Hidden Markov Model I

- Assume there are two types of weather "Sunny" and "Rainy". We assume, a prior, that the probabilities of two types of days are 0.7 and 0.3, e.g., Pr(Sunny) = 0.7, Pr(Rainy) = 0.3.
- Every morning, you do two things: walk our dog ("W") or reading ("R"). Assume the following conditional probabilities:

$$Pr(W|S unny) = 0.8, Pr(R|S unny) = 0.2.$$

 $Pr(W|Rainy) = 0.2, Pr(R|Rainy) = 0.8.$

Assume we know your morning activity for a number of days: {W, W, R, R, W, W, R, W, W, W, W, W, W}. How can we estimate the weather condition for each day?

Using Bayes' rule, one can do for each day:

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Pr(Sunny|W) = \frac{Pr(W|Sunny)Pr(Sunny)}{Pr(W|Sunny)Pr(Sunny) + Pr(W|Rainy)Pr(Rainy)}
= \frac{0.8 * 0.7}{0.8 * 0.7 + 0.2 * 0.3} = 0.9
Pr(Sunny|R) = \dots
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- However, this assumes independence of observations and completely ignores
 the connections between weather changes, e.g., probability of today is Sunny
 given yesterday is Sunny, etc.
- With the consideration of those, today's weather Pr(Sunny|W) should also depend on yesterday's weather, in addition to the W/R status.
- Such an approach can be formalized by a "hidden Markov model" (HMM).

- Assume we observe sequential data $\mathbf{u} = \{u_1, u_2, \dots, u_T\}$ (your morning activities).
- u is generated by a **hidden**, unobserved state: $s = \{s_1, s_2, \dots, s_T\}$.
- Each s_t can take M states, with "initial probability" $\pi_k, k = 1, ..., M$: $Pr(s_1 = k) = \pi_k, \sum_k \pi_k = 1$.
- The distribution of u conditional on s is specified by a distribution $b_k(u)$: $u_t|s_t=k\sim b_k(u_t)$. This is called "emission probability".
- The changes of states between consecutive hidden state is specified by "transition probability": $a_{k,l} = Pr(s_{t+1} = l | s_t = k)$. Or you can write this as $a_{k \to l}$.
- Assume the underlying states follow a Markov chain, that is, given present, the future is independent of the past:

$$Pr(s_{t+1}|s_t, s_{t-1}, \ldots, s_1) = Pr(s_{t+1}|s_t).$$

To summarize: Observed data: u. Missing data: s. Parameters: $\lambda = \{\pi_k, b_k(u), a_{k,l}\}$.

Review: discrete time finite homogeneous Markov Chain— 4/22 —

- The possible states are included in a finite discrete set: $\{E_1, E_2, \dots, E_M\}$.
- From time t to t+1, make stochastic movement from one state to another.
- Markov Property: the state of s_{t+1} only depends on the state of s_t , not the states before time t:

$$Pr(s_{t+1}|s_t, s_{t-1}, \ldots, s_1) = Pr(s_{t+1}|s_t).$$

- Time-homogeneous transition probabilities property: $P(s_{t+1}|s_t)$ independent of t.
- Denote the transition probability matrix by **A**. Define N step transition as: $a_{k,l}(N) = Pr(s_{t+N} = l | s_t = k)$. It can be shown that $\mathbf{A}(N) = \mathbf{A}^N$.

A HMM can answer following questions:

- Parameter estimation: estimate the initial/emission/transition probabilities. $\hat{\lambda} = \operatorname{argmax}_{\lambda} Pr(\boldsymbol{u}|\lambda)$.
- What are the probabilities of the underlying states, given the observations: Pr(s|u).
- The most likely path: given the observed data, what are the most likely underlying states for all observations: $\hat{s} = \operatorname{argmax}_{s} Pr(s|\lambda, u)$.
- Predict future, e.g., $Pr(u_{t+1}|\mathbf{u}, \hat{\lambda})$.

Examples of HMM applications:

- Speech recognition.
- DNA sequence analysis, e.g., gene finding, sequence alignment.
- Financial time series data.

- There's close connection between a HMM and a mixture model: both have hidden states/group assignment, initial and emission probabilities.
- Difference is that mixture model assumes **independent** observations, HMM assumes sequential observation with transition probability.

According to Markov property, we have:

Joint probability of hidden states:

$$P(s_1, s_2, \dots, s_T) = P(s_1)P(s_2|s_1)\dots P(s_T|s_{T-1})$$

= $\pi_{s_1}a_{s_1, s_2}\dots a_{s_{T-1}, s_T}$

• Conditional on the states, the observations are independent of each other:

$$P(u_i, u_i|s) = P(u_i|s)P(u_i|s)$$

So the joint probability of observations, given hidden states is:

$$P(\boldsymbol{u}|\boldsymbol{s}) = \prod_{i=1}^{T} P(u_i|s_i) = \prod_{i=1}^{T} b_{s_i}(u_i)$$

Joint probability of hidden states and observed data

$$P(\mathbf{u}, \mathbf{s}) = P(\mathbf{s})P(\mathbf{u}|\mathbf{s})$$

= $\pi_{s_1}b_{s_1}(u_1)a_{s_1, s_2}b_{s_2}(u_2)a_{s_2, s_3} \dots a_{s_{T-1}, s_T}b_{s_T}(u_T)$

Marginal probability of observed data:

$$P(\mathbf{u}) = \sum_{s} P(s)P(\mathbf{u}|s)$$

$$= \sum_{s} \pi_{s_1}b_{s_1}(u_1)a_{s_1,s_2}b_{s_2}(u_2)a_{s_2,s_3}\dots a_{s_{T-1},s_T}b_{s_T}(u_T)$$

- First need to make parametric assumption of the emission probabilities $b_k(u)$.
- A common assumption is that $b_k(u)$ is Normal, e.g., $b_k(u) = N(u : \mu_k, \sigma_k^2)$.
- Then the model parameters to be estimated are:

$$\lambda = \{\pi_k, \mu_k, \sigma_k, a_{k,l} : k, l = 1, \dots, M\}$$

- One can obtain the MLEs for λ from the marginal probability of observed data. However it's very difficult because the marginal probability involves summing over all possible paths (\sum_s) .
- Clever algorithm was invented to solve the problem.

• Define $L_k(t)$ be the conditional probability of being in state k at position t given the observed data u:

$$L_k(t) = P(s_t = k|\boldsymbol{u})$$

• Define $H_{k,l}(t)$ be the conditional probability of being in state k at position t and being in state l at position t+1 (i.e., seeing a transition from k to l at t), given the observed data u:

$$H_{k,l}(t) = P(s_t = k, s_{t+1} = l|\mathbf{u})$$

• Note that $L_k(t) = \sum_{l=1}^{M} H_{k,l}(t), \sum_{l=1}^{M} L_k(t) = 1.$

- Then the parameters can be estimated by EM:
 - E-step: Compute $L_k(t)$ and $H_{k,l}(t)$ given current parameters.
 - M-step: update parameters:

$$\mu_{k} = \frac{\sum_{t=1}^{T} L_{k}(t)u_{t}}{\sum_{t=1}^{T} L_{k}(t)}$$

$$\sigma_{k}^{2} = \frac{\sum_{t=1}^{T} L_{k}(t)(u_{t} - \mu_{k})^{2}}{\sum_{t=1}^{T} L_{k}(t)}$$

$$a_{k,l} = \frac{\sum_{t=1}^{T-1} H_{k,l}(t)}{\sum_{t=1}^{T-1} L_{k}(t)}$$

$$\pi_{k} = \frac{\sum_{t=1}^{T} L_{k}(t)}{T}$$

• Derivation steps are similar to that in M-component normal mixture model (try it yourself). The new items are the transition probabilities.

- In the M-step to update Normal distribution parameters, $L_k(t)$ plays the same role as the posterior probability (expected value) of a component (state) given the observation, i.e., $p_{t,k} = P(s_t = k|u_t)$.
- In comparison, $L_k(t) = P(s_t = k | u_1, u_2, \dots, u_T)$.
- If one ignores the connections among observations, e.g., s_t 's are independent and thus u_t 's are iid, then $L_k(t) = p_{t,k}$, and HMM reduce to a M-component Normal mixture model.
- In a mixture model, s_t only depends on u_t because observations are independent.
- In a HMM, s_t depends on the entire sequence of observations because of the underlying Markov process.

The forward-backward algorithm is designed to efficiently compute:

$$L_k(t) = P(s_t = k|\mathbf{u})$$

$$H_{k,l}(t) = P(s_t = k, s_{t+1} = l|\mathbf{u})$$

• Define the **forward probability** $\alpha_k(t)$ as the **joint probability** of observing the first t data u_i , i = 1, ..., t and being in state k at time t:

$$\alpha_k(t) = P(u_1, u_2, \dots, u_t, s_t = k)$$

The forward probability can be computed recursively:

$$\alpha_k(1) = \pi_k b_k(u_1) \quad 1 \le k \le M$$

$$\alpha_k(t) = b_k(u_t) \sum_{l=1}^{M} \alpha_l(t-1) a_{l,k} \quad 1 < t \le T, 1 \le k \le M.$$

$$a_{k}(t) = P(u_{1}, u_{2}, \dots, u_{t}, s_{t} = k)$$

$$= \sum_{l=1}^{M} P(u_{1}, u_{2}, \dots, u_{t}, s_{t} = k, s_{t-1} = l)$$

$$= \sum_{l=1}^{M} P(u_{1}, u_{2}, \dots, u_{t-1}, s_{t-1} = l)P(u_{t}, s_{t} = k \mid u_{1}, u_{2}, \dots, u_{t-1}, s_{t-1} = l)$$

$$= \sum_{l=1}^{M} \alpha_{l}(t-1)P(u_{t}, s_{t} = k \mid s_{t-1} = l)$$

$$= \sum_{l=1}^{M} \alpha_{l}(t-1)P(u_{t} \mid s_{t} = k, s_{t-1} = l)P(s_{t} = k \mid s_{t-1} = l)$$

$$= \sum_{l=1}^{M} \alpha_{l}(t-1)P(u_{t} \mid s_{t} = k)P(s_{t} = k \mid s_{t-1} = l)$$

$$= b_{k}(u_{t}) \sum_{l=1}^{M} \alpha_{l}(t-1)a_{l,k}$$

• Define the **backward probability** $\beta_k(t)$ as the **conditional probability** of observing the data after time t, u_i , i = t + 1, ..., T, given the state at time t is k.

$$\beta_k(t) = P(u_{t+1}, \dots, u_T \mid s_t = k) \quad 1 \le t \le T - 1$$

Again, the backward probability can be computed by following recursive formula:

$$\beta_k(T) = 1$$

$$\beta_k(t) = \sum_{l=1}^{M} a_{k,l} b_l(u_{t+1}) \beta_l(t+1) \quad 1 \le t < T$$

$$\beta_{k}(t) = P(u_{t+1}, \dots, u_{T} \mid s_{t} = k)$$

$$= \sum_{l=1}^{M} P(u_{t+1}, \dots, u_{T}, s_{t+1} = l \mid s_{t} = k)$$

$$= \sum_{l=1}^{M} P(u_{t+1}, \dots, u_{T} \mid s_{t+1} = l, s_{t} = k) P(s_{t+1} = l \mid s_{t} = k)$$

$$= \sum_{l=1}^{M} P(u_{t+1}, \dots, u_{T} \mid s_{t+1} = l) a_{k,l}$$

$$= \sum_{l=1}^{M} P(u_{t+2}, \dots, u_{T} \mid s_{t+1} = l, u_{t+1}) P(u_{t+1} \mid s_{t+1} = l) a_{k,l}$$

$$= \sum_{l=1}^{M} P(u_{t+2}, \dots, u_{T} \mid s_{t+1} = l) b_{l}(u_{t+1}) a_{k,l}$$

$$= \sum_{l=1}^{M} a_{k,l} b_{l}(u_{t+1}) \beta_{l}(t+1)$$

Compute $L_k(t)$ using forward and backward probabilities:

$$L_k(t) \equiv P(s_t = k \mid \boldsymbol{u}) = \frac{P(\boldsymbol{u}, s_t = k)}{P(\boldsymbol{u})} = \frac{\alpha_k(t) \beta_k(t)}{P(\boldsymbol{u})}$$

Proof:

$$P(u, s_t = k) = P(u_1, ..., u_T, s_t = k)$$

$$= P(u_1, ..., u_t, s_t = k) P(u_{t+1}, ..., u_T \mid u_1, ..., u_t, s_t = k)$$

$$= P(u_1, ..., u_t, s_t = k) P(u_{t+1}, ..., u_T \mid s_t = k)$$

$$= \alpha_k(t) \beta_k(t)$$

Compute $H_{k,l}(t)$ using forward and backward probabilities:

$$H_{k,l}(t) = P(s_t = k, s_{t+1} = l | \mathbf{u}) = \frac{P(s_t = k, s_{t+1} = l, \mathbf{u})}{P(\mathbf{u})}$$
$$= \frac{1}{P(\mathbf{u})} \alpha_k(t) \ a_{k,l} \ b_l(u_{t+1}) \ \beta_l(t+1)$$

Proof:

$$P(s_{t} = k, s_{t+1} = l, \mathbf{u}) = P(u_{1}, \dots, u_{t}, \dots, u_{T}, s_{t} = k, s_{t+1} = l)$$

$$= P(u_{1}, \dots, u_{t}, s_{t} = k)P(u_{t+1}, s_{t+1} = l \mid s_{t} = k, u_{1}, \dots, u_{t})$$

$$P(u_{t+2}, \dots, u_{T} \mid s_{t+1} = l, s_{t} = k, u_{1}, \dots, u_{t+1})$$

$$= \alpha_{k}(t)P(u_{t+1}, s_{t+1} = l \mid s_{t} = k)P(u_{t+2}, \dots, u_{T} \mid s_{t+1} = l)$$

$$= \alpha_{k}(t)P(s_{t+1} = l \mid s_{t} = k)P(u_{t+1} \mid s_{t+1} = l, s_{t} = k)\beta_{l}(t+1)$$

$$= \alpha_{k}(t)P(s_{t+1} = l \mid s_{t} = k)P(u_{t+1} \mid s_{t+1} = l)\beta_{l}(t+1)$$

$$= \alpha_{k}(t) a_{k,l} b_{l}(u_{t+1}) \beta_{l}(t+1)$$

The joint observed data likelihood is:

$$P(\boldsymbol{u}) = \sum_{k=1}^{M} \alpha_k(t) \beta_k(t)$$

Proof:

$$P(\mathbf{u}) = \sum_{k=1}^{M} P(u_1, \dots, u_t, \dots, u_T, s_t = k)$$

$$= \sum_{k=1}^{M} P(u_1, \dots, u_t, s_t = k) P(u_{t+1}, \dots, u_T \mid s_t = k, u_1, \dots, u_t)$$

$$= \sum_{k=1}^{M} P(u_1, \dots, u_t, s_t = k) P(u_{t+1}, \dots, u_T \mid s_t = k)$$

$$= \sum_{k=1}^{M} \alpha_k(t) \beta_k(t)$$

To summarize, estimation of model parameters requires iterating following steps, under the current estimates of parameters:

1. Compute the forward and backward probabilities:

$$\alpha_k(1) = \pi_k b_k(u_1) \quad 1 \le k \le M$$

$$\alpha_k(t) = b_k(u_t) \sum_{l=1}^{M} \alpha_l(t-1) a_{l,k} \quad 1 < t \le T, 1 \le k \le M.$$

$$\beta_k(T) = 1$$

$$\beta_k(t) = \sum_{l=1}^{M} a_{k,l} b_l(u_{t+1}) \beta_l(t+1) \quad 1 \le t < T$$

- 2. Compute whole data likelihood: $P(\mathbf{u}) = \sum_{k=1}^{M} \alpha_k(t)\beta_k(t)$.
- 3. Compute $L_k(t)$ and $H_{k,l}(t)$ from forward/backward probabilities:

$$L_k(t) = \frac{\alpha_k(t) \beta_k(t)}{P(\boldsymbol{u})}$$

$$H_{k,l}(t) = \frac{1}{P(\boldsymbol{u})} \alpha_k(t) a_{k,l} b_l(u_{t+1}) \beta_l(t+1)$$

4. Update parameters using $L_k(t)$ and $H_{k,l}(t)$ (assuming Normal emission probabilities):

$$\mu_{k} = \frac{\sum_{t=1}^{T} L_{k}(t)u_{t}}{\sum_{t=1}^{T} L_{k}(t)}, \quad \sigma_{k}^{2} = \frac{\sum_{t=1}^{T} L_{k}(t)(u_{t} - \mu_{k})^{2}}{\sum_{t=1}^{T} L_{k}(t)},$$

$$a_{k,l} = \frac{\sum_{t=1}^{T-1} H_{k,l}(t)}{\sum_{t=1}^{T-1} L_{k}(t)}, \quad \pi_{k} = \frac{\sum_{t=1}^{T} L_{k}(t)}{T}$$

- HMM is used to model sequential data.
- Difference between HMM and mixture model: mixture model assumes iid observations, HMM assumes underlying sequential correlation among hidden states.
- Important components in a HMM: initial, emission and transition probabilities.
- Goals of HMM: estimate hidden states and model parameters, find best path, future prediction.
- Parameter estimation via EM and forward-backward algorithm.
- Next lecture: dynamic programming and Viterbi algorithm to find the best path.