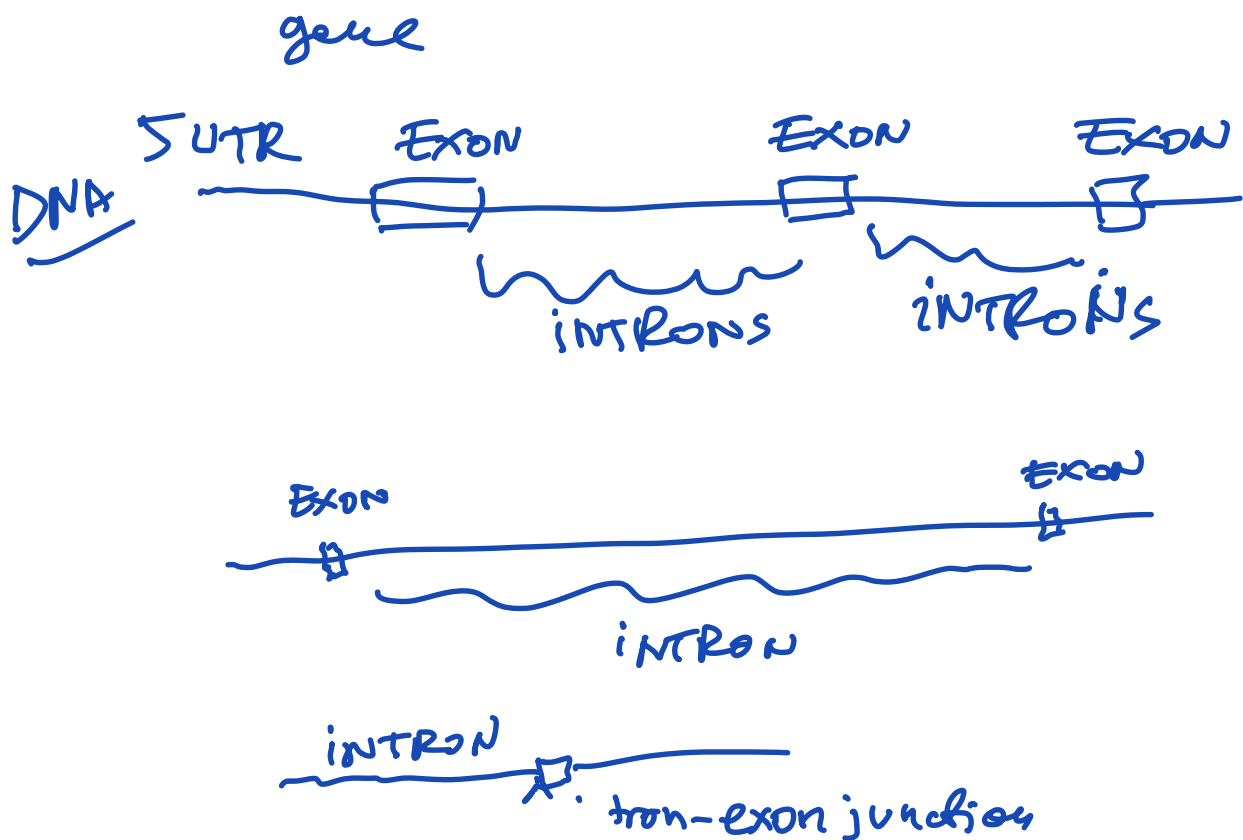


MARKOV CHAINS & HIDDEN MARKOV MODELS

SORIN ISTRAIL

we can find out genes on
the genomes



$\frac{1}{2}$ in ○ Head
Tail
SOME EXPERIMENTS can hear
 you but cannot see her
 $H, T, T, T, H, T, \dots$
 is the coin fair? 50-50 Head-Tail

coin H 75%
 T 25%

Can we infer the bias of the coin?
 How do we know?

① $H, T, H, H, T, T, T, H, T, H, H$

② $H, \underline{T, T, T, T, T}, H, \underline{T, T, T}, H$

50% = unbiased

①

1	2	3	4	5	6	7
(1,0)	(1,1)	(2,1)	(3,1)	(3,2)	(3,3)	(3,4)

②

↑	↑					
H	T					
(1,0)	(1,1)	(1,2)	(1,3)	(1,4)	(1,5)	...

The T of the coin (2) is biased
we cannot do this inference
when we have a long seq.
of tosses.

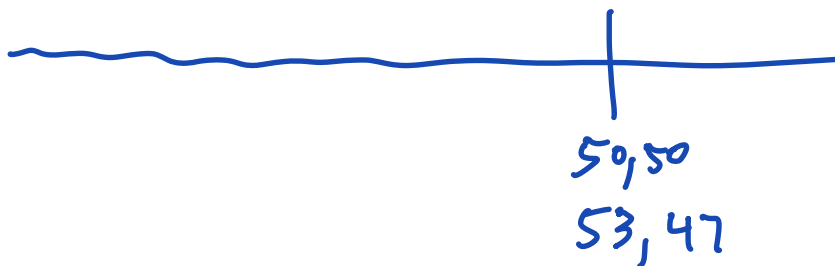
We can estimate what
the bias is: $75-25$ or
 $80-20$
 $55-45$

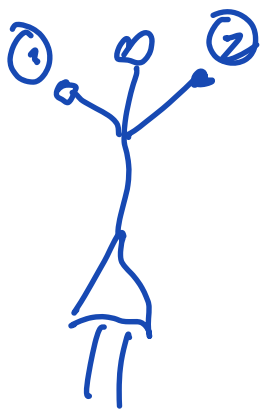
Dishonest Casino Problem

A random sequence:

H, T, H, H, T, T, T, ...

at any point you leave
to have roughly 50-50
H-T





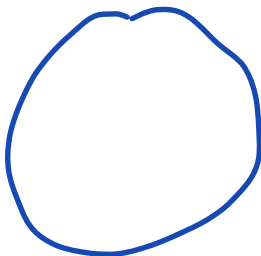
two coins coin: H or T
 we coin #3 — H Flips ①
 T Flips ②

H, T, H, H, T
 ① ① ② ① ②

- ① P_H^1, P_T^1 $P_H^1 + P_T^1 = 1$
- ② P_H^2, P_T^2 $P_H^2 + P_T^2 = 1$
- ③ P_H^3, P_T^3 $P_H^3 + P_T^3 = 1$

Can we find out: all these 6
 numbers $P_H^1, P_T^1, P_H^2, P_T^2, P_H^3, P_T^3$
 ?

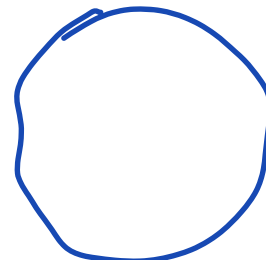
HMM we can solve this pb.



Fair



#3



biased

MARKOV CHAINS

$N = \#$ of states

$$S = \{s_1, \dots, s_N\}$$

$A =$ the state transition
probability matrix

A

s_1		
s_i	$a(i,j)$	
s_N		

$a(i,j) = \text{prob.}$
of transitioning
from state
 s_i to state s_j

Providence

$s_1 = \text{Sunny}$
 $s_2 = \text{Rainy}$

A Markov Model

$N = 2$

$$S = \{s_1, s_2\}$$

day by day

	s_1	s_2
s_1	0.8	0.2
s_2	0.9	0.1

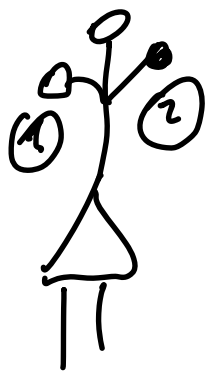
States: are observable

later: Hidden States

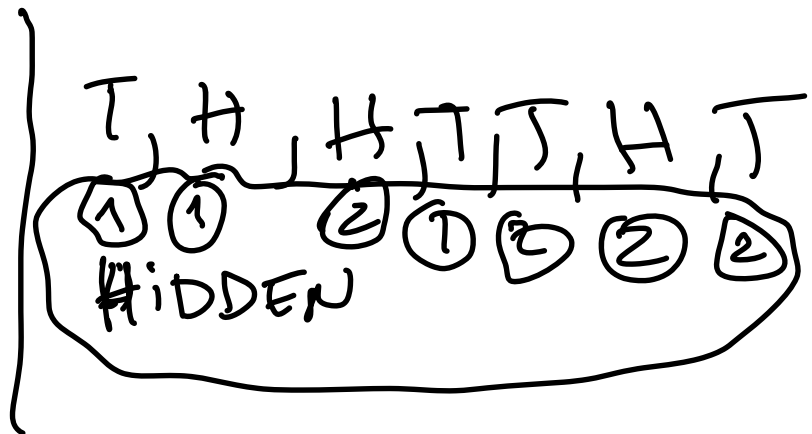
"decode": reveal the hidden states

Hidden Markov Models (HMM)

Back to the example



coin
#3



$M=2$

alphabet $V = \{T, H\}$
sequence of symbols or

sequence of observations

DEF (HMM)

(N, M, A, B, π)

1) $N = \text{number of } \underline{\text{states}}$

$$S = \{s_1, \dots, s_N\}$$

2) $M = \text{number of } \underline{\text{symbols observed per state}}$

$$V = \{v_1, v_2, \dots, v_M\}$$

3) $A = \text{the } \underline{\text{transition probability matrix}}$

$$A = [a_{ij}] \quad \begin{matrix} 1 \leq i \leq N \\ 1 \leq j \leq N \end{matrix}$$

$$a_{ij} = \text{Prob} [q_{t+1} = s_j | q_t = s_i]$$

$t = 1, 2, 3, \dots$ - time units

4) states emit symbols
from $V = \{v_1, \dots, v_m\}$

$$B = [b_j(k)]$$

$$1 \leq k \leq m$$

$$1 \leq j \leq N$$

$$b_j(k) = \text{Prob} \left[\begin{array}{l} \text{observing} \\ \text{symbol} \\ v_k \text{ at} \\ \text{time } t \end{array} \middle| \begin{array}{l} q_t = s_j \\ t = j \end{array} \right]$$

9) The initial state
distribution

$$\pi = \{\pi_i\} \quad 1 \leq i \leq N$$

$$\pi_i = \text{Prob} \{q = s_i\}$$

A, B, π are probabilistic
matrices and vector

$$A: \sum_{j=1}^N a_{ij} = 1, \quad 0 \leq a_{ij} \leq 1$$

$$\pi: \sum_{i=1}^N \pi_i = 1, \quad 0 \leq \pi_i \leq 1 \quad \text{constraints}$$

or

$$B: \sum_{k=1}^K b_j(k) = 1, \quad 0 \leq b_j(k) \leq 1$$

probabilistic distributions

what is time unit?

Marker Chaining

$X_1, X_2, X_3, \dots$

$t = 1, 2, 3, \dots$

$t=1 \quad q_1$

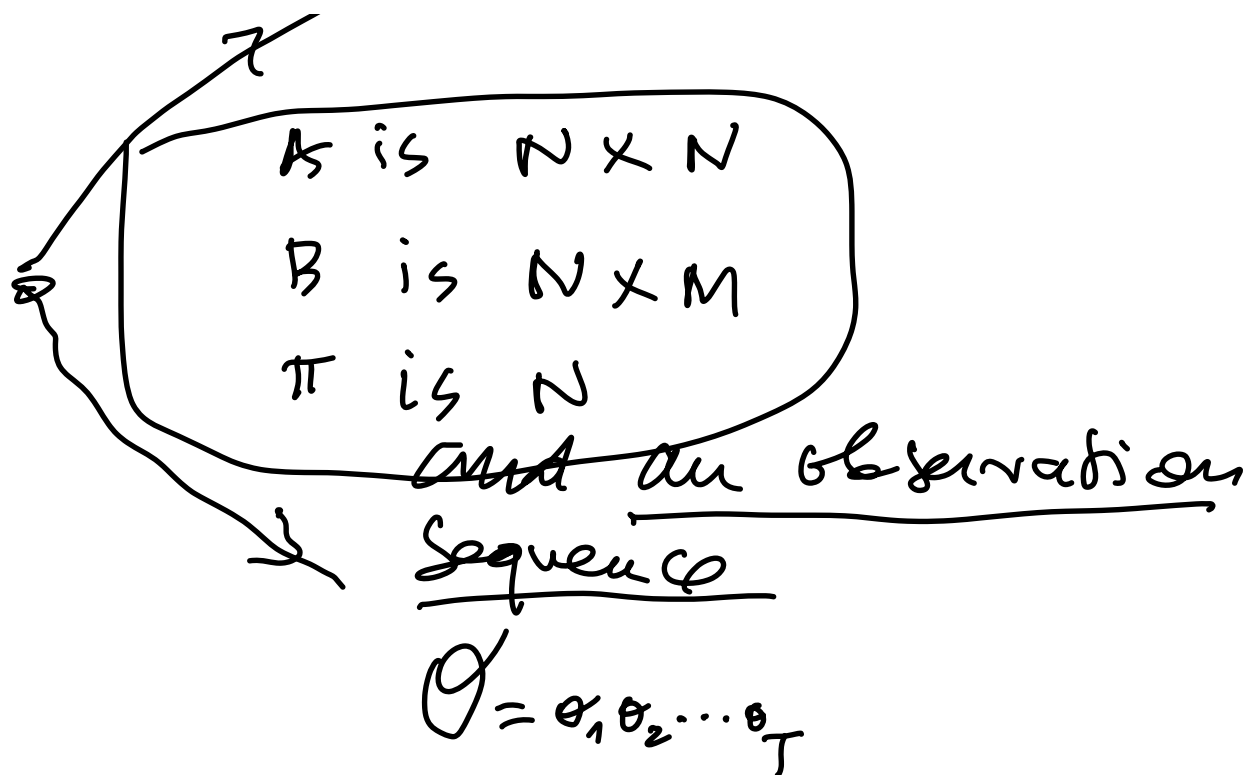
$t=2 \quad q_2$

$\vdots$

3 FUNDAMENTAL COMPUTATIONAL
PROBLEMS FOR HMMs

INPUT: an HMM $\lambda = (A, B, \pi)$

✓



INPUT: $\lambda = (A, B, \pi),$
 θ

OUTPUT:

PROBLEM 1: The EVALUATION PB

INPUT: $\lambda = (A, B, \pi)$

$\theta = \theta_1 \theta_2 \dots \theta_T$, $\theta_i \in V = \{v_1, \dots, v_n\}$
 $1 \leq i \leq T$

OUTPUT:

$P(\theta | \lambda) =$
Probability of observing
the observation sequence
 θ in the model λ .

PROBLEM 2: The DECODING PB

INPUT: $\lambda = (A, B, \pi),$
 θ

OUTPUT: A sequence of
states $Q = \underline{q_1 q_2 \dots q_T}$

which optionally "explains"
the observation sequence

PROBLEM 3 : The LEARNING PB

INPUT : θ

OUTPUT: to find values of
the parameters of
 $\lambda(A, B, \pi)$

Such that $P(\theta | \lambda) = \text{MAX}$

SOLUTION TO PB 1

Problem 1 is the Evaluation PB, i.e.
 given seq θ and model λ ,
 compute the probability of observing
 θ in the model λ .

We wish to compute:

$$Pr(\theta | \lambda)$$

$$\theta = \sigma_1 \dots \sigma_T$$

Remember symbols are emitted by
 states

Ex. $\theta = a a b c c c$ $V = \{a, b, c\}$

$\begin{array}{ccccccc}
s_{10} & s_1 & s_5 & s_2 & s_{10} & & \\
\downarrow & \downarrow & \downarrow & \downarrow & \downarrow & & \\
a & a & b & c & c & c &
\end{array}$

$\begin{array}{ccccccc}
\uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \\
s_1 & s_3 & s_{10} & s_1 & s_2 & s_{25} &
\end{array}$

For any sequence of states of length T
 we can emit a sequence of symbols
 of length T .

This example $O = aabccc$
can be emitted in any sequence
of states of length 6

We have N states

How many sequences of states of
length 6 are there?

$$N^6$$

in general N^T for a
sequence of length T .

N^T is exponential

If $N = 50$, $T = 100$

$N^T =$ more than atoms in
universe

$$T = 1,000,000 \cdot \text{our}$$

AMAZINGLY: Alg $O(N^2)$

Consider one fixed state sequence

$$Q = q_1, \dots, q_T, \quad q_1 = \text{initial state}$$

the prob of observing $\theta = \sigma_1 \dots \sigma_T$ in the state sequence Q

$$P(\theta | Q) = P(\sigma_1 | q_1) P(\sigma_2 | q_2) \dots P(\sigma_T | q_T)$$

statistical independence:

a state s_i emits a symbol completely independent of any other state emission of symbols

$$P(\theta | Q) = b_{q_1}(\sigma_1) b_{q_2}(\sigma_2) \dots b_{q_T}(\sigma_T)$$

what is Prob $P(Q)$?

$$P(Q) = \pi_{q_1} a_{q_1 q_2}^a a_{q_2 q_3}^a \dots a_{q_{T-1} q_T}^a$$

We computed $P(Q)$ and $P(\theta|Q)$

We want now $P(\theta, Q)$ joint prob.

$$P(\theta, Q) = P(\theta|Q) \cdot P(Q)$$

by Prob. theorem

$P(\theta|\lambda)$ = the objective
we want to compute

$$P(\theta|\lambda) = P(\theta) =$$

$$P(\theta) = \sum_{\substack{\text{all seq } \alpha \\ \text{states } Q}} P(\theta|Q) \cdot P(Q)$$

$$= \sum_{q_1, q_2, \dots, q_T} b_{q_1}(\sigma_1) b_{q_2}(\sigma_2) \dots b_{q_T}(\sigma_T) \cdot \pi_{q_1} a_{q_1 q_2} a_{q_2 q_3} \dots a_{q_{T-1} q_T}$$

$$P(\theta) = \sum_{q_1, q_2, \dots, q_T} \pi_{q_1} b_{q_1}(\sigma_1) a_{q_1 q_2} b_{q_2}(\sigma_2) \dots a_{q_{T-1} q_T} b_{q_T}(\sigma_T)$$

$O(N^T)$ alg : impractical.

The FORWARD ALGORITHM

SOL TO PR)

(Dynamic Programming)

Def a variable called forward variable

$$\alpha_t(i) = P(\underbrace{\sigma_1 \sigma_2 \dots \sigma_t}_{\text{prefix}}, q_t = s_i)$$

= The probability of the partial observation sequence $\sigma_1 \sigma_2 \dots \sigma_t$ until time t and being in state s_i at time t .

THE FORWARD ALGORITHM

(1) INITIALIZATION

$$\alpha_1(i) = \pi_i b_i(\sigma_1), \quad 1 \leq i \leq N$$

(2) RECURRENCE

$$\alpha_{t+1}(j) = \left[\sum_{i=1}^N \alpha_t(i) a_{ij} \right] b_j(\sigma_{t+1})$$

$$\begin{aligned} 1 \leq t \leq T-1 \\ 1 \leq j \leq N \end{aligned}$$

(3) TERMINATION

$$P(\theta | x) = \sum_{i=1}^N \alpha_T(i)$$

$O(N^2 T)$ practical
 fast
 exact

Solve the Viterbi Algorithm.

variable δ

The new

$$-X \left[q_1, q_2, \dots, q_{t-1}, q_t = S_i \right]$$

$$\delta_t(i) = \max_{q_1, q_2, \dots, q_{t-1}} \left(\text{probability} \right)$$

= the best score (high σ)
 along the state sequence
 accounts for the prefix $\sigma_1 \dots \sigma_{t-1}$
 of the obs. sequence, and which
 ends in state $q_t = S_i$

TRACE BACK = to find the seq of states that "best explain" the obs seq

$$P(\theta, \underline{Q}) = \text{MAX most probable path } \underline{Q}$$

RABINER TUTORIAL ON HMMs

SOL TO PB 1. More details

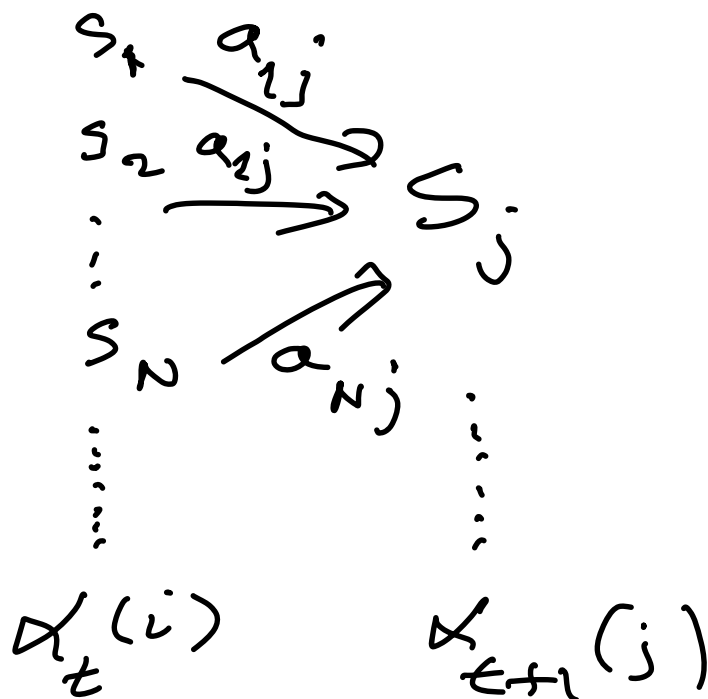
THE RECURRENCE:

RECURRENCE [Step (2)]

$$\alpha_{t+1}(j) = \left[\sum_{i=1}^N \alpha_t(i) a_{ij} \right] b_j(\sigma_{t+1})$$

$$\begin{aligned} 1 \leq t \leq T-1 \\ 1 \leq j \leq N \end{aligned}$$

Step (2)



- s_j can be reached at time $t+1$ from N possible states

s_i , $1 \leq i \leq N$, at time t .

- Since $\alpha_t(i)$ is the prob. of the joint event " $\sigma_1 \sigma_2 \dots \sigma_t$ " is observed and "at time t in state s_i "
- Then $\alpha_t(i) \cdot a_{ij}$ is the prob. of the joint event " $\sigma_1 \sigma_2 \dots \sigma_t$ " observed and state s_j is reached at time $t+1$ via s_i at time t
- Summing up all N states s_i gives the joint prob.

of " $\sigma_1 \sigma_2 \dots \sigma_t$ " observed
and state S_j at time
 $t+1$.

- The concluding $\alpha_{t+1}^{(j)}$
accounts also for the
symbol σ_{t+1} to be emitted

so we need to multiply
by its prob $b_j(\sigma_{t+1})$

- This recurrence is performed
for all states $1 \leq j \leq N$
and for all $t = 1, 2, \dots, T-1$
-

STEP ③

③ TERMINATION

$$P(\theta | \lambda) = \sum_{i=1}^N \alpha_T(i)$$

$P(\theta | \lambda)$ is computed via

$$\alpha_T(i) = P[\sigma_1, \sigma_2, \dots, \sigma_T, \sigma_T = s_i]$$

$$\sum_{i=1}^N \alpha_T(i) \text{ gives you } P(\theta).$$

Time complexity is $O(N^2 T)$

SOLUTION TO PR 2

THE VITERBI ALGORITHM

a new variable

$$\delta_t(i) = \max_{\substack{q_1, q_2, \dots, q_{t-1} \\ \sigma_1, \sigma_2, \dots, \sigma_t}} P[q_1, \dots, q_t = s_i]$$

$$\theta = \sigma_1 \sigma_2 \dots \sigma_T$$

$\delta_t(i)$ = the best score
(highest probability)
along the single
path of states,
at time t ,

which accounts for
the first t observations
 $o_1 o_2 \dots o_t$ and
ends in state S_i

The Recurrence:

$$\delta_{t+1}(j) = \left[\max_i \left\{ \sum_{t'} P_{(i)} a_{ij} \right\} \delta_{t'}(i) \right] b_j(o_{t+1})$$

We will need to backtrack
we will use a new
variable for backtracking

THE VITERBI ALGORITHM

① INITIALIZATION

$$\delta_1(i) = \pi_i b_i(\sigma_1)$$

$$1 \leq i \leq N$$

$$\psi_1(i) = 0$$

$$1 \leq i \leq N$$

② RECURRENCE

$$\delta_t(j) = \max_{1 \leq i \leq N} [\delta_{t-1}(i) a_{ij}] b_j(\sigma_t)$$

$$1 \leq j \leq N$$

$$2 \leq t \leq T$$

in FB1

Σ

$$\psi_t(j) = \underset{1 \leq i \leq N}{\operatorname{Argmax}} \left[\delta_{t-1}(i) a_{ij} \right]$$

$$2 \leq t \leq T$$

$$1 \leq j \leq N$$

③ TERMINATION

$$p^* = \underset{1 \leq i \leq N}{\operatorname{MAX}} \left[\delta_T(i) \right]$$

$$q_T^* = \underset{1 \leq i \leq N}{\operatorname{Argmax}} \left[\delta_T[i] \right]$$

q^* = store index

④ BACKTRACKING

$$q_t^* = \psi_{t+1}(q_{t+1}^*)$$

$$t = T-1, T-2, \dots, 2, 1$$

The matrix V is for
backpointers.

Backtracking reconstruct
the "alignment" between
 O and O_{opt}

$$\begin{array}{l} O = o_1, o_2, \dots, o_T \\ O_{opt} = z_1, z_2, \dots, z_T \end{array} \quad \left(\begin{array}{c} o_i \\ \uparrow \\ z_i \end{array} \right)$$

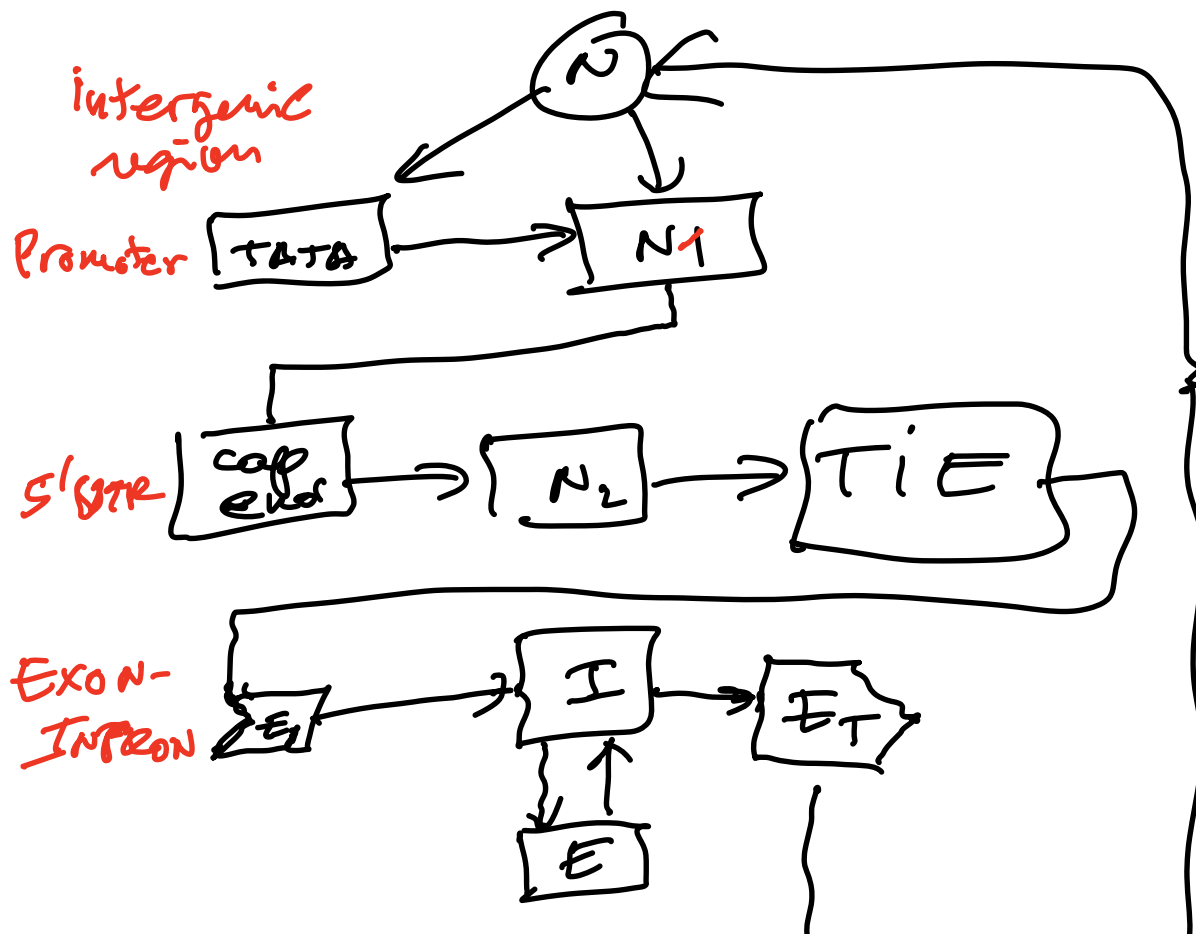
Reconstruction through
backtracking goes from
right to left

(like in the ^{seq}align
backtracking)

$Q_{opt} = q_1 \dots q_T$ is the
"decoding"

revealed the hidden
states

Gene Finding



Post
translational
regional

