







HMM inference. Forward algorithm

Better solution: decompose the computation along the time

• Let $\alpha_t(i) = P(O_1, O_2, ..., O_t, s_t = i | \lambda)$

Algorithm:

- 1. $\alpha_1(i) = \pi_i b_i(O_1)$ for all *i*
- 2. Repeat for $t = 2, 3, \dots$ *T*-1 and all *j*

$$\alpha_{t+1}(j) = \left[\sum_{i} \alpha_{t}(i)a_{ij}\right]_{i} b_{j}(O_{t+1})$$

3. Then $P(O \mid \lambda) = \sum_{i} \alpha_{T}(i)$

This is called forward algorithm

- Implements a dynamic programming approach
- It is polynomial in the number of states and steps

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HMM inference. Backward algorithm

- We can do the computation backward in time as well
- Let $\beta_t(i) = P(O_{t+1}, O_{t+2}, ..., O_T | s_t = i, \lambda)$

Backward algorithm :

- 1. $\beta_T(i) = 1$ for all *i* 2. Repeat for t = T - I, T - 2, ..., I and for all *i* $\beta_t(i) = \sum_j a_{ij} b_j(O_{t+1}(j)) \beta_{t+1}(j)$ 3. Then $P(O \mid \pi) = \sum_i \pi_i \beta_1(i)$ • The algorithm
 - is polynomial in the number of states and steps

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Finding the most likely sequence

- **Goal:** compute the most likely sequence of states
- **Problem:** How to define the most likely sequence?
- Solution 1: piece together most likely states
- A probability of being in state *i* at time *t*, given the complete observation sequence

• **Problem:** what if the transition between the two most likely states is 0?

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Viterbi algorithm• Initialization: (for all i) $\delta_1(i) = \pi_i b_i(O_1)$ $\psi_1(i) = 0$ • Recursion (for all j) $\delta_t(j) = \max_{1 \le i \le N} \left[\delta_{t-1} a_{ij} \right] b_j(O_t)$ $\psi_t(j) = \max_{1 \le i \le N} \left[\delta_{t-1} a_{ij} \right]$ • Termination $P^* = \max_{1 \le i \le N} \left[\delta_T \right]$ $s_T^* = \arg \max_{1 \le i \le N} \left[\delta_T \right]$ • Sequence Recovery (T-1, ..., 1) $s_t^* = \psi_{t+1} \left[s_{t+1}^* \right]$

HMM learning • Learning with hidden variables (the same idea as used in EM) • **Baum-Welch algorithm** is a special case of EM • Find the ML set of parameters , maximizing $P(O | \lambda)$ • **HMM re-estimation step** (one cycle of EM): $\gamma_t(i) = P(s_t = i | O, \lambda)$ $\xi_t(i, j) = P(s_t = i, s_{t+1} = j | O, \lambda)$ $\gamma_t(i) = \frac{\alpha_t(i)\beta_t(i)}{P(O | \lambda)}$ $\xi_t(i, j) = \frac{\alpha_t(i)\alpha_{ij}b_j(O_{t+1})\beta_{t+1}(i)}{P(O | \lambda)}$ $\widetilde{\pi}_i = \gamma_1(i)$ $\widetilde{\alpha}_{ij} = \frac{\sum_{t=1}^T \xi_t(i, j)}{\sum_{t=1}^T \gamma_t(i)}$ $\widetilde{b}_j(k) = \frac{\sum_{t=1,O_t=k}^T \gamma_t(j)}{\sum_{t=1}^T \gamma_t(j)}$ ES 2750 Machine Learning

HMMs in practice

- HMMs have been widely used in various contexts
- **Speech recognition** (single word recognition)
 - words correspond to sequences of observations
 - we estimate a HMM for each word
 - the output model is a mixture of Gaussians over spectral features
- Bio-sequence analysis
 - a single HMM model for each type of protein (sequence of amino acids)
 - gene identication (parsing the genome)
 - etc.

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