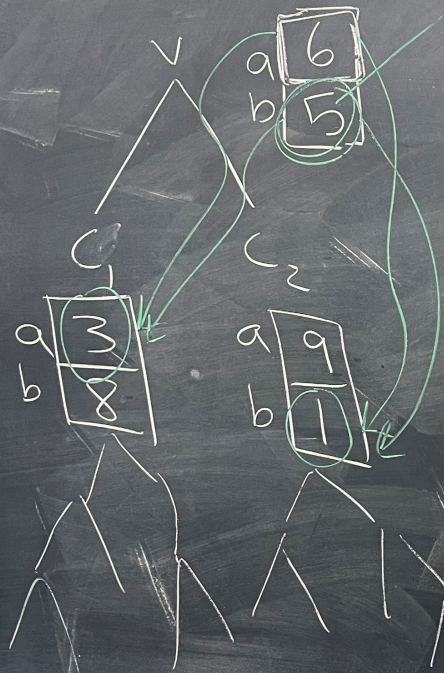
iterative step

let $A_v(x) =$

optimal cost
 of subtree rooted
 at v , with
 label x

[c_1 & c_2 children of v]

$$A_v(x) = \min \{ A_{c_1}(y) + \sigma(x, y) \} + \min \{ A_{c_2}(z) + \sigma(x, z) \}$$



choose min at root

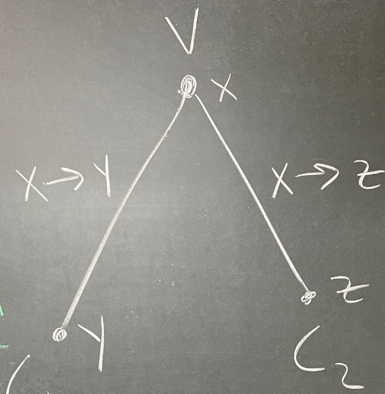
$$A_v(a) = \min \{ A_{c_1}(a) + \sigma(a, a), A_{c_1}(b) + \sigma(a, b) \} + \min \{ A_{c_2}(a) + \sigma(a, a), A_{c_2}(b) + \sigma(a, b) \}$$

$$= \min \{ 3 + 0, 8 + 2 \} + \min \{ 9 + 0, 1 + 2 \}$$

$$= 3 + 3 = 6$$

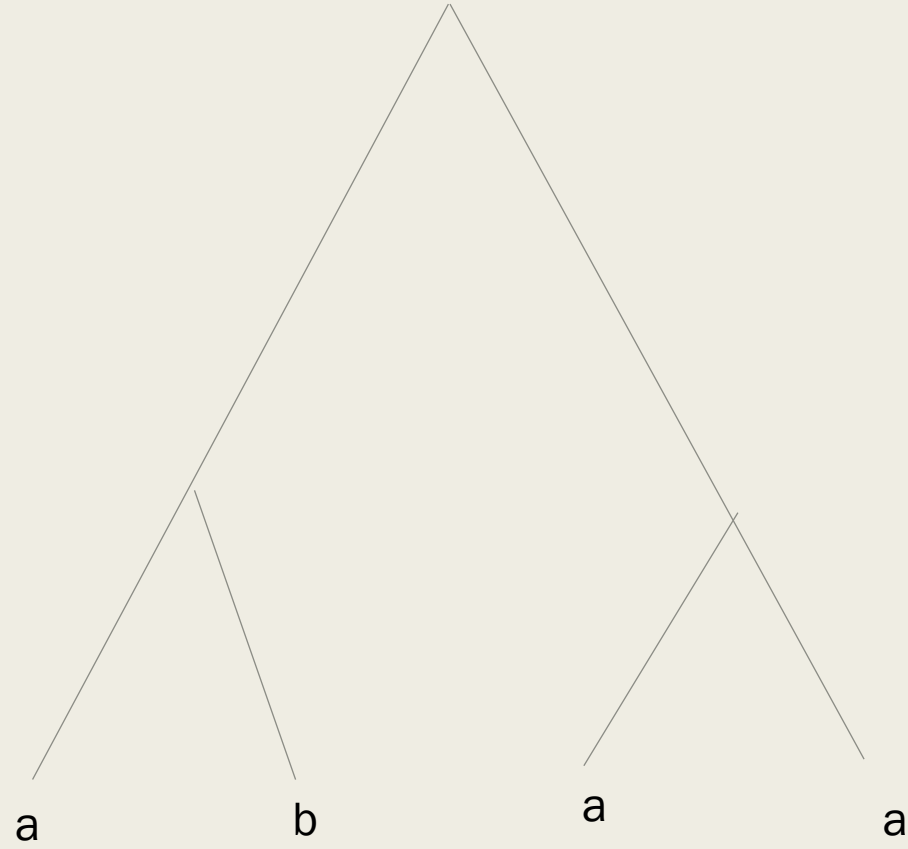
$$A_v(b) = 4 + 1 = 5$$

exercise!



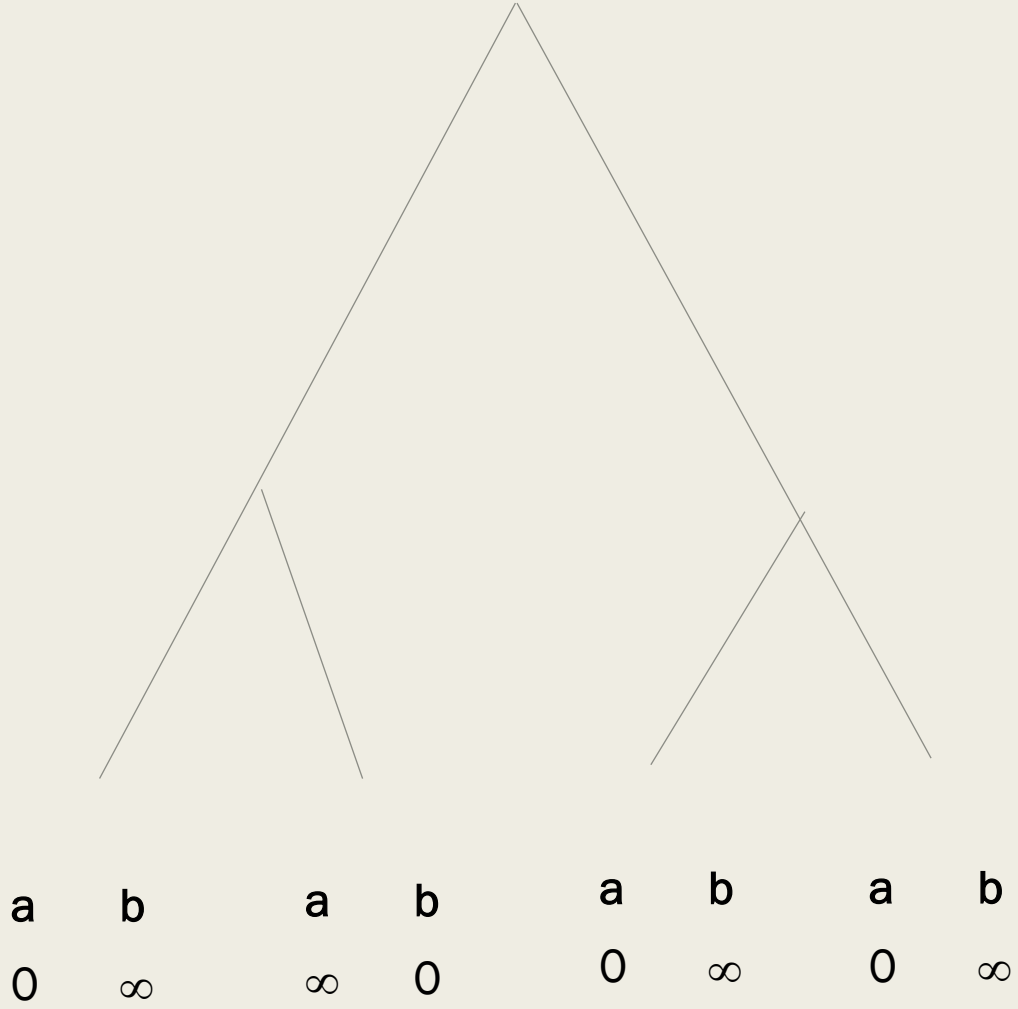
child 1

child 2

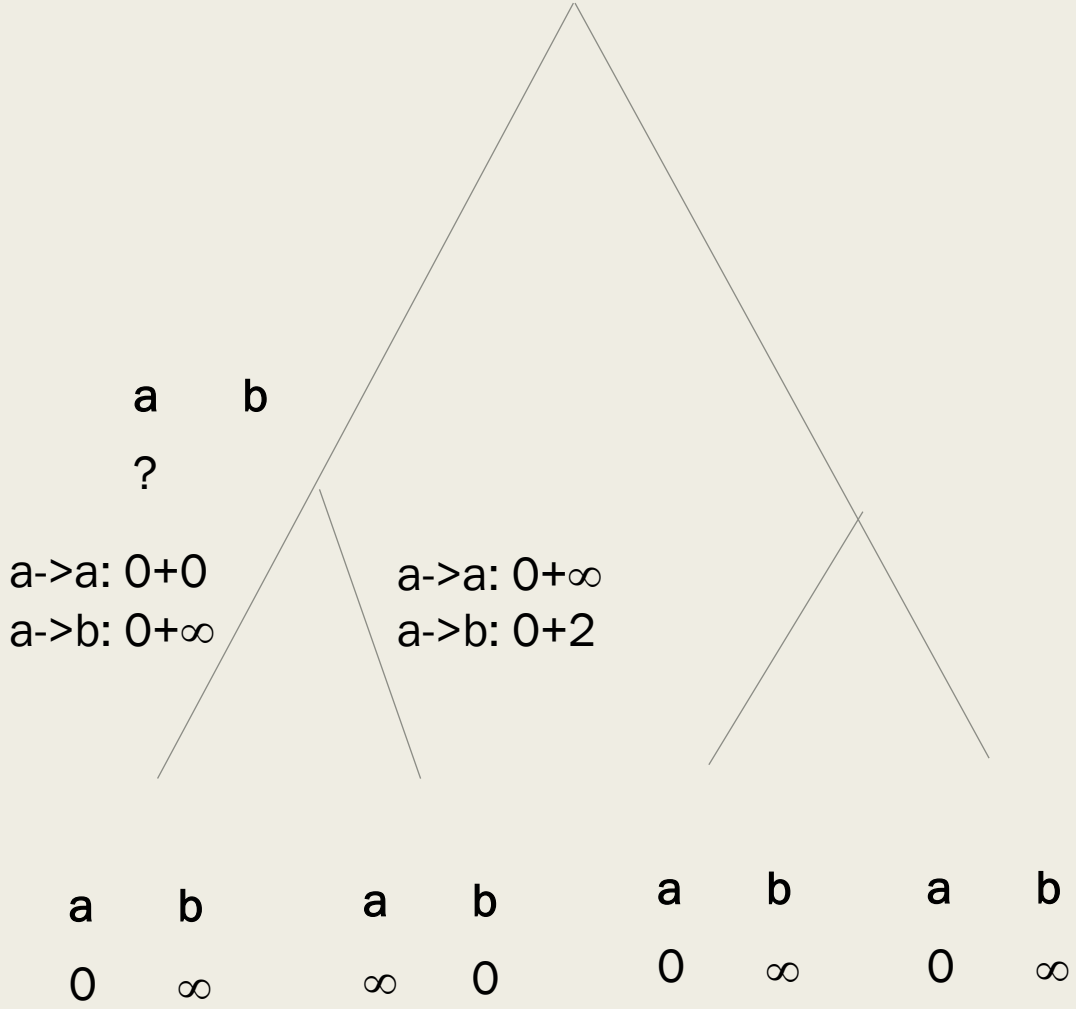


σ	a	b
a	0	2
b	1	0

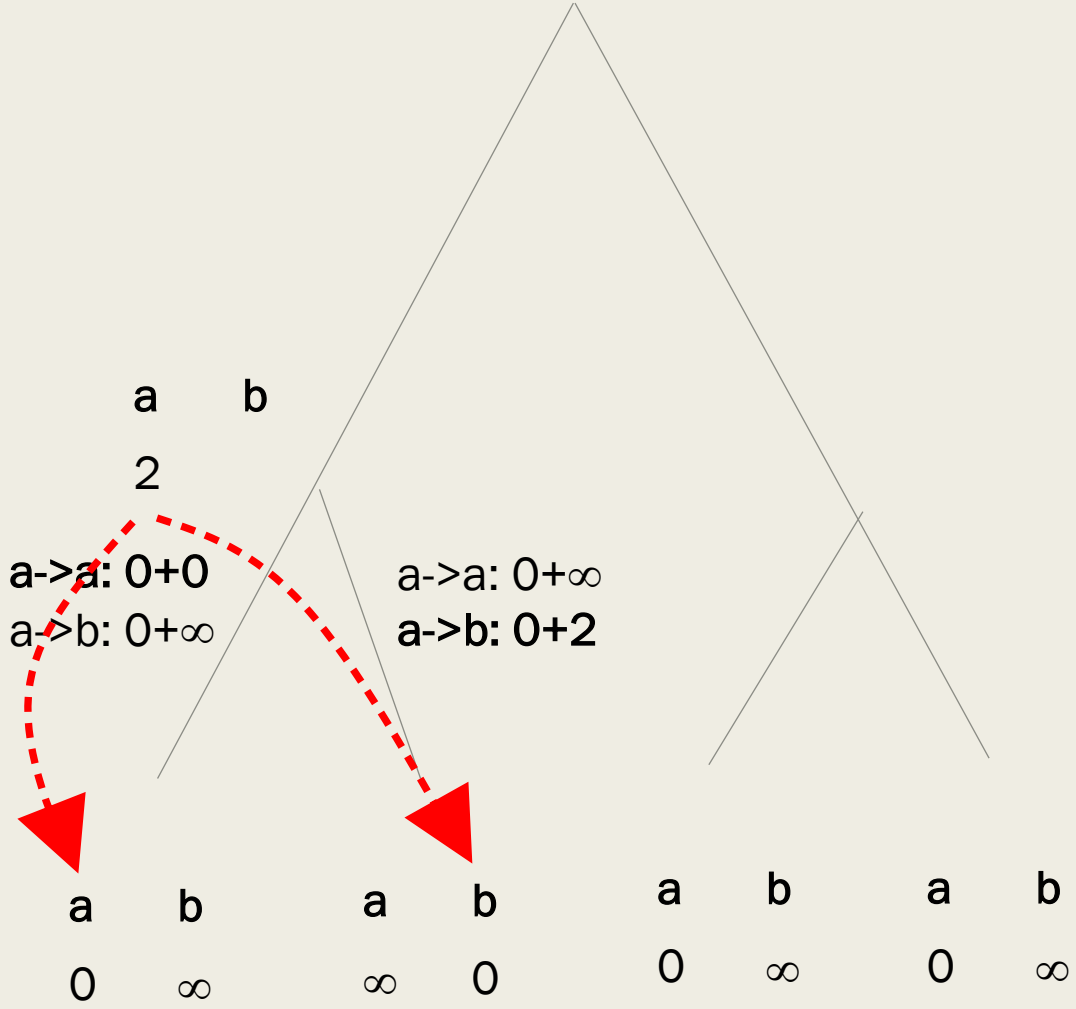
σ	a	b
a	0	2
b	1	0



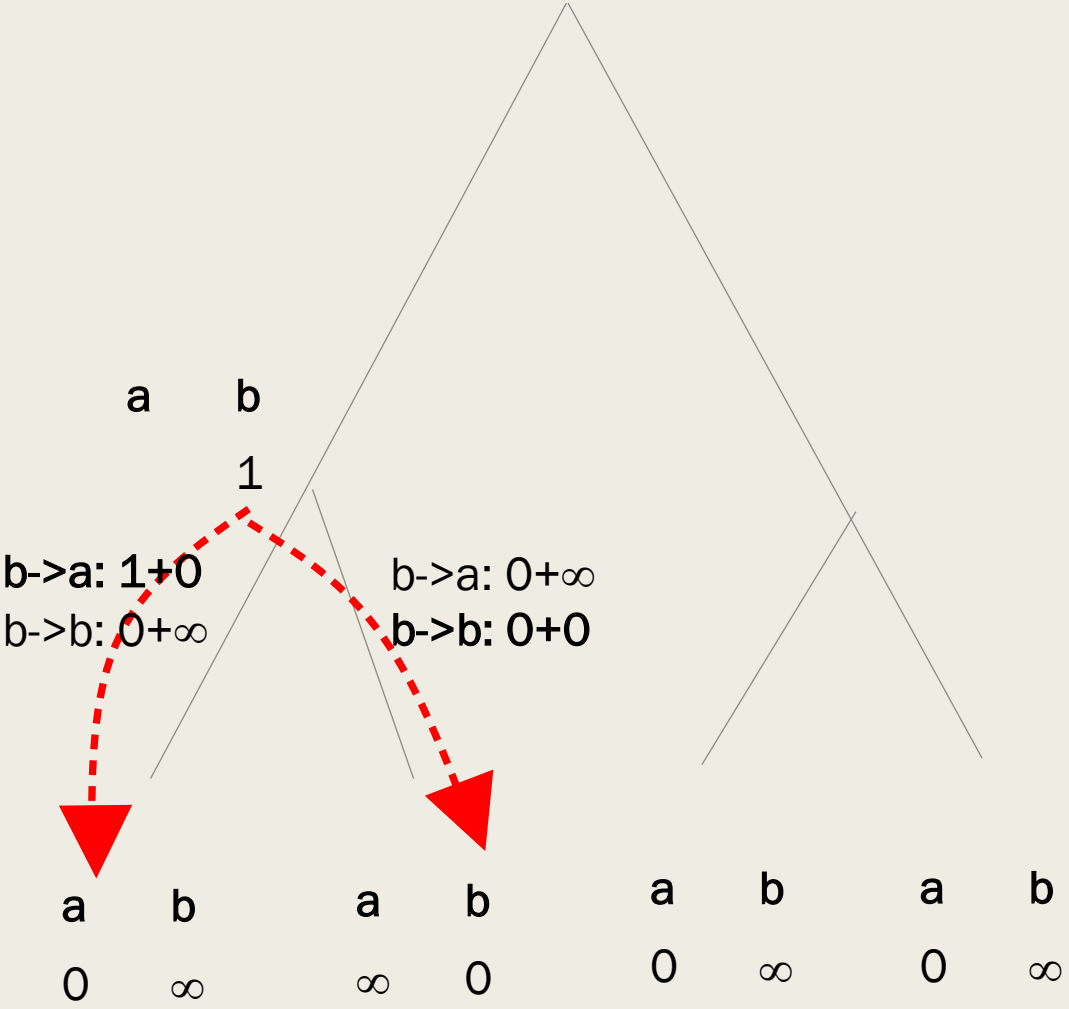
σ	a	b
a	0	2
b	1	0



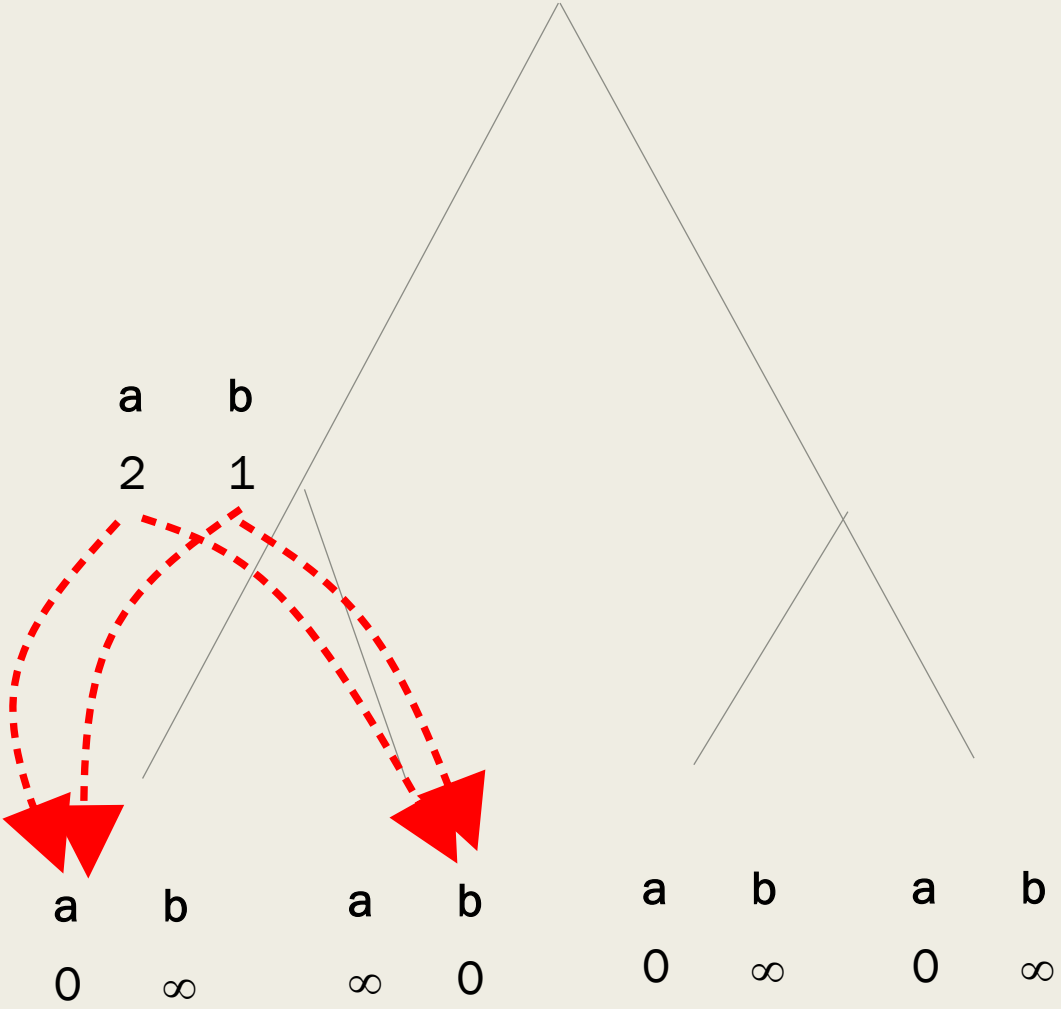
σ	a	b
a	0	2
b	1	0



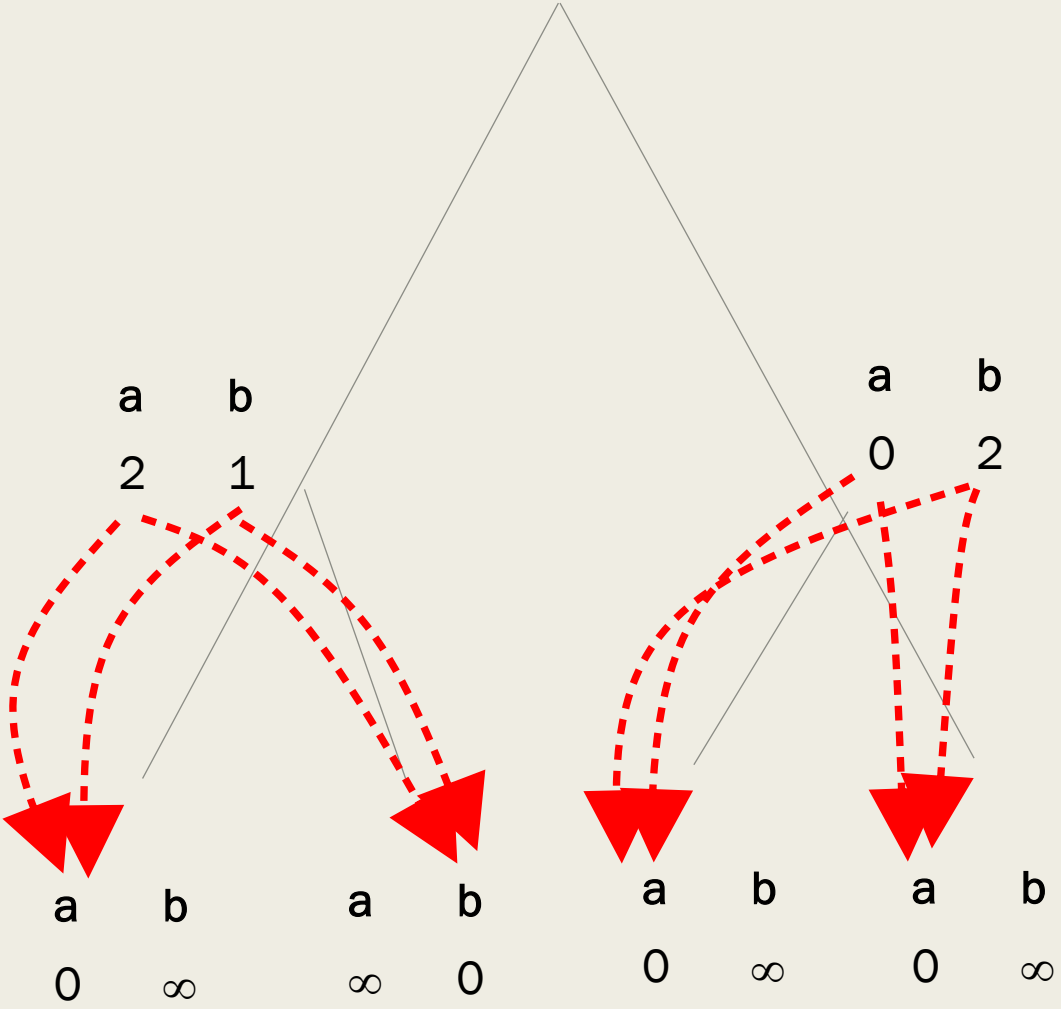
σ	a	b
a	0	2
b	1	0

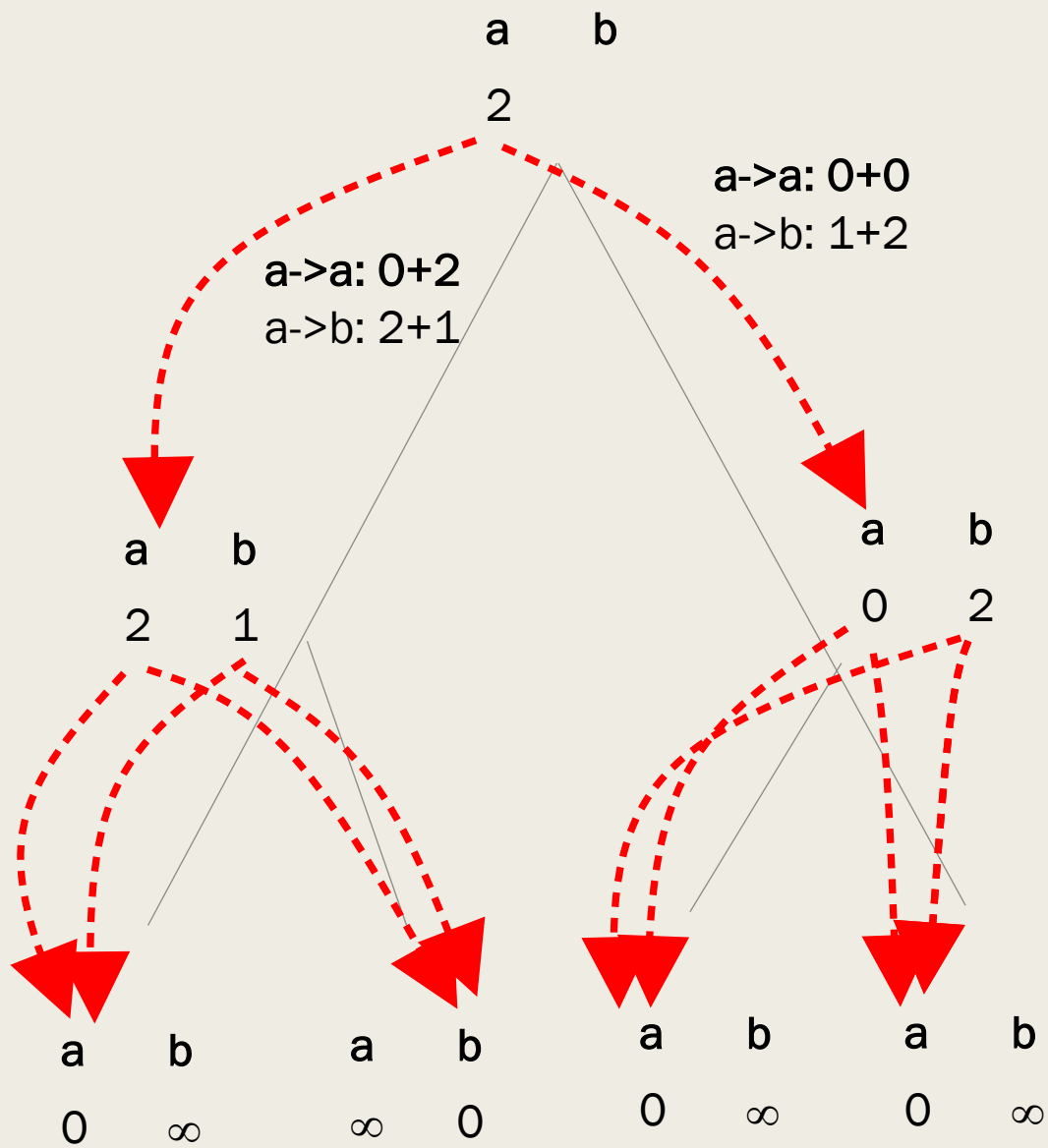


σ	a	b
a	0	2
b	1	0

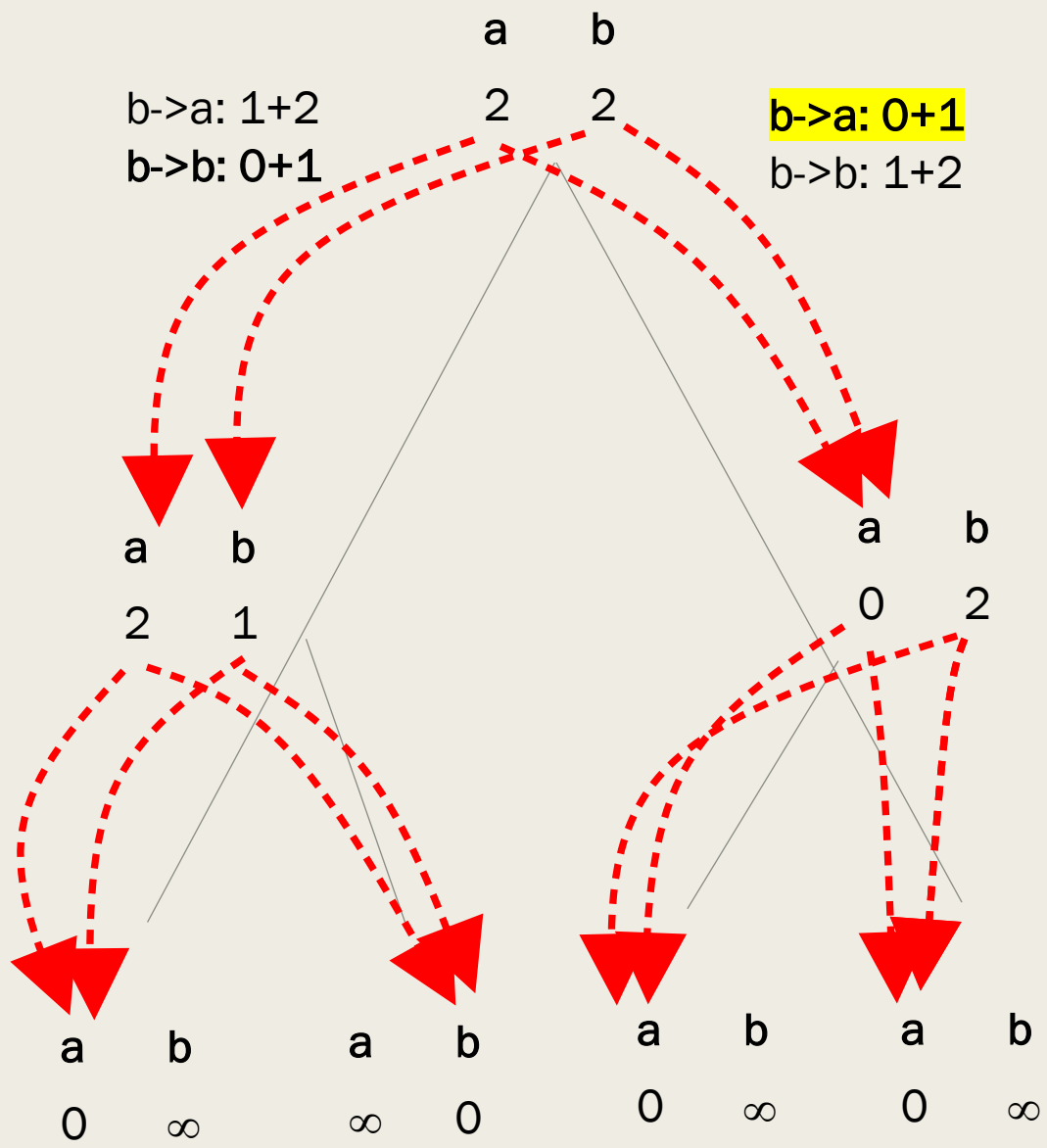


σ	a	b
a	0	2
b	1	0





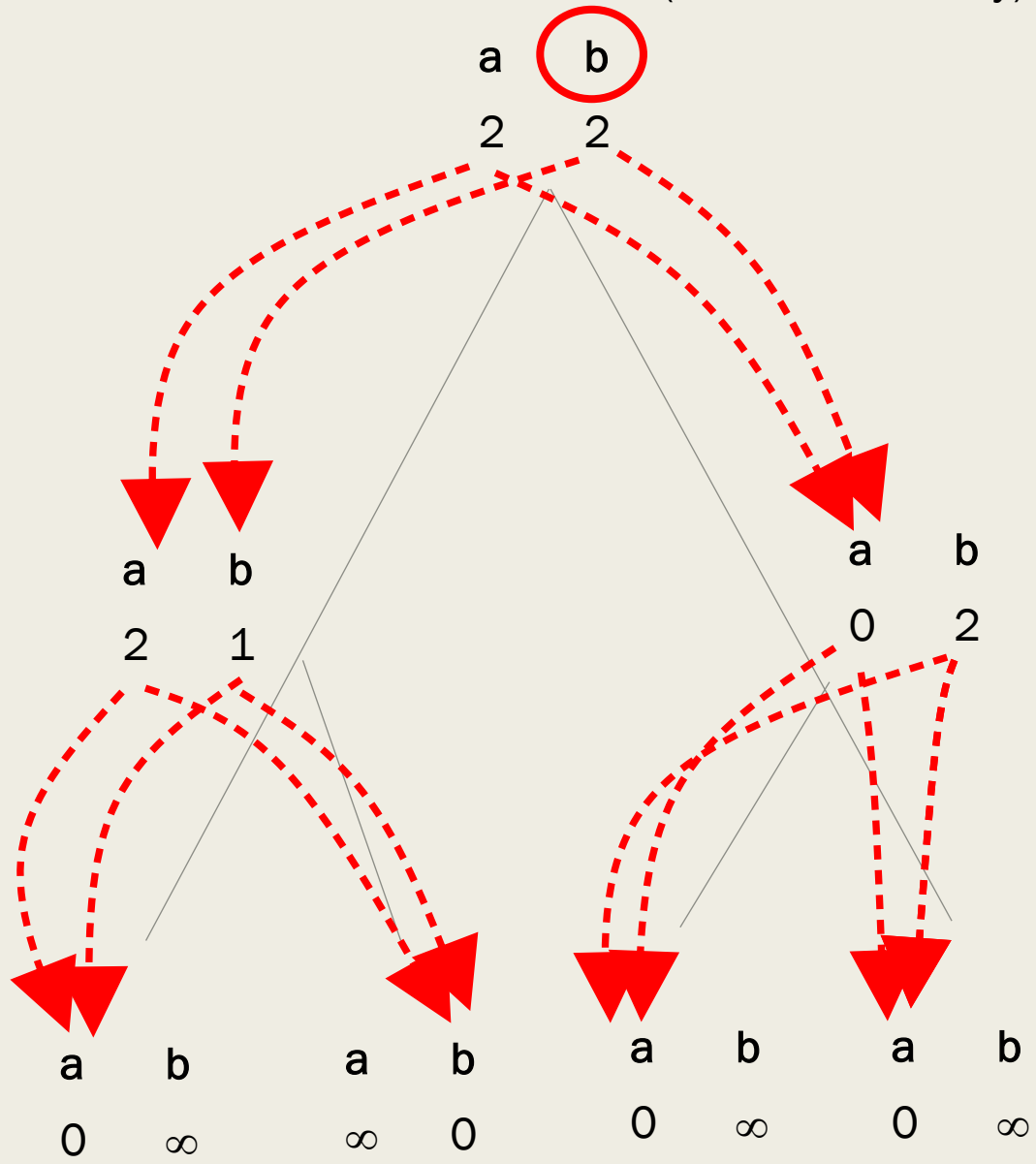
σ	a	b
a	0	2
b	1	0



σ	a	b
a	0	2
b	1	0

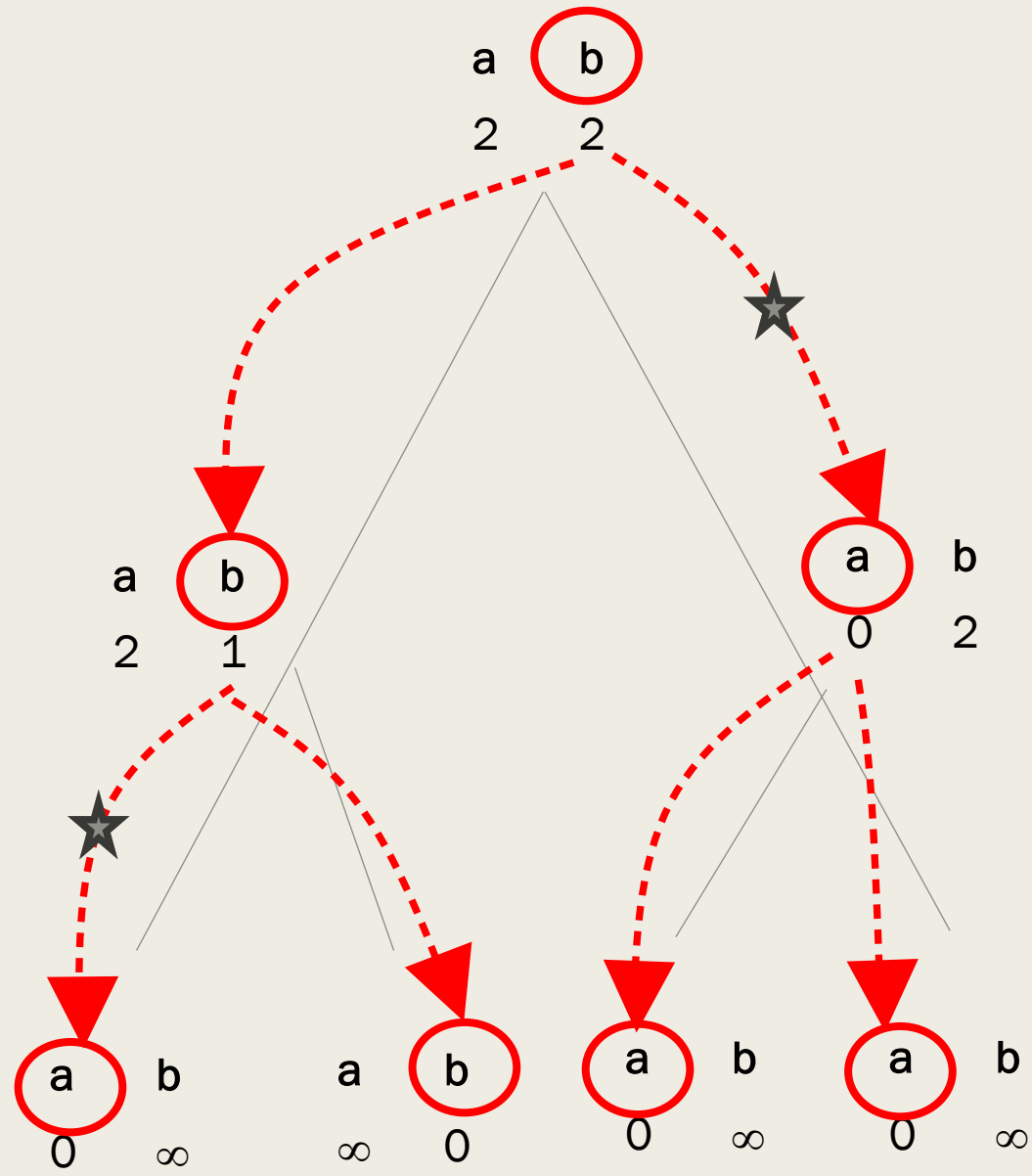
Corrected!

(choose arbitrarily)



σ	a	b
a	0	2
b	1	0

Corrected!



σ	a	b
a	0	2
b	1	0

★ mutation

Corrected!

Sankoff's algorithm handout

Initialization: Let $A_v(x)$ be the minimum parsimony score of assigning character x to vertex v . To begin $A_{\text{leaf}}(x) = 0$ if the leaf is assigned character x , and ∞ otherwise. This prevents us from ever tracing back to a non-assigned leaf label.

Bottom-up recursive step: Let c_1 and c_2 be the two children of vertex v . For all x in our character state set, let

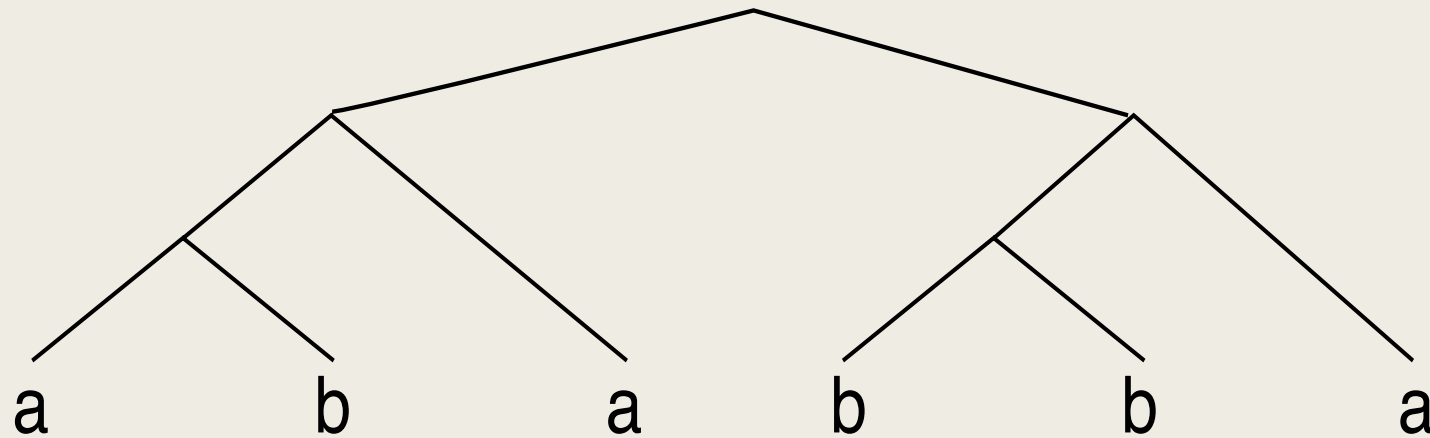
$$A_v(x) = \min_y \{A_{c_1}(y) + \sigma(x, y)\} + \min_z \{A_{c_2}(z) + \sigma(x, z)\}.$$

Keep track of a back-pointer to the minimum y and z .

Top-down traceback: Choose root state x such that $A_{\text{root}}(x)$ is the minimum. Follow back-pointers to find the assigned state of every internal vertex.

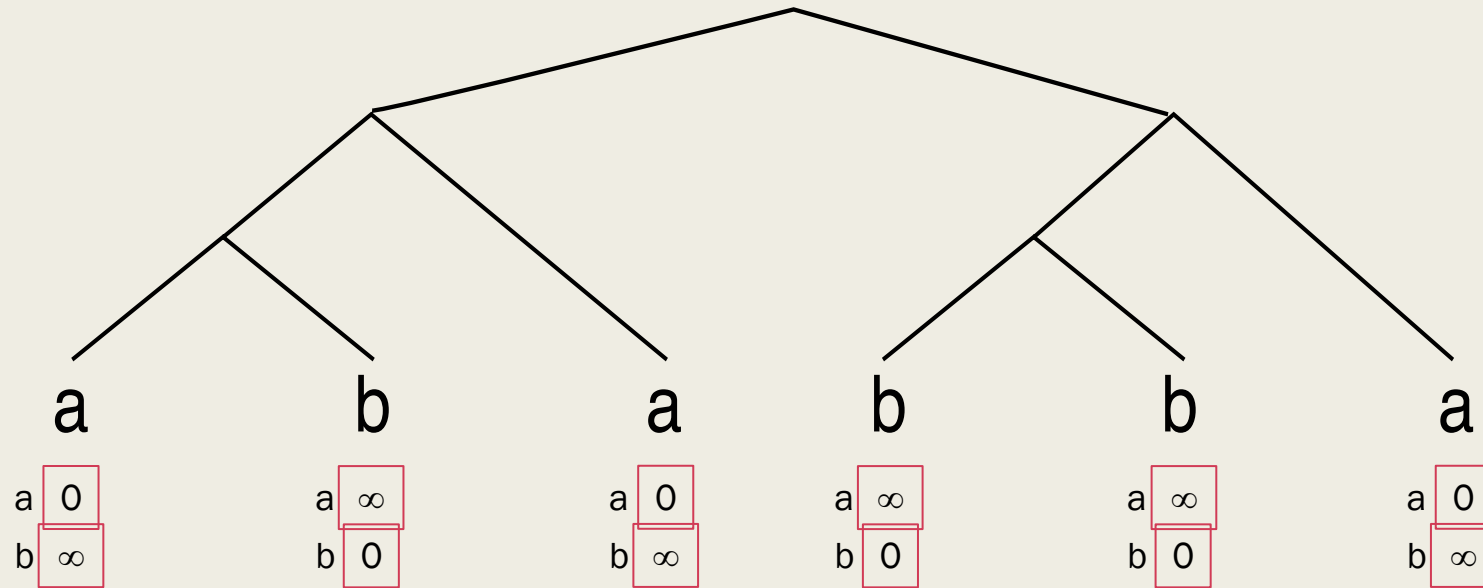
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



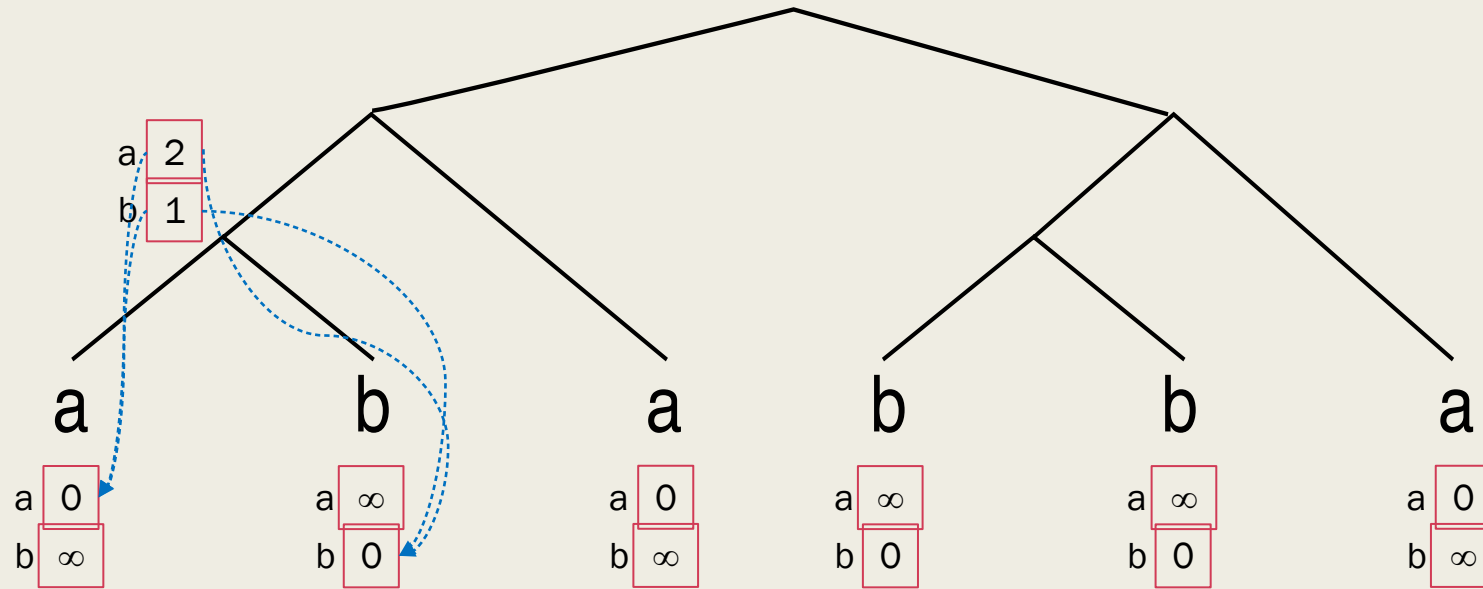
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



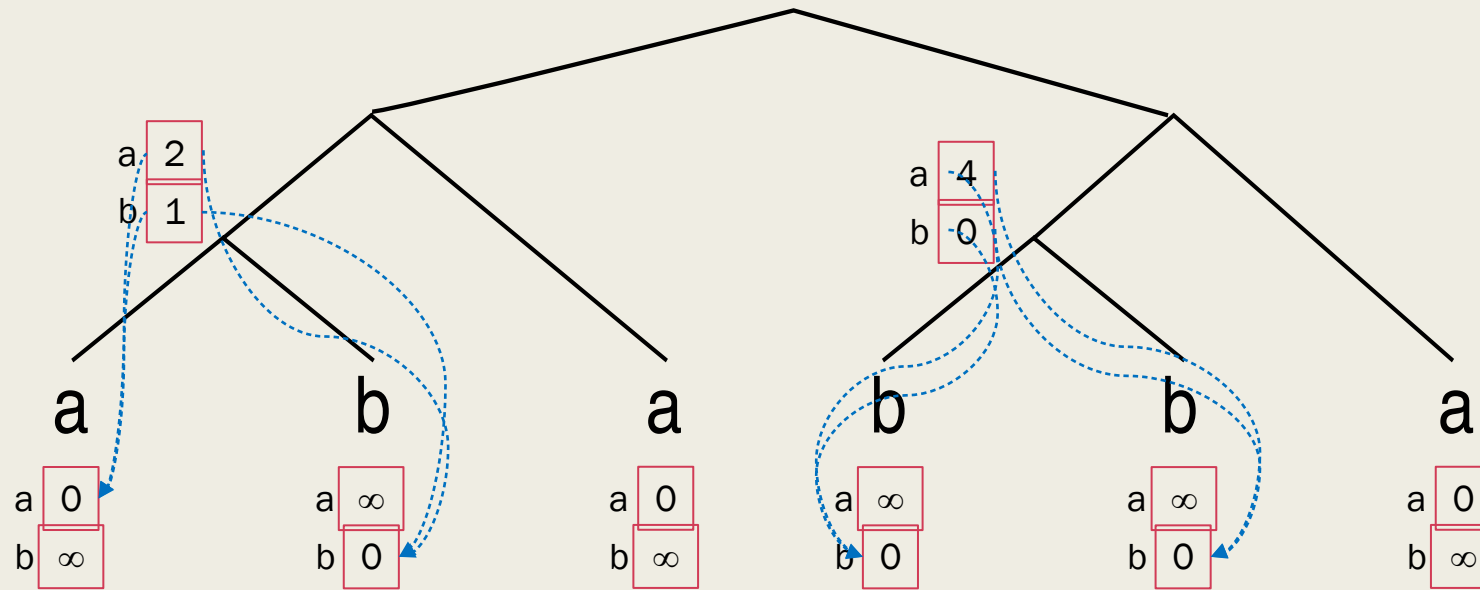
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



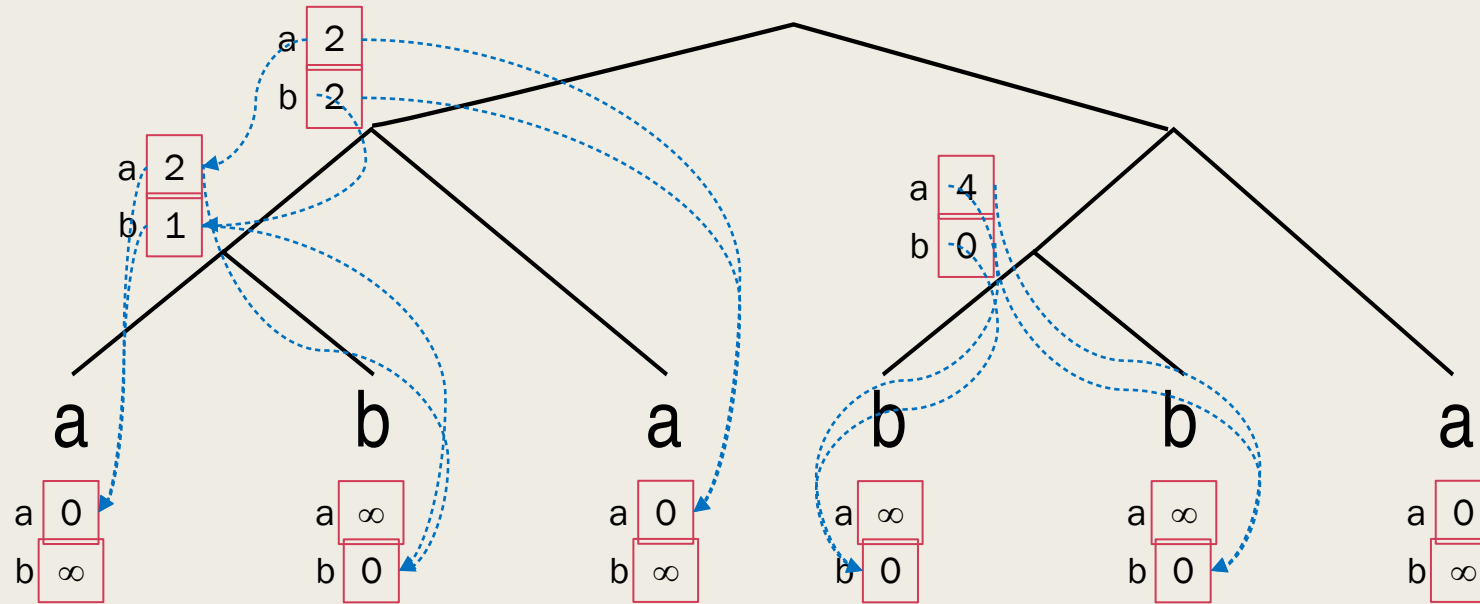
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



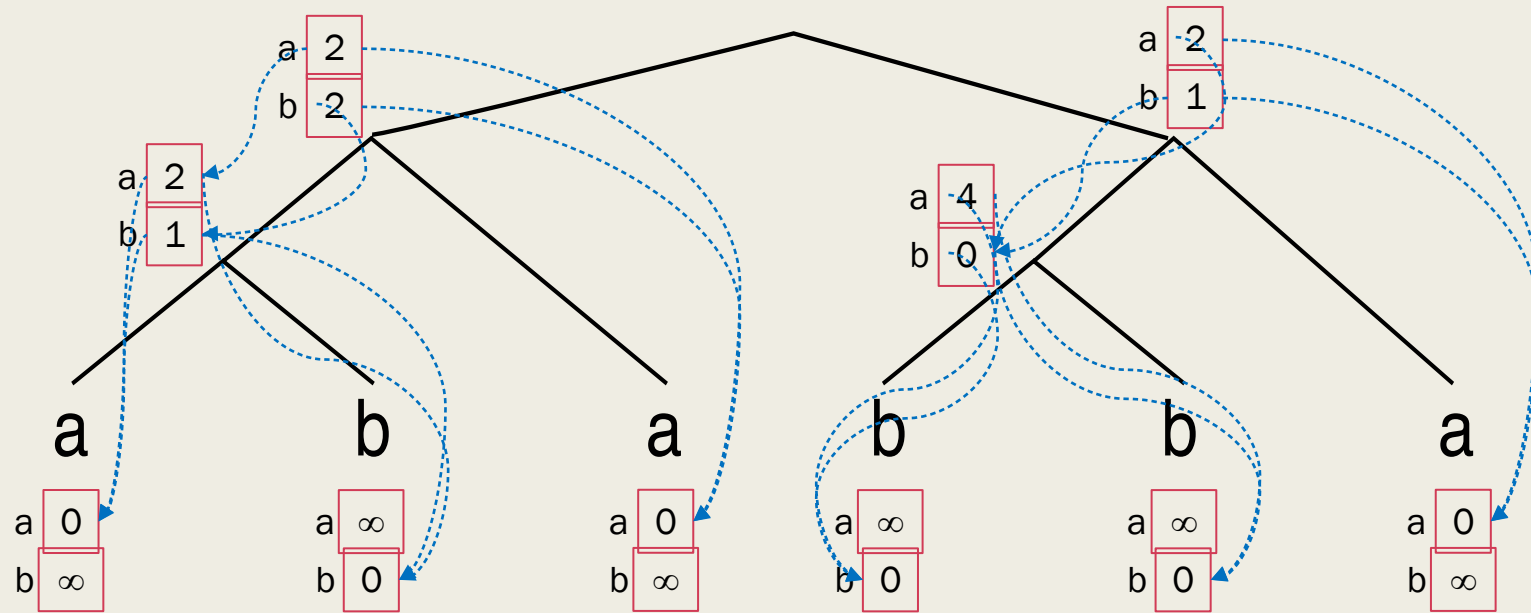
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



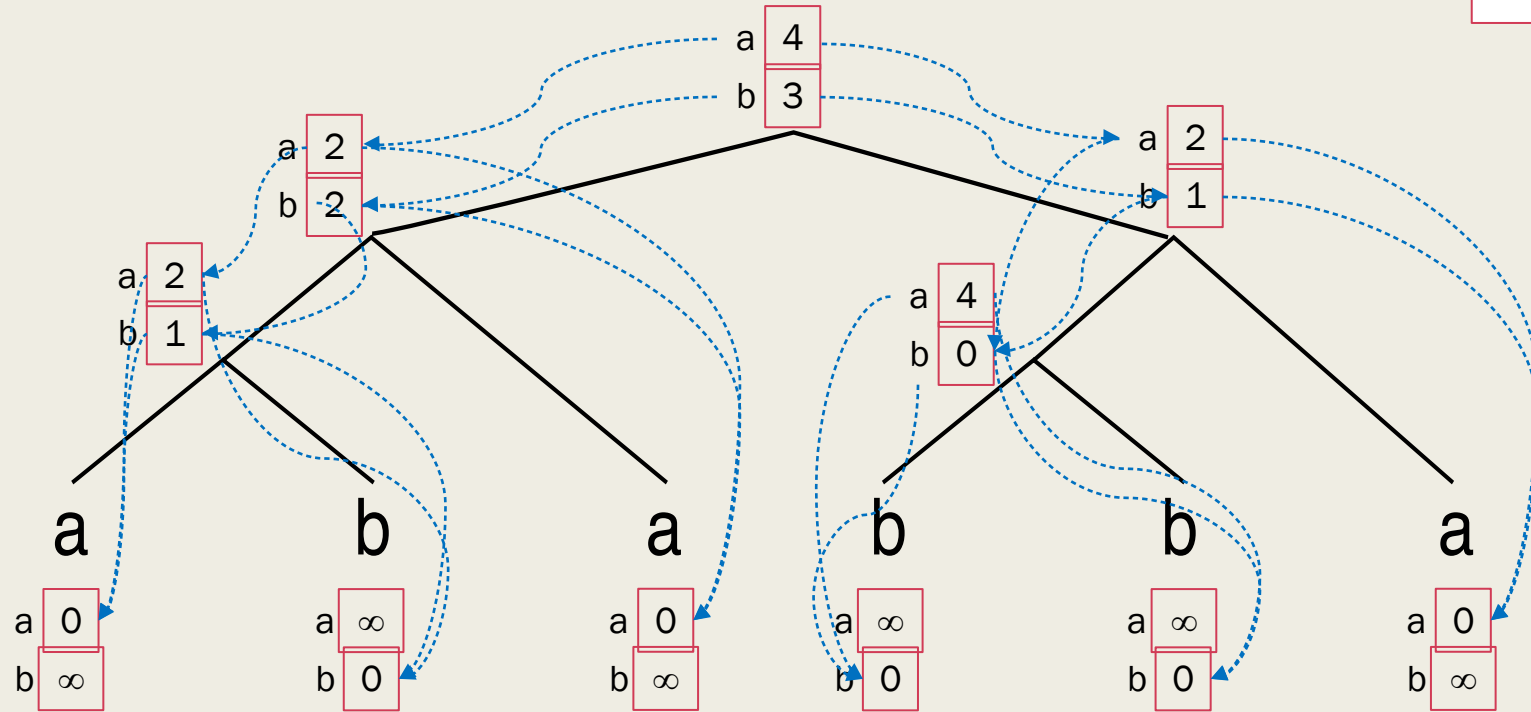
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



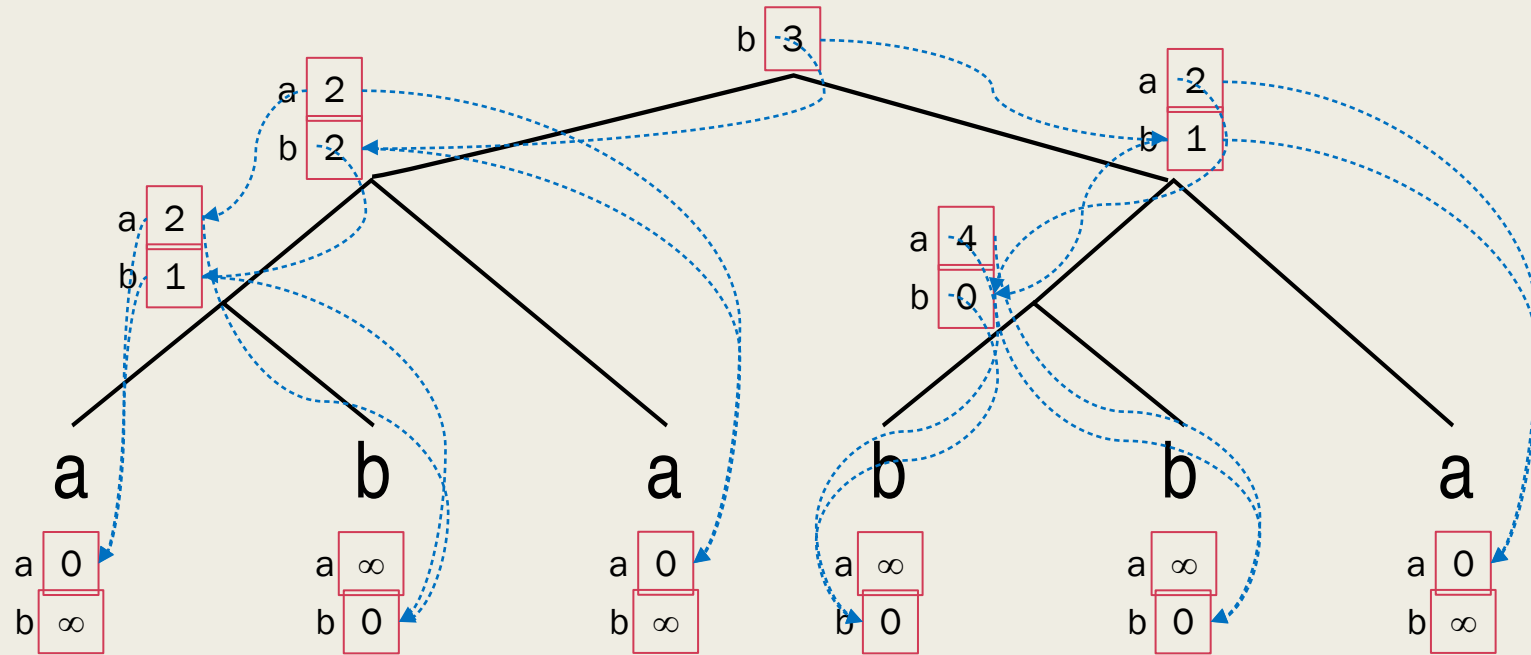
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



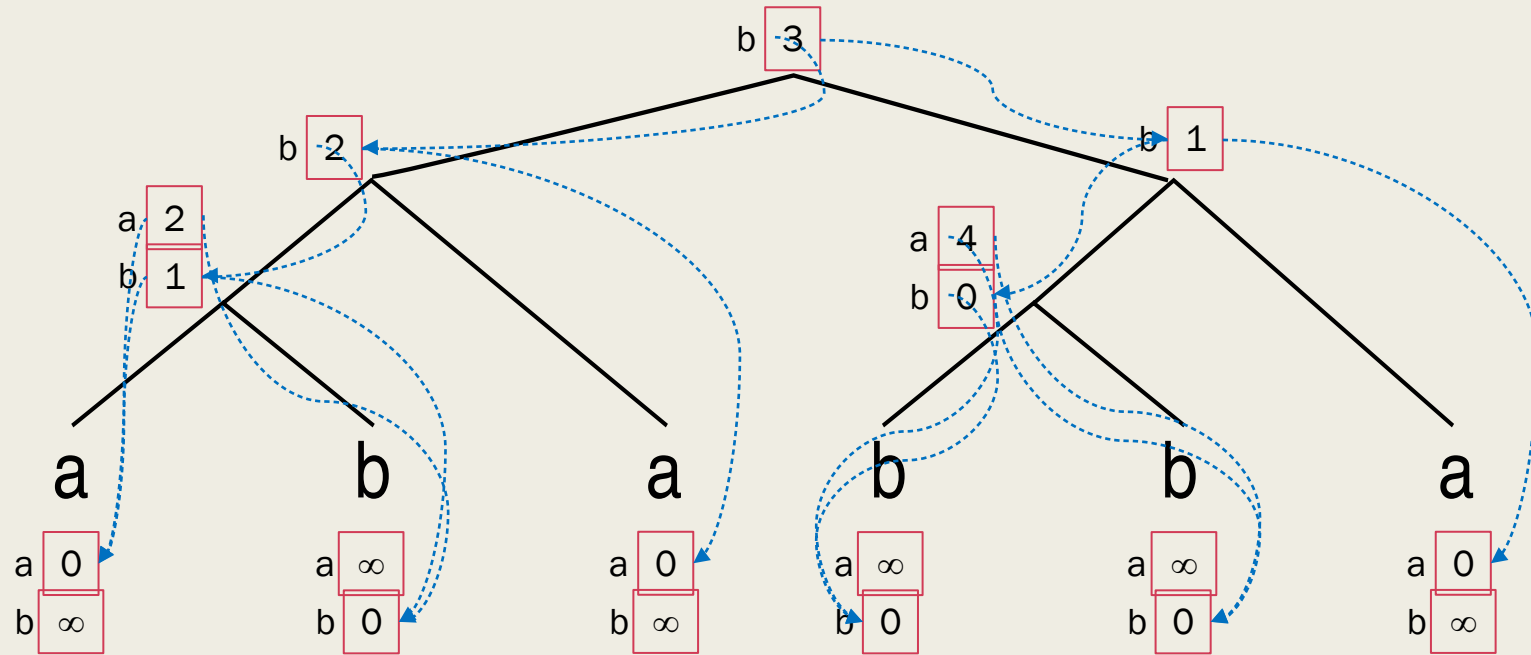
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



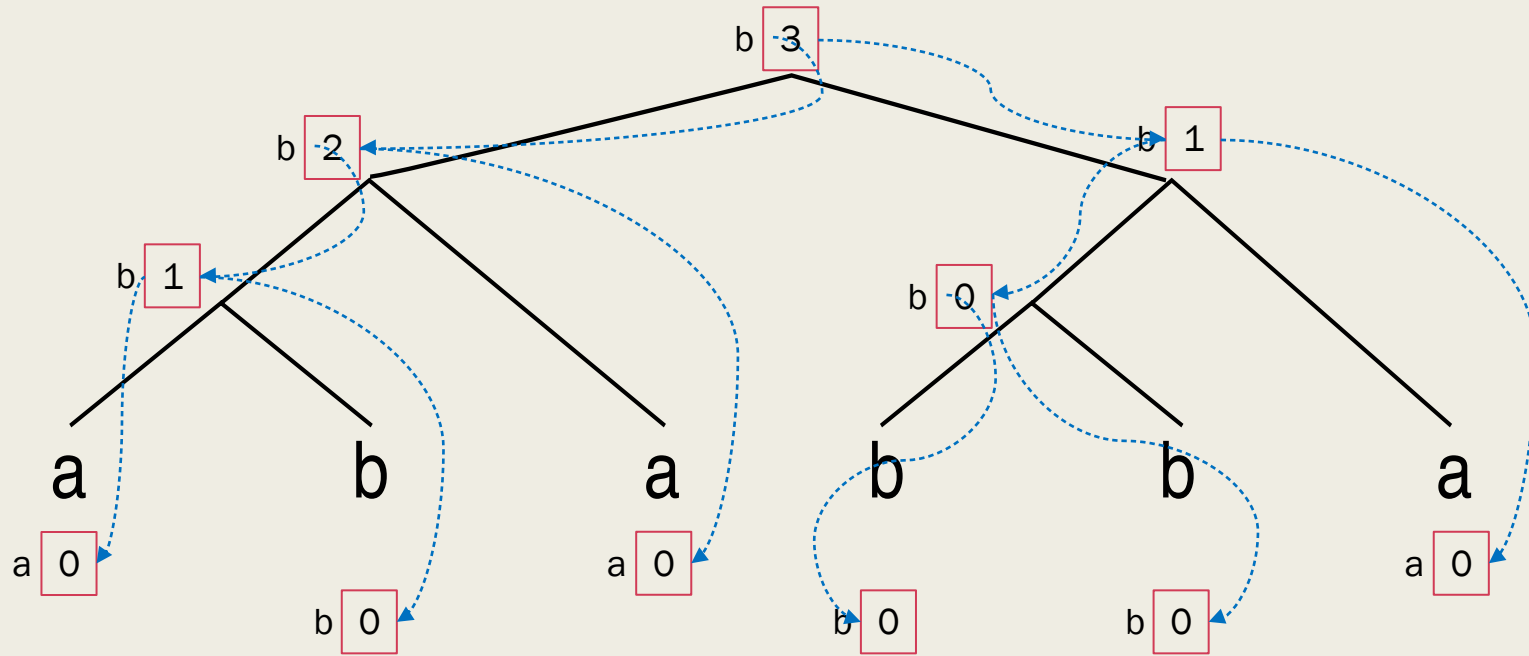
Sankoff algorithm

σ	a	b
a	0	2
b	1	0



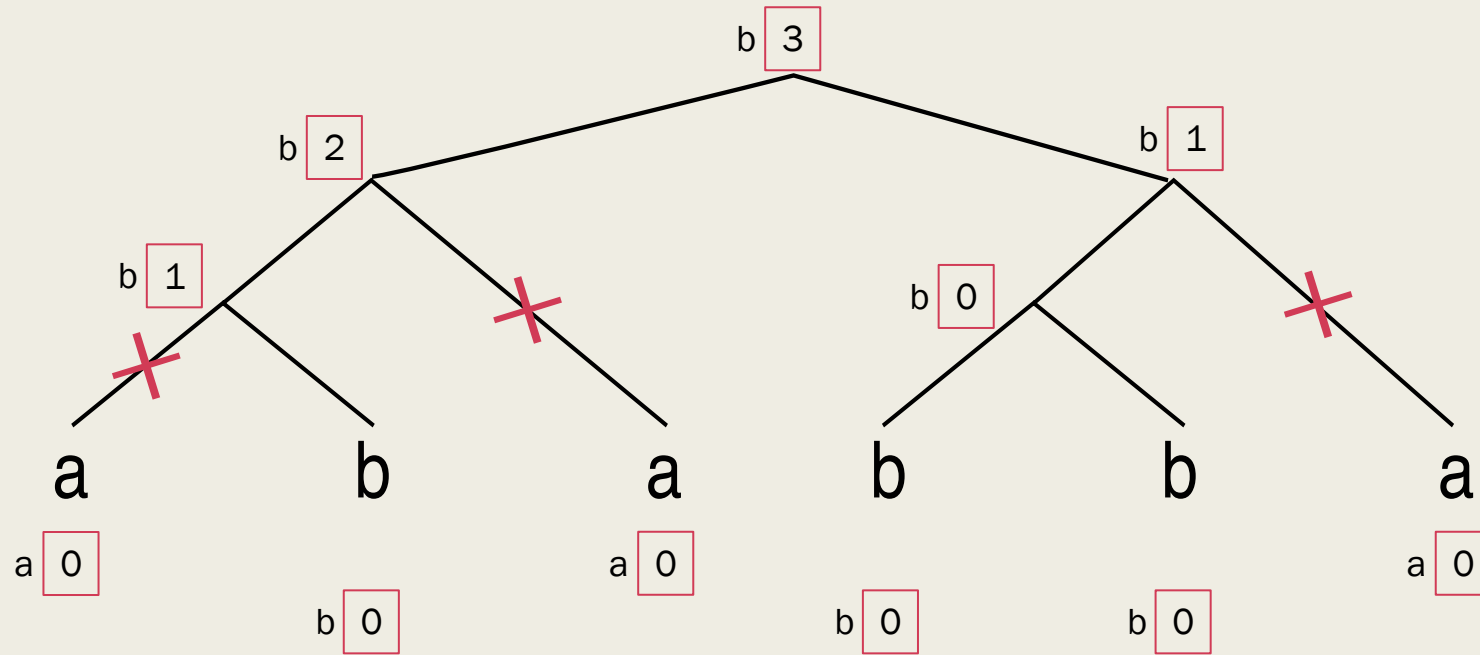
Sankoff algorithm

σ	a	b
a	0	2
b	1	0

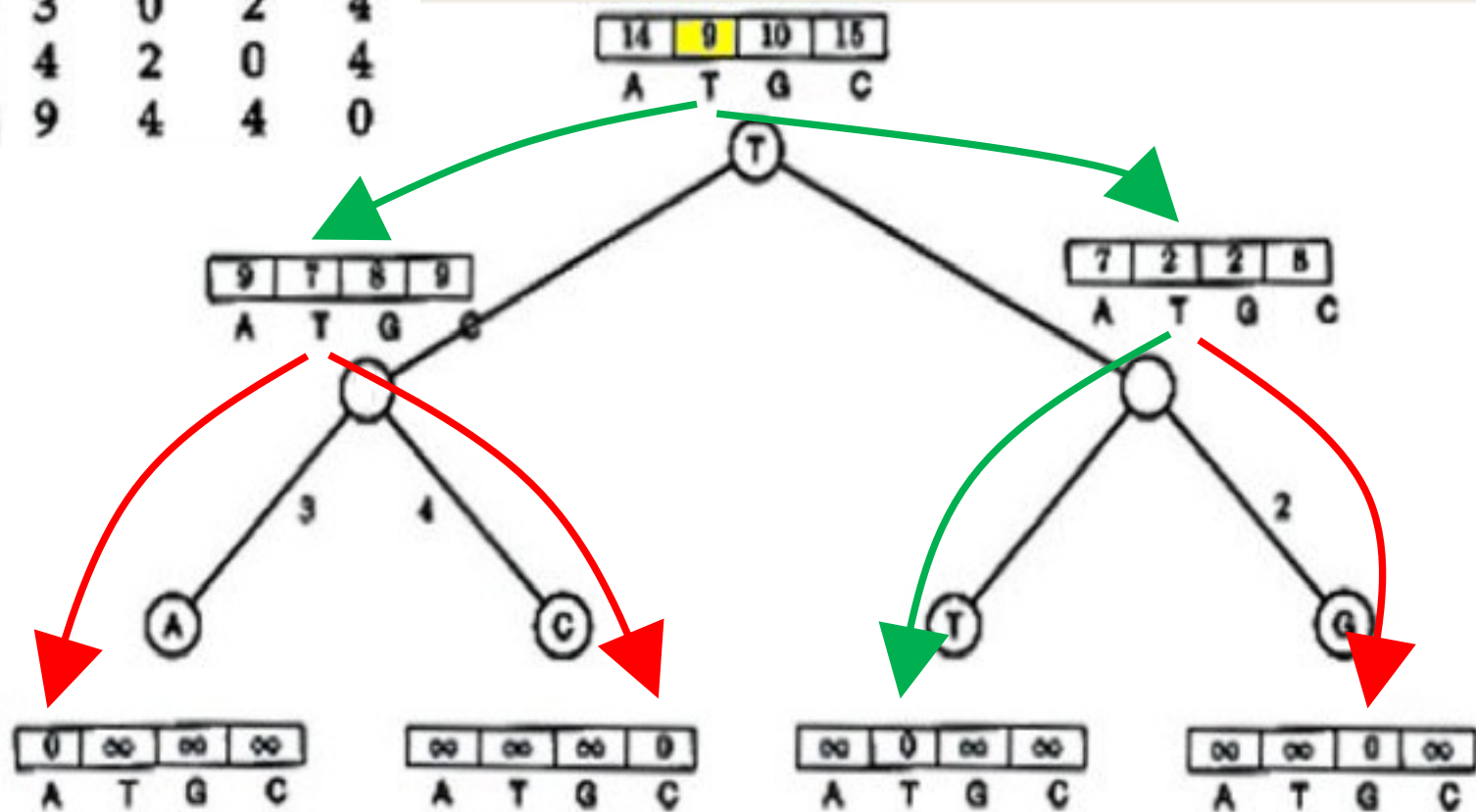


Sankoff algorithm

σ	a	b
a	0	2
b	1	0



δ	A	T	G	C
A	0	3	4	9
T	3	0	2	4
G	4	2	0	4
C	9	4	4	0



Runtime

Suppose there are N leaves and K states

What is the complexity of Fitch's algorithm?

What is the complexity of Sankoff's algorithm?