Hidden Markov model and its application in sequence analysis

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CpG islands

- Example: CpG islands
- CpG site: CpG is a pair of nucleotides C and G, appearing successively in this order along one DNA strand. "CpG" stands for cytosine and guanine separated by a phosphate, which links the two nucleosides together in DNA.
- CpG is relative rare in vertebrate genomes due to certain biochemical properties.
- Frequency of CpG dinucleotides in the human genome: Based on 42% GC content (like the human genome).
 - 1. Expected frequency: 0.21*0.21 = 4.41%
 - 2. Actual frequnency: 1%

CpG islands detection

CpG islands

Regions of the DNA that have a higher concentration of CpG sites, typically several hundreds of nucleotides long. What are CpG islands interesting? They are typically associated with the promoter regions of genes. About half of the genes in human have a CpG island present in their promoters.

CpG island detection problem
Input: A long DNA sequences $X = (x_1, \dots, x_L) \in \Sigma^*$ Question: Locate all CpG lands within X.

An occasionally dishonest casino dealer

Suppose a dealer in a casino rolls a die. The dealer use a fair die most of the time, but occasionally he switches to a loaded die. The loaded die has probability 0.5 of a six and probability 0.1 for the numbers 1 to 5.

Hidden Markov Model

Definition: A hidden Markov model (HMM) is a triplet $M = (\Sigma, Q, \Theta)$, where

- ▶ Σ is an alphabet of symbols, e.g. $\Sigma = \{1, 2, 3, 4, 5, 6\}$.
- Q is a finite set of states, capable of emiting symbols from ∑, e.g.
 $Q = \{F, L\}.$
- Θ is a set of probabilities, consisting of

 1) State transition probabilities, a_{kl} for all k, l ∈ Q. and 2) Emission
 probabilities.

Hidden Markov Model

State transition probabilities, a_{kl} for all $k, l \in Q$.

$$a = \begin{bmatrix} a_{FF} & a_{FL} \\ a_{LF} & a_{LL} \end{bmatrix} = \begin{bmatrix} 0.95 & 0.05 \\ 0.1 & 0.9 \end{bmatrix}$$

• Emission probabilities: $e_k(b)$ for all $k \in Q$ and $b \in \Sigma$.

$$e_F = (1/6, 1/6, 1/6, 1/6, 1/6, 1/6)$$

$$e_L = (0.1, 0.1, 0.1, 0.1, 0.1, 0.5)$$
(1)
(2)

Denote $\Theta = (a, e_F, e_L)$.

Path and joint prob

▲ A path $\Pi = (\pi_1, \dots, \pi_L)$ in the model *M* is a sequence of states, modeled by a Markov chain.

$$P(\pi_i = l | \pi_{i-1} = k) = a_{kl}$$

Joint probability:

$$P(X,\Pi) = a_{\pi_0,\pi_1} \prod_{i=1}^{L} e_{\pi_i}(x_i) a_{\pi_i,\pi_{i+1}}$$

where $\pi_0 = begin$ and $\pi_{L+1} = end$.

• A sequence $X = (x_1, \cdots, x_L) \in \Sigma^*$:

$$P(x_i = b | \pi_i = k) = e_k(b)$$

Decoding problem

The decoding problem
 Input: A HMM M = (Σ, Q, Θ) and a sequence X = Σ*, for which the generating path Π = (π₁, · · · , π_L) is unknown.
 Question: Find the most probable path ÎI for X, i.e.

$$\hat{\Pi} = \underset{\Pi}{\arg\max} P(X, \Pi)$$

Decoding problem: Viterbi algorithm

Consider a path ending at $k \in Q$, and the probability of Π generating the prefix (x_1, \dots, x_i) of X.

$$v_k(i) = \max_{\Pi:\pi_i=k} P(x_1, \cdots, x_i, \Pi)$$

$$v_{begin}(0) = 1; \quad v_k(0) = 0 \ \forall k \neq begin$$

● For $i = 0, \cdots, L - 1$ and $l \in Q$,

$$v_l(i+1) = e_l(x_{i+1}) \max_{k \in Q} \{v_k(i)a_{kl}\}$$



$$P(X,\hat{\Pi}) = \max_{k \in Q} \left\{ v_k(L)a_{k,end} \right\}$$

The posterior decoding problem

The posterior decoding problem
 Input: A HMM M = (Σ, Q, Θ), and a sequence X ∈ Σ*, for which the generating path Π = (π₁, · · · , π_L) is unknown.
 Question: For all 1 ≤ i ≤ L and k ∈ Q, find P(π_i = k|X).

Posterior decoding: forward algorithm

Denote by $f_k(i)$ the prob of emitting the prefix (x_1, \dots, x_i) and eventually reaching state $\pi_i = k$:

$$f_k(i) = P(x_1, \cdots, x_i, \pi_i = k)$$

$$f_{begin}(0) = 1; \quad f_k(0) = 0 \ \forall k \neq begin$$

● For
$$i = 0, \cdots, L - 1$$
 and $l \in Q$,

$$f_l(i+1) = e_l(x_{i+1}) \sum_{k \in Q} f_k(i) a_{kl}$$



$$P(X) = \sum_{k \in Q} f_k(L) a_{k,end}$$

Posterior decoding: backward algorithm

Denote by $b_k(i)$ the prob of emitting the suffix $(x_{i=1}, \dots, x_L)$ given $\pi_i = k$:

$$b_k(i) = P(x_{i+1}, \cdots, x_L | \pi_i = k)$$

Initialize

$$b_k(L) = a_{k,end} \quad \forall k \in Q$$

For $i = L - 1, \dots, 1$ and $l \in Q$,
 $b_l(i) = \sum_{l \in Q} a_{kl} e_l(x_{i+1}) b_k(i+1)$
Finally

$$P(X) = \sum_{l \in Q} a_{begin, l} e_l(x_1) b_l(1)$$

The posterior decoding problem

The posterior decoding problem

$$P(X, \pi_i = k) = P(x_1, \cdots, x_i, \pi_i = k) P(x_{i+1}, \cdots, x_L | x_1, \cdots, x_i, \pi_i = k)$$
(3)

$$=P(x_1, \cdots, x_i, \pi_i = k)P(x_{i+1}, \cdots, x_L | \pi_i = k)$$
(4)

$$=f_k(i)b_k(i) \tag{5}$$

Hence,

$$P(\pi_i = k | X) = \frac{f_k(i)b_k(i)}{P(X)}$$

Parameter estimation

Parameter estimation for HMM
 Input: Given *n* example sequences X⁽¹⁾, ..., X⁽ⁿ⁾ ∈ Σ* of length L⁽¹⁾, ..., L⁽ⁿ⁾, respectively, which were generated from a HMM
 M = (Σ, Q, Θ) with unknown Θ.
 Question: Find the most probable Ô, that is

$$\hat{\Theta} = \underset{\Theta}{\operatorname{arg\,max}} P(X^{(1)}, \cdots, X^{(n)} | \Theta)$$

Parameter estimation I: state is known

When the state sequence is known Suppose we know the state sequences $\Pi^{(1)}, \dots, \Pi^{(n)} \in \Sigma^*$ corresponding to $X^{(1)}, \dots, X^{(n)} \in \Sigma^*$, respectively. We can then scan the sequences and compute

1. A_{kl} - the number of transitions from state k to l.

2. $E_k(b)$ - the number of times that an emission of the symbol *b* occurred in state *k*.

Then the ML estimators will be

$$a_{kl} = \frac{A_{kl}}{\sum_{q \in Q} A_{kq}}$$

$$e_k(b) = \frac{E_k(b)}{\sum_{\sigma \in \Sigma} E_k(\sigma)}$$
(6)
(7)

Parameter estimation I: state is unknown

- When the state sequence is unknown: Baum-Welch algorithm
 - 1. Initialization: Assign random values to Θ .
 - 2. Expectation

EM-algorithm: Expectation

Calculate the expected number of state transitions from statet k to l. Note that

$$P(\pi_i = k, \pi_{i+1} = l | X, \Theta) = \frac{f_k(i)a_{kl}e_l(x_{i+1})b_l(i+1)}{P(X)}$$

Let $\{f_k^{(j)}(i), b_k^{(j)}\}$ denote the forward and backward probabilities of the sequence $X^{(j)}$. Then the expectation is

$$A_{kl} = \sum_{j=1}^{n} \frac{1}{P(X^{(j)})} \sum_{i=1}^{L^{(j)}} f_k^{(j)}(i) \ a_{kl} \ e_l(x_{i+1}^{(j)}) \ b_l^{(j)}(i+1)$$

EM-algorithm: Expectation

Calculate the expected number of emissions of the symbol b that occurred at the state k,

$$E_k(b) = \sum_{j=1}^n \frac{1}{P(X^{(j)})} \sum_{\{i:x_i^{(j)}=b\}} f_k^{(j)}(i) \ b_k^{(j)}(i)$$

EM-algorithm: Expectation

- Initialize
- Repeat until convergence
 - Expectation
 - Maximization: Update Θ using A_{kl} and $E_k(b)$.