

# Algorithms in Genome Analysis, Spring 2023

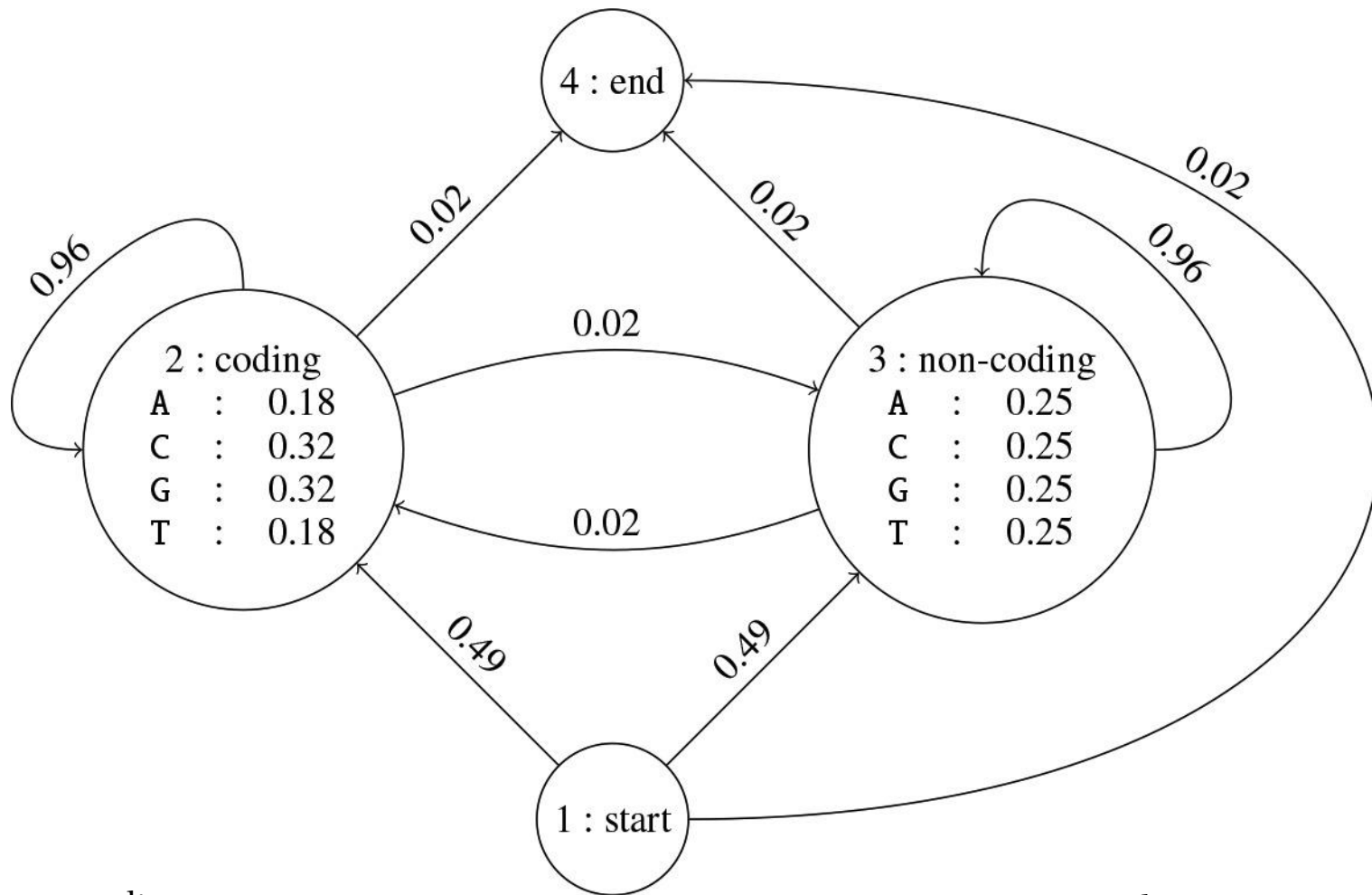
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# Week 5

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Hidden Markov Models (HMMs)

Max probable path = segmentation into coding and non-coding regions



coding

coding

CACCGCTGCCGAAATTCTATGGTTAGTACTGTCATGATCTTCCGAGTCCGAATCAGA

# HMM

- Directed graph with set  $H$  of hidden states emitting symbols and set  $T$  of transitions
- Emission probabilities  $p(c | h)$  in state  $h$  sum to 1
- Transition probabilities  $p(h | h')$  to exit state  $h'$  sum to 1
- Unique source (no transitions in) and sink (no transitions out)
- Given an input sequence, what is the most probable path from source to sink?
- Many other interesting questions to be studied under this framework

# Viterbi

- Dynamic programming algorithm solving the most probable path problem
- $v(i, h)$  = max probability of a path emitting  $S[1..i]$  so that  $S[i]$  is emitted at state  $h$
- $v(i, h) = p(S[i] | h) * \max_{(h', h) \in T} v(i - 1, h') * p(h | h')$
- Initialization  $v(0, start) = 1$
- Finalization  
 $v(|S| + 1, end) = \max_{(h, end) \in T} v(|S|, h) * p(end | h)$   
Most probable path can be traced back from  $v(|S| + 1, end)$
- For numerical accuracy, log probabilities are used so that all multiplications become summations

	$h_{\text{start}}$		$x$		$h$		$y$		$z$		$h_{\text{end}}$
0	1		0		0		0		0		0
1											
2											
$\vdots$											
$i-1$											
$i$											
$n$											

$$v(i, h) = \mathbb{P}(s_i | h) \max(v(i-1, x)\mathbb{P}(h | x), v(i-1, y)\mathbb{P}(h | y), v(i-1, z)\mathbb{P}(h | z))$$

# Training

- With labeled training data one can use the observed frequencies to fix the emission and transition probabilities
- Without labels, common ways to proceed are
  - **Viterbi training**: Set initial probabilities based on background knowledge, find a most probable path  $P$  for input sequence  $S$ , label  $S$  according to  $P$  and use this as labeled data. Iterate.
  - **Expectation maximization (EM)**: As above, but take all paths into account in proportion to their probabilities, calculating expected probability for each emission / transition. Iterate.

# EM through Baum-Welsh 1/2

- Uses **forward-backward** variant of viterbi to find EM estimates
- $f(i, h)$  = sum of probabilities of a paths emitting  $S[1..i]$  so that  $S[i]$  is emitted at state  $h$
- $f(i, h) = p(S[i] | h) * \sum_{(h',h) \in T} f(i - 1, h') * p(h | h')$
- $b(i, h)$  = sum of probabilities of a paths emitting  $S[i..|S|]$  so that  $S[i]$  is emitted at state  $h$
- $b(i, h) = p(S[i] | h) * \sum_{(h,h') \in T} b(i + 1, h') * p(h' | h)$
- Initialization:  $f(0, start) = b(|S| + 1, end) = 1$
- Finalization:
  - $f(|S| + 1, end) = \sum_{(h,end) \in T} f(|S|, h) * p(end | h)$
  - $b(0, start) = \sum_{(start,h) \in T} b(1, h) * p(h | start)$



# EM through Baum-Welch 2/2

- Note:  $f(|S| + 1, end) = b(0, start)$  is the total probability of emitting  $S$
- The total probability  $T(h, i)$  of emitting  $S[i]$  at state  $h$  is

$$\sum_{(h', h) \in T} f(i - 1, h') * p(h|h') * b(i, h)$$

- Expected emission count  $EC(h, c)$  of seeing  $c$  being emitted at state  $h$  is thus

$$\frac{\sum_{\{i: S[i]=c\}} T(h, i)}{f(|S| + 1, end)}$$

- One can then set  $p(c | h) = \frac{EC(h, c)}{\sum_{\{c' \in \Sigma\}} EC(h, c')}$
- Derivation for  $p(h | h')$  is left as an exercise

	$h_{\text{start}}$	$h'$	$x$		$h$	$y$		$z$		$h_{\text{end}}$
0	1		0		0	0		0		0
1										
2										
$i-1$										
$i$										
$n$										

$$T(h, i) = \sum_{(h, h') \in T} f(i, h) * p(h' | h) * b(i + 1, h')$$

Diagram illustrating the calculation of  $v(i, h)$  at time step  $i$  and hidden state  $h$ . The value  $v(i, h)$  is determined by the maximum of three options, each weighted by a function  $f$ :

- Option 1:  $f \cdot v(i-1, x) \cdot \mathbb{P}(h | x)$
- Option 2:  $f \cdot v(i-1, y) \cdot \mathbb{P}(h | y)$
- Option 3:  $f \cdot v(i-1, z) \cdot \mathbb{P}(h | z)$

The overall value is also influenced by the transition function  $T(h, i)$  and the emission function  $b(i+1, h')$ .