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The Concepts of Hidden Markov Model in Speech Recognition

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1. Introduction

The speech recognition field is one of the most challenging fields that has faced scientists for a long time. The complete solution is still far from reach. The efforts are concentrated with huge funds from the companies to different related and supportive approaches to reach the final goal. Then, apply it to the enormous applications that are still waiting for the successful speech recognisers that are free from the constraints of speakers, vocabularies or environment. This task is not an easy one due to the interdisciplinary nature of the problem and as it requires speech perception to be implied in the recogniser (Speech Understanding Systems) which in turn point strongly to the use of intelligence within the systems.

The bare techniques of recognisers (without intelligence) are following wide varieties of approaches with different claims of success by each group of authors who put their faith in their favourite way. However, the sole technique that gains the acceptance of the researchers to be the state of the art is the Hidden Markov Model (HMM) technique. HMM is agreed to be the most promising one. It might be used successfully with other techniques to improve the performance, such as hybridising the HMM with Artificial Neural Networks (ANN) algorithms. This does not mean that the HMM is pure from approximations that are far from reality, such as the successive observations independence, but the results and potential of this algorithm is reliable. The modifications on HMM take the burden of releasing it from these poorly representative approximations hoping for better results.

In this report we are going to describe the backbone of the HMM technique with the main outlines for successful implementation. The representation and implementation of HMM varies in one way or another but the main idea is the same as well as the

results and computation costs, it is a matter of preferences to choose one. Our preference here is that adopted by Ferguson [1] and Rabiner et al. [2]-[5].

In this report we will describe the Markov Chain, and then investigate a very popular model in the speech recognition field (the Left-Right HMM Topology). The mathematical formulations needed to be implemented will be fully explained as they are crucial in building the HMM. The prominent factors in the design will also be discussed. Finally we conclude this report by some experimental results to see the practical outcomes of the implemented model.

2. Markov Chains

The HMM algorithms are basically inspired by the more than 90 year old mathematical models known as Markov Chains. To understand the behaviour of the Markov Chain it is advisable to start with a simple real life example.

Let us consider a simple weather forecast problem and try to emulate a model that can predict tomorrow's weather based on today's conditions. In this example we have three stationary all day weather, which could be sunny (S), cloudy (C), or Rainy (R). From the history of the weather of the town under investigation we have the following table (Table-1) of probabilities of having certain weather tomorrow and being in certain condition today:

		Tomorrow		
		Sunny(S)	Cloudy(C)	Rainy(R)
Today	Sunny(S)	.7	.2	.1
	Cloudy(C)	.05	.8	.15
	Rainy(R)	.15	.25	.6

Table-1 Weather expectation probabilities.

In this case what we are looking for is the weather conditional probability P(Tomorrow | Today). We realise that tomorrow's weather depends on today's conditions as well as the previous several days, but we accept the assumption that tomorrow's weather depends only on today's condition as it is in consistency with the first order Markov chain. This assumption greatly simplifies the problem of formu-

lating the model even in the actual speech recognition case and we will use it when we come to tackle the real problem.

We refer to the weather conditions by state q that are sampled at instant t and the problem is to find the probability of weather condition of tomorrow given today's condition $P(q_{t+1} \mid q_t)$.

The approximation for n instants history is:

$$P(q_{t+1} \mid q_t, q_{t-1}, q_{t-2}, \ldots, q_{t-n}) \approx P(q_{t+1} \mid q_t).$$

This is the first order Markov chain as the history is considered to be one instant only. The finite state diagram of the weather probabilistic table is shown in Fig.(1).

Let us now ask this question: Given today is sunny (S) what is the probability that the next following five days are S, C, C, R and S, given the above model?

The answer resides in the following formula using first order Markov chain:

$$P(q_1=S, q_2=S, q_3=C, q_4=C, q_5=R, q_6=S) =$$

$$P(S)\cdot P(q_2=S \mid q_1=S)\cdot P(q_3=C \mid q_2=S)\cdot P(q_4=C \mid q_3=C)\cdot P(q_5=R \mid q_4=C)\cdot P(q_6=S \mid q_5=R)$$

$$= 1 \times 0.7 \times 0.2 \times 0.8 \times 0.15 \times 0.15$$

$$= 0.00252$$

The initial probability P(S) = 1, as it is assumed that today is sunny.

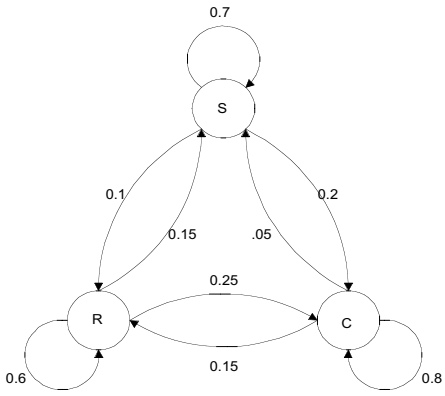


Fig.(1) Finite state representation of the weather forecast problem.

3. Hidden Markov Model (HMM)

In the particular problem presented in the previous section, the states were observable and they represented the weather conditions (S, C, R). They also represented the observation sequence. This kind of model formulation is very limited due to the need of

observable state sequence, which is unknown in most problems. The more general case is by considering the state sequence to be hidden (unobservable) and the observations are probabilistic functions of the state. This notion implies the double stochastic process. More precisely, the HMM is a probabilistic pattern matching technique in which the observations are considered to be the output of stochastic process and consists of an underlying Markov chain. It has two components: a finite state Markov chain and a finite set of output probability distributions. The first fruitful investigation of HMM was done by Baum et al. [6]–[8] in the late 60s and early 70s. The technique was applied to the speech recognition field by Baker [9].

To understand the HMM we prefer to start with simple example inspired from that given by Rabiner et al. [3]. Assume that we have two persons, one doing an experiment and the other is an outside observer. Let us consider that we have N urns (states) numbered from S_1 to S_N and in each urn there are M coloured balls (observations) distributed in different proportions. Also we have a black bag belongs to each urn, each bag contains 100 counters numbered by three numbers. These numbers are the current urn number S_i and the following two urns numbers S_{i+1} and S_{i+2} in probability proportions of .8, .15, and .05 respectively. The counters of the bag belonging to the urn just before the last are carrying one of two numbers only; S_{N-1} and S_N in probabilities of .9 and .1 respectively. We assume that the starting urn (state) is always urn1 (S_1) and we end up in urnN (S_N). The last urn needs no bag as we suggest to stay there when we reach it till the end of the experiment. We start the experiment at time t =1 by drawing a ball from urn1 and register the colour then return it back to the urn. Then draw a counter from the corresponding urn bag. The expected possible numbers on the counters are: 1 (stay in urn1), or 2 (move to the next urn), or 3 (jump to the third urn). We continue with the same procedure of drawing a counter then a ball from the corresponding urn and registering the ball colours till we reach state N and till the end of the experiment at instant T.

The outcome of this experiment is a series of coloured balls (observations) which could be considered as a sequence of events governed by the probability distribution of the balls inside each urn and by the counters existing in each bag. The outside observer has no idea about which urn a ball at any instant has drawn from (hidden states), what he knows is only the observation sequence of the coloured balls (observations). Several things could be concluded from this experiment:

1. The starting urn is always (S_1).

2. The urn which has been left can not be visited again (i.e. moving from left to right direction).
3. Movements are either by one or two urns to the right.
4. The last urn visited is always urnN (S_N).

A chain of 5 urns (states) is shown in Fig.(2).

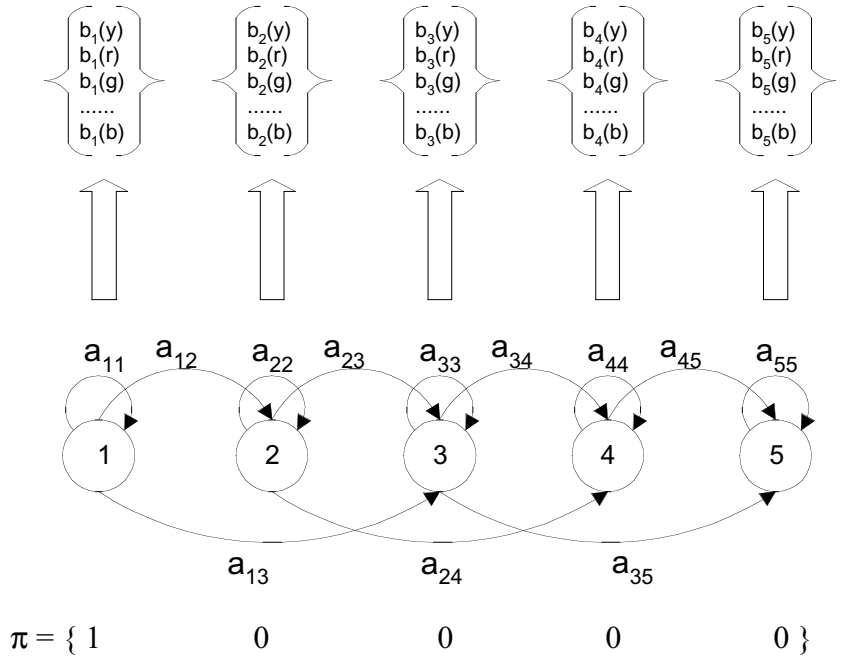


Fig.(2) States chain of the urn experiment using 5 urns. Each numbered circle represents a state and the arrows shows the states' flow during the whole process.

Fig.(2) shows the notations which we intend to use for the rest of the report and they are defined as follows:

a_{ij} represents the probability of state transition (probability of being in state S_j given state S_i)

$$a_{ij} = P(q_{t+1}=S_j \mid q_t=S_i) \quad (1)$$

$b_j(w_k)$ is the w_k symbol (ball colour) probability distribution in a state S_j

w is the alphabet and k is the number of symbols in this alphabet.

$\pi = \{1\ 0\ 0\ 0\ 0\}$ is the initial state distribution.

In this special case of states chain topology $\pi_i = P(q_1 = S_i) = \begin{cases} 1 & \text{for } i=1 \\ 0 & \text{for } 1<i\le N \end{cases}$

The model is completely defined by these three sets of parameters a, b, and π and the model of N states and M observations can be referred to by:

$$\lambda = (A, B, \pi) \quad (2)$$

where $A = \{a_{ij}\}$, $B = \{b_j(w_k)\}$ $1 \le i, j \le N$ and $1 \le k \le M$.

The model that we have just described is a special type of HMM which is normally used in speech recognition. It is called Left-Right HMM as derived from its way of behaviour and its topology (moving from left to right during state transition). The reason for using the L-R topology of HMM is due to its inherent structure that can model the temporal flow of speech signals over time.

It might be not very obvious how the HMM is related to the speech signal modelling [10]. This could be envisaged by looking at the speech production mechanism. Speech is produced by the slow movements of the articulatory organ. The speech articulators taking up a sequence of different positions and consequently producing the stream of sounds that form the speech signal. Each articulatory position could be represented by a state of different and varying duration. Accordingly, the transition between different articulatory positions (states) can be represented by $A = \{a_{ij}\}$. The observations in this case are the sounds produced in each position and due to the variations in the evolution of each sound this can be also represented by a probabilistic function $B = \{b_j(w_k)\}$.

The correspondence between the model parameters and what they represent in the speech signal is not unique and could be viewed differently. The important thing is to envisage the physical meanings of the states and observations in each view.

4. HMM Constraints for Speech Recognition Systems

HMM could have different constraints depending on the nature of the problem that wanted to be modelled. The main constraints needed in the implementation of speech recognisers can be summarised in the following assumptions [11]:

1 – First order Markov chain:

In this assumption the probability of transition to a state depends only on the current state:

$$P(q_{t+1}=S_j \mid q_t=S_i, q_{t-1}=S_k, q_{t-2}=S_w, \ldots, q_{t-n}=S_z) \approx P(q_{t+1}=S_j \mid q_t=S_i) \quad (3)$$

2 – Stationary states' transition:

This assumption testifies that the states' transition are time independent, and accordingly we will have:

$$a_{ij} = P(q_{t+1}=S_j \mid q_t=S_i) \quad \text{for all } t \tag{4}$$

3 – Observations independence:

This assumption presumes that the observations come out within certain state depend only on the underlying Markov chain of the states, without considering the effect of the other observations. Although this assumption is a poor one and deviates from reality, it works fine in modelling speech signal.

This assumption implies that:

$$P(O_t \mid O_{t-1}, O_{t-2}, \ldots, O_{t-p}, q_t, q_{t-1}, q_{t-2}, \ldots, q_{t-p}) = P(O_t \mid q_t, q_{t-1}, q_{t-2}, \ldots, q_{t-p}) \tag{5}$$

where p represents the considered history of the observation sequence.

Then we will have:

$$b_j(O_t) = P(O_t \mid q_t=j) \tag{6}$$

4 – Left-Right topology constraint:

$$a_{ij} = 0 \quad \text{for all} \quad i+2 < j < i \tag{7}$$

$$\pi_i = P(q_1 = S_i) = \begin{cases} 1 & \text{for} \quad i=1 \\ 0 & \text{for} \quad 1<i\leq N \end{cases} \tag{8}$$

(i.e., $\pi = \{1 \ 0 \ \ldots \ 0\}$)

5 – Probability constraints:

Our problem is dealing with probabilities then we have the following extra constraints:

$$\sum_{j=1}^{N} a_{ij} = 1 \tag{9}$$

$$\sum_{j=1}^{N} \pi_j = 1 \tag{10}$$

$$\int_O b_i(O_t)\,dO = 1 \tag{11}$$

If the observations are discrete then the last integration will be a summation.

5. The principal cases of HMM

There are three main cases to be dealt with to formulate a successful HMM. These are:

Case 1: Evaluation

Given:

- a model $\lambda = (A, B, \pi)$ ready to be used.
- testing observation sequence $O = O_1, O_2, O_3, \ldots, O_{T-1}, O_T$.

Action:

- compute $P(O \mid \lambda)$; the probability of observation sequence given the model.

Case 2: Decoding

Given:

- a model $\lambda = (A, B, \pi)$ ready to be used.
- testing or training observation sequence $O = O_1, O_2, O_3, \ldots, O_{T-1}, O_T$.

Action:

- track the optimum state sequence $Q = q_1, q_2, q_3, \ldots, q_{T-1}, q_T$ that most likely produces the given observations, using the given model.

Case 3: Training

Given:

- a model $\lambda = (A, B, \pi)$ ready to be used.
- training observation sequence $O^k = O_1^k, O_2^k, O_3^k, \ldots, O_{T-1}^k, O_T^k$ where k is the number of examples for training the model.

Action:

- Tune the model parameters to maximise $P(O \mid \lambda)$.

Case 1 is an evaluation procedure as we are seeking to find the probability of producing given observation O by a given model λ. This could be used to find out the best model among many who produces the given observation.

Case 2 is a decoding procedure to detect or unhide the state sequence of a given observation. The observations could be training examples if we want to study the behaviour of each state from different aspects, such as states' duration or spectral characteristics of each state. Some techniques utilise the state duration in their

evaluation procedure and in this case the observation will be the test example to detect the states' duration.

Case 3 is the training procedure to optimise the model parameters to obtain the best model that represents certain set of observations belonging to one spoken entity.

The way is paved now to tackle an important goal of our task, namely derivation of the mathematical formulas to the three previous cases.

5–1. Case 1 Formulation

Let us take a simple case then generalise to the complete one. Consider that we have 3 states and 5 observations in a process and we want to find P(O | λ). To explain the whole flow of the process the trellis diagram of Fig.(3) is of big help. The state at each instant is represented by a small circle, and the arrows represent the state transitions.

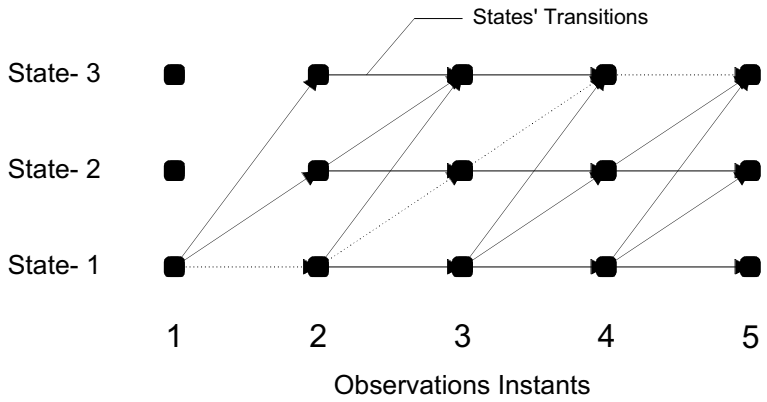


Fig.(3) Trellis Diagram of 3 States, and 5 Instants L-R Model

From Fig.(3) we can see all the possibilities that the events might take during the whole process. The dotted lines shows one possibility in which the $P(O^1 \mid Q^1, \lambda)$ is:

$$P(O^1 \mid Q^1, \lambda) = \prod_{t=1}^{5} P(O_t^1 \mid q_t^1 \tag{12}$$

$$P(O^1 \mid Q^1, \lambda) = b_{q1}^1 . b_{q2}^1 . b_{q3}^1 . b_{q4}^1 . b_{q4}^1 = b_1^1 . b_1^1 . b_2^1 . b_3^1 . b_3^1 \tag{12a}$$

$$P(Q^1 \mid \lambda) = \pi_1 \sum_{t=1}^{T-1} a_{q_t q_{t+1}} \tag{13}$$

$$P(Q^1 \mid \lambda) = \pi_1 . a_{q1q2} . a_{q2q3} . a_{q3q4} . a_{q4q5} = \pi_1 . a_{11} . a_{12} . a_{23} . a_{33} \tag{13a}$$

$$P(O^1, Q^1 \mid \lambda) = P(O^1 \mid Q^1, \lambda) P(Q^1 \mid \lambda) \tag{14}$$

$$P(O^1, Q^1 \mid \lambda) = b_1^1 . b_1^1 . b_2^1 . b_3^1 . b_3^1 . \pi_1 . a_{11} . a_{12} . a_{23} . a_{33} \tag{14a}$$

This procedure has to be done for all possible states' sequence (paths). The super-scripts of O and Q indicate the possibility number. Then the probabilities of all the paths has to be summed to get the probability of how likely the model produces the given observation sequence.

$$P(O/\lambda) = \sum_{i=1}^{p} P(O^i, Q^i \mid \lambda) \tag{15}$$

$$P(O/\lambda) = \sum_{i=1}^{p} P(O^i \mid Q^i, \lambda) P(Q^i \mid \lambda) \tag{15a}$$

where p is the number of states' paths possibilities.

The total number of possibilities increases exponentially with the increasing number of states and observation instances. The Left-Right topology is substantially reducing the number of possible paths over the full connection topology (ergodic models in which every state could be reached from any other state at any instant).

Further reduction in the computational cost can be achieved by the Forward-Backward Procedure [12]. This technique greatly reduces the computational cost with simple iterative mathematical formulas. Actually it is a compound procedure composed of forward procedure and backward procedure. In the evaluation case we only need one of them and the forward procedure will be our preference.

5-1.1 Forward Procedure

Initially consider a new forward probability variable $\alpha_t(i)$, at instance t and state i, has the following formula:

$$\alpha_t(i) = P(O_1, O_2, O_3, \ldots, O_t, q_t = S_i \mid \lambda) \tag{16}$$

This probability function could be solved for N states and T observations iteratively:

1 – Initialisation

$$\alpha_1(i) = \pi_i . b_i(O_1) \quad 1 \le i \le N \tag{17}$$

2 – Induction

$$\alpha_{t+1}(j) = \left[\sum_{i=1}^{N} \alpha_t(i) a_{ij} \right] b_j(O_{t+1}) \quad \begin{matrix} 1 \le t \le T-1 \\ 1 \le j \le N \end{matrix} \tag{18}$$

Fig.(4) shows the induction step graphically. It is clear from this figure how state S_j at instance t+1 reached from N possible states at instance t.

3 – Termination

$$P(O \mid \lambda) = \sum_{i=1}^{N} \alpha_T(i) \tag{19}$$

This stage is just a sum of all the values of the probability function $\alpha_T(i)$ over all the states at instant T. This sum will represent the probability of the given observations to be driven from the given model. That is how likely the given model produces the given observations. The proof of the termination formula will be given later on.

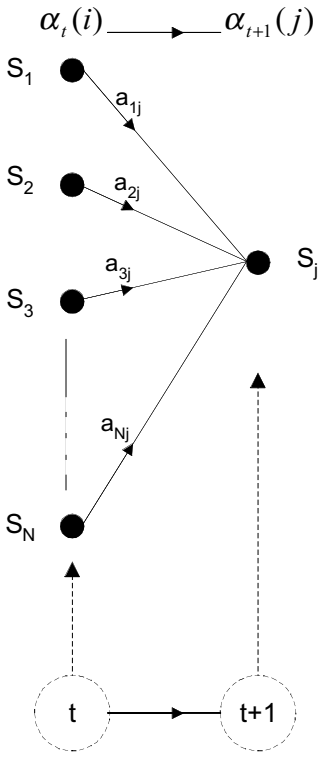


Fig.(4) Forward Probability Function Rpresentation

5-1.2 Backward Procedure

This procedure is similar to the forward procedure but it takes into consideration the state flow as if in backward direction from the last observation entity, instant T, till the first one, instant 1. That means that the access to any state will be from the states that are coming just after that state in time and as shown in Fig.(5).

To formulate this approach let us consider the backward probability function $\beta_t(i)$ and define as:

$$\beta_t(i) = P(O_{t+1}, O_{t+2}, \ldots, O_T \mid q_t = S_i, \lambda). \tag{20}$$

In analogy to the forward procedure we can solve for $\beta_t(i)$ in the following two steps:

1 – Initialisation:

$$\beta_T(i) = 1, \quad 1 \le i \le N. \tag{21}$$

These initial values for β's of all states at instant T are arbitrarily selected.

2 – Induction:

$$\beta_t(i) = \sum_{j=1}^{N} a_{ij}.b_j(O_{t+1}).\beta_{t+1}(j), \quad t = T-1, T-2, \ldots, 1, \quad 1 \le i \le N \tag{22}$$

Equation (22) can be well understood with help of Fig.(5). We are still looking from left to right in calculating the partial probability function (from t to T). However, at each instant we consider that we have β at *t+1* and we need to calculate it at time *t*; as if we are moving backward in time.

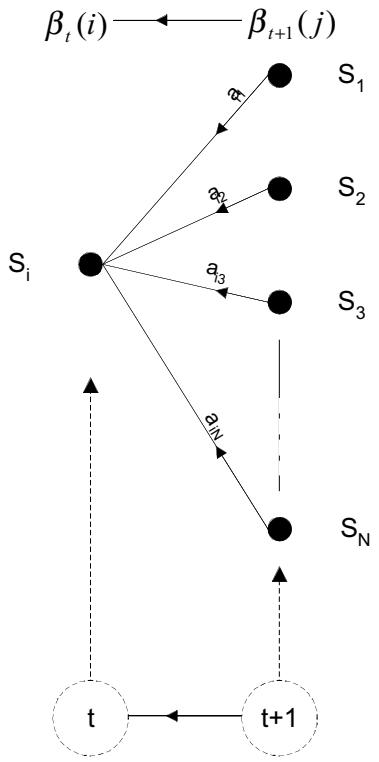


Fig.(5) Backward Probability Function Rpresentation

5-1.3 Computing P(O | λ) from Forward and Backward Probability Functions

The probability function of the model P(O | λ) can be computed from both α and β functions. Fig.(6) demonstrates this computation graphically. At instant t, the event of being in state q_i and moving to state q_j at instance t+1 is calculated by $\alpha_t(i)$ which accounts for the path termination in state q_i. The transition to state q_j is weighted by the product $a_{ij} \cdot b_j(O_{t+1})$. At instance t+1 the event of observation sequence to the instant T starting from state S_j, while being at state S_i during instant t, is represented by the backward probability function $\beta_{t+1}(j)$.

Then P(O | λ) is directly concluded to be:

$$P(O \mid \lambda) = \sum_{i=1}^{N} \sum_{j=1}^{N} \alpha_t(i).a_{ij}.b_j(O_{t+1}).\beta_{t+1}(j) \tag{23}$$

Substitute (22) in (23) to get:

$$P(O \mid \lambda) = \sum_{i=1}^{N} \alpha_t(i)\beta_t(i) \tag{23a}$$

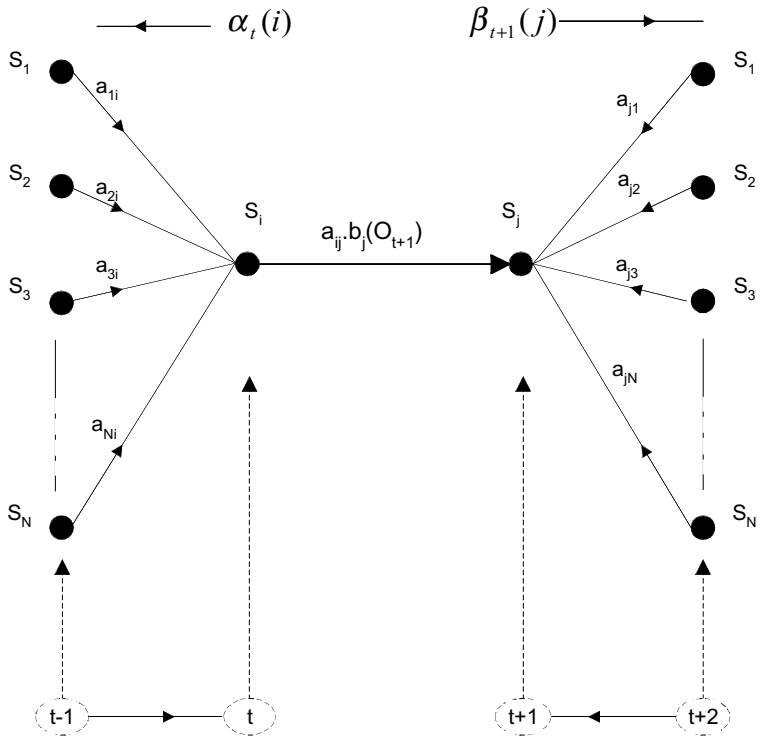


Fig.(6) Forward - Backward Probability Functions to find P(O/λ)

5-1.4 Proof of Termination Formula in Forward Probability Function

From (18) we have:

$$\alpha_{t+1}(j) = \left[\sum_{i=1}^{N} \alpha_t(i) a_{ij}\right] b_j(O_{t+1}) \quad \begin{matrix} 1 \le t \le T-1 \\ 1 \le j \le N \end{matrix} \tag{18}$$

Let t = T-1 and substitute in 18 to get:

$$\alpha_T(j) = \left[\sum_{i=1}^{N} \alpha_{T-1}(i).a_{ij}\right].b_j(O_T) \tag{18a}$$

$$\alpha_T(j) = \sum_{i=1}^{N} \alpha_{T-1}(i).a_{ij}.b_j(O_T) \tag{18b}$$

From (23) we have:

$$P(O \mid \lambda) = \sum_{i=1}^{N}\sum_{j=1}^{N} \alpha_t(i).a_{ij}.b_j(O_{t+1}).\beta_{t+1}(j) \tag{23}$$

Substitute t = T-1 in (23) to get:

$$P(O \mid \lambda) = \sum_{i=1}^{N}\sum_{j=1}^{N} \alpha_{T-1}(i).a_{ij}.b_j(O_T).\beta_T(j) \tag{23b}$$

We have from (21)

$$\beta_T(i) = 1, \quad 1 \le i \le N. \tag{21}$$

Substitute for $\beta_T(i)$ in (23b) and rearrange the equation to get:

$$P(O \mid \lambda) = \sum_{j=1}^{N}\left[\sum_{i=1}^{N} \alpha_{T-1}(i).a_{ij}.b_j(O_T)\right] \tag{24}$$

The term inside the square brackets is the same as that in (18b), substitute it and you will get the final needed formula:

$$P(O \mid \lambda) = \sum_{i=1}^{N} \alpha_T(i) \tag{19}$$

5-2. Case 2 Formulation

This case deals with the uncovering the hidden states of the model given the observation sequence and the model. This means that we have to find the optimal state sequence $Q = (q_1, q_2, q_3, \ldots, q_{T-1}, q_T)$ associated with the given observation sequence $O = (O_1, O_2, O_3, \ldots, O_{T-1}, O_T)$ presented to the model $\lambda = (A, B, \pi)$. The criteria of optimality here is to search for a single best state sequence through modified dynamic

programming technique called Viterbi Algorithm [13]. We need to maximise P(Q | O,λ) to detect the best state sequence. This could be achieved via maximising the joint probability function P(Q,O | λ) using the Bayesian Rule which states that:

$$P(Q \mid O,\lambda) = \frac{P(Q,O \mid \lambda)}{P(O \mid \lambda)} \tag{25}$$

The denominator has nothing to share in maximising *P(Q | O,λ)* as it does not include the state sequence factor *Q*. To go through the Viterbi Algorithm method let us define the probability quantity $\delta_t(i)$ which represents the maximum probability along the best probable state sequence path of a given observation sequence after t instants and being in state *i.* This quantity can be mathematically represented by:

$$\delta_t(i) = \max_{q_1,q_2,\ldots,q_{t-1}} P[q_1 q_2 \ldots q_{t-1}, q_t = S_i, O_1 O_2 \ldots O_t \mid \lambda] \tag{26}$$

The best state sequence can be backtracked by another function $\psi_t(j)$. The complete algorithm can be described by the following steps:

Step 1: Initialisation

$$\delta_1(i) = \pi_i b_i(O_1), \quad 1 \le \mathrm{i} \le \mathrm{N} \tag{27}$$

$$\psi_1(i) = 0 \tag{28}$$

Step 2: Recursion

$$\delta_t(j) = \max_{1 \le i \le N}[\delta_{t-1}(i)a_{ij}]b_j(O_t), \quad 2 \le \mathrm{t} \le \mathrm{T}, \quad 1 \le \mathrm{j} \le \mathrm{N} \tag{29}$$

$$\psi_t(j) = \arg\max_{1 \le i \le N}[\delta_{t-1}(i)a_{ij}], \quad 2 \le \mathrm{t} \le \mathrm{T}, \quad 1 \le \mathrm{j} \le \mathrm{N} \tag{30}$$

Step 3: Termination

$$P^* = \max_{1 \le i \le N}[\delta_T(i)] \tag{31}$$

$$q_T^* = \arg\max_{1 \le \mathrm{i} \le \mathrm{N}}[\delta_{\mathrm{T}}(i)] \tag{32}$$

Step 4: Backtracking

$$q_t^* = \psi_{t+1}(q_{t+1}^*), \quad \mathrm{T}-1 \ge \mathrm{t} \ge 1 \tag{33}$$

It is clear that (29) of Viterbi recursion is similar to (18) of forward induction, except the interchange of summation by maximisation. One thing could be noted here is that Viterbi Algorithm can also be used to calculate the P(O | λ) approximately by considering the use of P^* instead. This is acceptable as it gives comparable results and can be justified through the modified equation (15) to do the summation on the

most probable state sequence, which has the major weight among all the possible states' paths.

5-3. Case 3 Formulation

This case is dealing with the training issue, which is the most difficult one in all the three cases. The task of this case is to adjust the model parameters, (A, B, π), according to certain optimality criteria. There are many techniques to achieve the task of this case and we will describe here the well-known Baum-Welch Algorithm, called also Forward-Backward Algorithm. It is an iterative method to reach the local maximas of the probability function $P(O \mid \lambda)$. Each time the model parameters are adjusted to get a new model which is proved by Baum et. al. that the new model is either better or reach a critical condition at which the iteration has to be stopped as the local minima has reached. The model always converges but the global maximisation cannot be assured. Fig.(7) shows the non-linear optimisation of this problem and how the global optimality seeking is difficult to locate and greatly depending on the initial point of search.

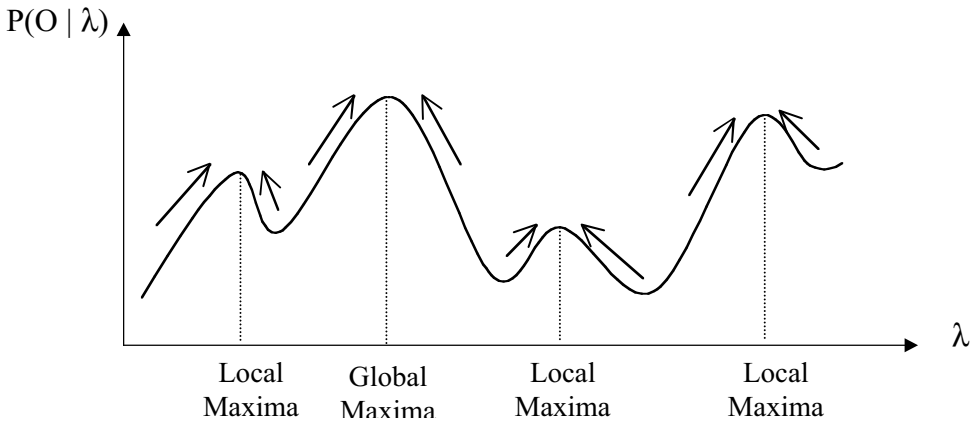


Fig.(7) Optimum Search Possibilities.

To go through the training procedure let us first define a posteriori probability function $\gamma_t(i)$, the probability of being in state i at instant t, given the observation sequence O and the model λ as:

$$\gamma_t(i) = P(q_t = S_i \mid O, \lambda) \tag{34}$$

$$\gamma_t(i) = \frac{P(O, q_t = S_i \mid \lambda)}{P(O \mid \lambda)} \tag{35}$$

Since $P(O, q_t = S_i \mid \lambda) = \alpha_t(i)\beta_t(i)$ (36)

and from (23a)

$$P(O \mid \lambda) = \sum_{i=1}^{N} \alpha_t(i)\beta_t(i) \tag{23a}$$

then $\gamma_t(i) = \dfrac{\alpha_t(i)\beta_t(i)}{\sum_{i=1}^{N} \alpha_t(i)\beta_t(i)}$ (37)

Let us define another probability function $\xi_t(i,j)$, the probability of being in state i at instant t and going to state j at instant t+1, given the model λ and the observation sequence O.

$\xi_t(i,j)$ can be mathematically defined as:

$$\xi_t(i,j) = P(q_t = S_i, q_{t+1} = S_j \mid O, \lambda) \tag{38}$$

Multiply both sides of (39) by $P(O \mid \lambda)$ to get:

$$\xi_t(i,j).P(O \mid \lambda) = P(q_t = S_i, q_{t+1} = S_j \mid O, \lambda) \cdot P(O \mid \lambda) \tag{39}$$

From Bayesian Rule

$$P(q_t = S_i, q_{t+1} = S_j \mid O, \lambda) \cdot P(O \mid \lambda) = P(O, q_t = S_i, q_{t+1} = S_j \mid \lambda) \tag{40}$$

The right-hand side of (40) can be represented by the forward α and backward β functions, with the help of Fig.(6), as follows:

$$P(O, q_t = S_i, q_{t+1} = S_j \mid \lambda) = \alpha_t(i) a_{ij} b_j(O_{t+1}) \beta_{t+1}(j) \tag{41}$$

Substitute (23a) and (41) in (39) and rearrange to get:

$$\xi_t(i,j) = \frac{\alpha_t(i) a_{ij} b_j(O_{t+1}) \beta_{t+1}(j)}{\sum_{i=1}^{N} \alpha_t(i)\beta_t(i)} \tag{42}$$

Also, from (23) we can have:

$$\xi_t(i,j) = \frac{\alpha_t(i)a_{ij}b_j(O_{t+1})\beta_{t+1}(j)}{\sum_{i=1}^{N}\sum_{j=1}^{N}\alpha_t(i)a_{ij}b_j(O_{t+1})\beta_{t+1}(j)} \tag{43}$$

The relation between $\gamma_t(i)$ and $\xi_t(i,j)$ can be easily deduced from their definitions:

$$\gamma_t(i) = \sum_{j=1}^{N}\xi_t(i,j) \tag{44}$$

Now, if $\gamma_t(i)$ is summed over all instants (excluding instant T) we get the expected number of times that state S_i has left, or the number of times this state has been visited over all instants. On the other hand if we sum $\xi_t(i,j)$ over all instants (excluding T) we will get the expected number of transitions that have been made from i to j.

From the behaviour of $\gamma_t(i)$ and $\xi_t(i,j)$ the following re-estimations of the model parameters could be deduced:

$$\begin{aligned}\hat{\pi}_i &= \text{expected number of instances the starting state is } S_i \\ &= \gamma_1(i)\end{aligned} \tag{45}$$

$$\begin{aligned}\hat{a}_{ij} &= \frac{\text{expected number of transitions from } S_i \text{ to } S_j}{\text{expected number of transitions from } S_i} \\ &= \frac{\sum_{t=1}^{T-1}\xi_t(i,j)}{\sum_{t=1}^{T-1}\gamma_t(i)} = \frac{\sum_{t=1}^{T-1}\alpha_t(i)a_{ij}b_j(O_{t+1})\beta_{t+1}(j)}{\sum_{t=1}^{T-1}\alpha_t(i)\beta_t(i)}\end{aligned} \tag{46}$$

$$\begin{aligned}\hat{b}_j(k) &= \frac{\text{expected number of instances in state } S_j \text{ and having observation } O_t = w_k}{\text{expected number instances in state } S_j} \\ &= \frac{\sum_{t=1,\,O_t=w_k}^{T}\gamma_t(j)}{\sum_{t=1}^{T}\gamma_t(j)} = \frac{\sum_{t:O_t=w_k}\alpha_t(j)\beta_t(j)}{\sum_{t=1}^{T}\alpha_t(j)\beta_t(j)}\end{aligned} \tag{47}$$

After the re-estimation of the model parameters we will have another model $\hat{\lambda}$ which is more likely, than model λ, producing observation sequence *O*. This means that:

$$P(O \mid \hat{\lambda}) > P(O \mid \lambda)$$

This process of re-estimation can be continued till no improvement in $P(O \mid \lambda)$ is reached, that is we reach local maxima.

6. Discrete Hidden Markov Model (DHMM)

The HMM modelling methods applied so far are for a process that has a discrete observation sequence. These observations could be the outcome indices of Vector Quantization technique (VQ) [14],[15]. VQ is a technique of clustering time series signals, in our case speech signals, into certain number of bins (clusters). Each bin represents the data belonging to certain population with similar (or minimum difference) spectral characteristics. The centre of gravity of each bin is assigned to a certain index and considered as representative of the cluster population in any process on the signal. The long sequence of speech samples will be represented by a stream of indices representing frames of different window lengths. Hence, VQ is considered as a process of redundancy removal, which minimises the number of bits required to identify each frame of speech signal. VQ was initially used successfully with Dynamic Time Warping (DTW) to recognise spoken words, and then proved to be successful with HMM as well. The role of VQ in HMM is to prepare discrete symbols from a finite alphabet. Each speech input will be quantized by the VQ reference bins. Each quantized input will be then considered as an observation. There are many other methods to represent the observations that are beyond the scope of this report, but a very good reference to recommend is [16].

The type of HMM that models speech signals based on VQ technique to produce the observations is called Discrete Hidden Markov Model (DHMM). It is an efficient and reliable technique that has comparable results to the more computational DTW technique. In addition the phones, phonemes, and subwords could be modelled easily with DHMM while it is very difficult with DTW as the latter needs to detect the segments boundary for comparison. However, VQ is responsible for los-

ing some information from the speech signal even when we try to increase the codewords. This loss is due to the quantization error (distortion). This distortion can be reduced by increasing the number of codewords in the codebook but cannot be eliminated.

7. Continuous Hidden Markov Model (CHMM)

It is more sophisticated methodology to develop improved HMM models of the speech signals. This method needs more memory than DHMM to represent the model parameters but does not suffer from the distortion problem. On the other hand it needs more deliberate techniques to initialise the model as it might diverge easily with randomly selected initial parameters.

In CHMM the model parameters are also π, A, and B, but they are represented differently. The probability density function (pdf) of certain observations O being in a state is considered to be of Gaussian Distribution (other distributions also valid). Let us consider it to be $b_i(O)$ and has the following general form:

$$b_i(O) = \sum_{m=1}^{M} c_{im} \aleph(O; \mu_{im}, U_{im}), \quad 1 \le i \le N \tag{48}$$

where:

c_{jm}: is the m-th mixture gain coefficient in state i.

$\aleph$: is the pdf distribution which is considered to be Gaussian in our case.

μ_{im}: is the mean of the m-th mixture in state i.

U_{im}: is the covariance of the m-th mixture in state i.

O: is the observation sequence of the feature vectors of dimension d.

M: is the number of mixtures used.

N: is the number of states.

The following constraints have to be fulfilled to ensure the consistency of the model parameters estimation.

$$\sum_{m=1}^{M} c_{im} = 1, \quad 1 \le i \le N \tag{49a}$$

$$c_{im} \ge 0, \quad 1 \le i \le \mathrm{N}, \quad 1 \le m \le M \tag{49b}$$

These constraints will lead to proper pdf normalisation, that is

$$\int_{-\infty}^{\infty} b_i(O)dO = 1, \quad \text{for } 1 \le i \le N \tag{50}$$

The pdf of the observations will be of the form:

$$b_i(O) = \frac{1}{(2\pi)^{d/2}\sqrt{|U_i|}} e^{-\frac{1}{2}(O-\mu_i)'U_i^{-1}(O-\mu_i)} \tag{51}$$

where prime (′) superscript here is referring to the transpose of matrix.

The covariance matrix in (51) could be simplified by using a diagonal matrix with elements representing the variance of each mixture. This approximation greatly reduces the computational cost in spite of the necessity to increase the number of mixtures to make it work better.

The reestimation formulas in multimixture continuous density HMM will be as follows:

$$\hat{c}_{im} = \frac{\text{expected instances of being in state i and mixture m}}{\text{expected instances of being in state i}}$$

$$\hat{c}_{im} = \frac{\sum_{t=1}^{T} \gamma_t(i,m)}{\sum_{t=1}^{T}\sum_{m=1}^{M} \gamma_t(i,m)} \tag{52}$$

$$\hat{\mu}_{im} = \frac{\sum_{t=1}^{T} \gamma_t(i,m) \cdot O_t}{\sum_{t=1}^{T} \gamma_t(i,m)} \tag{53}$$

$$\hat{U}_{im} = \frac{\sum_{t=1}^{T} \gamma_t(i,m) \cdot (O_t - \mu_{im})(O_t - \mu_{im})'}{\sum_{t=1}^{T} \gamma_t(i,m)} \tag{54}$$

where $\gamma_t(i,m)$ is the probability of being in state i with m-th mixture at instant t. It is the same as $\gamma_t(i)$ when m=1.

The following equation represents the modified version of (38) to make it suitable for the multimixture case:

$$\gamma_t(i,m) = \left[\frac{\alpha_t(i)\beta_t(i)}{\sum_{i=1}^{N} \alpha_t(i)\beta_t(i)} \right] \left[\frac{\mathrm{c}_{\mathrm{im}} \aleph(O_t; \mu_{im}, U_{im})}{\sum_{m=1}^{\mathrm{M}} \mathrm{c}_{\mathrm{im}} \aleph(O_t; \mu_{im}, U_{im})} \right] \tag{55}$$

For the initial state and the state transition probability distributions they are the same as for DHMM as in (45) and (46).

8. Mixture Density Components Estimation using Maximum Likelihood (ML)

The ML estimation is an optimisation technique that can be used efficiently in estimating the different components of multimixture models. We are not going through the mathematical derivations of the ML but only describe the method to be used in our task.

Let us first make some definitions:

$b_i(O_t)$: probability of being in state i given observation sequence O_t.

c_{im}: probability of being in state i with mixture m (gain coefficient).

$b_{im}(O_t)$: probability of being in state i with mixture m and given O_t.

$\Phi(w_{im}|O_t)$: probability function of being in a mixture class w_{im} given O_t in state i.

T_i: is the total number of observations in state i.

T_{im}: is the number of observations in state i with mixture m.

N: number of states.

M: number of mixtures in each state.

Now we are ready to implement the algorithm through applying the following steps:

1. Take several versions of observations of a certain word, say digit zero, spoken several times by many speakers.
2. Apply Viterbi algorithm to detect the states of each version of the spoken word.

3. Put the whole observations belonging to each state from all the versions of the spoken word into separate cells. Now we have N cells and each one represents the population of a certain state derived from several observation sequences of the same word.
4. Apply vector quantization technique to split the population of each cell into M mixtures and getting w_M classes within each state.
5. Use the well-known statistical methods to find the mean μ_{im} and the covariance U_{im} of each class. The gain factor c_{im} can be calculated by:

$$c_{im} = \frac{\text{number of observations being in state i and mixture m}}{\text{total number of observations in state i}} \tag{56}$$

6. Calculate $\Phi(w_{im} \mid O_t)$ from the following formula:

$$\Phi(w_{im} \mid O_t) = c_{im} \cdot \frac{b_{im}(O_t)}{b_i(O_t)} \tag{57}$$

7. Find the next estimate of $\hat{c}_{im}, \hat{\mu}_{im}$ and $\hat{U}_{im}$ from the formulas given by ML:

$$\hat{c}_{im} = \frac{1}{T_i} \sum_{i=1}^{T_i} \Phi(w_{im} \mid O_t) \tag{58}$$

$$\hat{\mu}_{im} = \frac{1}{T_{im}} \sum_{i=1}^{T_i} \Phi(w_{im} \mid O_t) \cdot O_t \tag{59}$$

$$\hat{U}_{im} = \frac{1}{T_{i\,m}} \sum_{i=1}^{T_i} \Phi(w_{im} \mid O_t) \cdot (O_t - \hat{\mu}_{im})(O_t - \hat{\mu}_{im})' \tag{60}$$

$$\hat{b}_{im}(O_t) = \sum_{m=1}^{M} \hat{c}_{im} \aleph(O; \hat{\mu}_{im}, \hat{U}_{im}), \quad 1 \le i \le N \tag{61}$$

$$\hat{b}_i(O_t) = \sum_{m=1}^{M} \hat{c}_{im} \hat{b}_{im}(O_t) \tag{62}$$

8. Compute the next estimate of using the formula:

$$\hat{\Phi}(w_{im} \mid O_t) = \frac{\hat{c}_{im}\, \hat{b}_{im}(O_t)}{\sum_{n=1}^{M} \hat{c}_{in}\, \hat{b}_{in}(O_t)} \tag{63}$$

9. IF $|\Phi(w_{im} \mid O_t) - \hat{\Phi}(w_{im} \mid O_t)| \leq \varepsilon$ THEN END

ELSE Make the new value of $\Phi(w_{im} \mid O_t)$ equal the newly predicted one.

$$\Phi(w_{im} \mid O_t) = \hat{\Phi}(w_{im} \mid O_t) \tag{64}$$

GO TO STEP 7.

where ε is a very small threshold value.

9. Implementation Factors

There are several factors that may in one way or another have effects on the implemented model. We are going to describe the more important factors and how to reduce their effects.

9-1. Scaling Factor

The scaling factor is a major issue in implementing the HMM because of the underflow that may easily occur when calculating the probability function $P(O \mid \lambda)$. This is due to the long sequence of multiplications of less than one-value probability functions. For instance in using the forward procedure to calculate $\alpha_i(t)$ in (18) we can see easily how many multiplications of probability functions we have to make to calculate any spoken entities.

The straightforward technique of scaling is started by defining the scaling coefficient c(t)[2]:

$$c(t) = \frac{1}{\sum_{i=1}^{N} \alpha_t(i)} \tag{65}$$

Now let us compute $\alpha_i(t)$ from (18) and then multiply it by *c(t).* This will lead to:

$$\sum_{i=1}^{N} c(t) \cdot \alpha_t(i) = 1, \quad 1 \leq t \leq T \tag{66}$$

The same thing can be done with $\beta_t(i)$ and form the product $c(t)\cdot\beta_t(i)$. The re-estimation formula of (46) can be rewritten with scaling to become:

$$\hat{a}(i,j) = \frac{\sum_{t=1}^{T-1} C_t \alpha_t(i) a_{ij} b_j(O_{t+1}) \beta_{t+1}(j) D_{t+1}}{\sum_{t=1}^{T-1}\sum_{r=1}^{N} C_t \alpha_t(i) a_{ir} b_r(O_{t+1}) \beta_{t+1}(r) D_{t+1}} \tag{67}$$

$$\text{where } C_t = \prod_{\tau=1}^{t} c_\tau \tag{68}$$

$$\text{and } D_t = \prod_{\tau=t}^{T} c_\tau \tag{68a}$$

The numerator and the denominator of (67) consist of the product $C_t D_{t+1} = \prod_{\tau=1}^{T} c_\tau$, which can be factored out and retain the original equation of (46). This scaling technique can also be applied successfully to (47).

The scaling coefficients can be used to find *log* $P(\mathrm{O} \mid \lambda)$ by the following method:

Consider that we have c_t for t=1,2,3, …,T and we obtained C_T from (68), then from (65) we will get:

$$C_T = \prod_{t=1}^{T} c_t = \left[\sum_{i=1}^{N} \alpha_T(i)\right]^{-1} \tag{69}$$

Using (19) we will have:

$$C_T = \prod_{t=1}^{T} c_t = \frac{1}{P(O \mid \lambda)} \tag{70}$$

Take the log of the last two terms:

$$\log\left(\prod_{t=1}^{T} c_t\right) = -\log[P(O \mid \lambda] \tag{71}$$

By using log properties we can obtain:

$$\log[P(O \mid \lambda] = -\sum_{t=1}^{T} \log(c_t) \tag{72}$$

Equation (72) shows that log*(P(O | λ)* can be computed but not *P(O | λ)* as the latter will be out of the dynamic range of the computer.

Viterbi Algorithm also shows itself here again to be a successful technique in calculating log*(P(O | λ)* even without bothering about the scaling problem.

To follow Viterbi Algorithm let us assume that:

$$\phi_t(i) = \log[\pi_i b_i(O_1)] \tag{73}$$

Take the log of both sides of (29) and use (72) to get:

$$\phi_t(i) = \max_{1 \le i \le N}[\phi_{t-1}(i) + \log(a_{ij})] + \log[b_j(O_t)] \tag{74}$$

Now $\log[P(O \mid \lambda)] = \max_{1 \le i \le N}[\phi_T(i)]$.

9-2. Multiple Observation Sequence factor [2]

The main disadvantage of the Left-Right topology of HMM is that the observations cannot be concatenated into one string and submitted to the model for training. This is due to the one direction left-right move and once a state is left we can not go back to it. Accordingly the model will get stuck in the last state after passing the first observation sequence and no modelling is possible for the other sequences.

The model has to be modified to accept multiple submissions to allow the model to be trained by many versions of the same spoken entity.

Let us define the set of observations of the k multiples of observations of a spoken entity by:

$$O = [O^1, O^2, O^3, \ldots, O^k]$$

where

$$O^k = [O_1^k, O_2^k, O_3^k, \ldots, O_{T_k}^k]$$

The goal of HMM is to maximise P(O | λ) by adjusting the parameters of λ. In multiple observations P(O | λ) is defined by:

$$P(O \mid \lambda) = \prod_{k=1}^{K} P(O^k \mid \lambda) \tag{75}$$

in a more abstract way

$$P(O \mid \lambda) = \prod_{k=1}^{K} P_k \tag{75a}$$

The multiple observation sequence implication can be done by normalising the numerators and denominators of (46) and (47) by P_k to get:

$$\hat{a}_{ij} = \frac{\sum_{k=1}^{K} \frac{1}{P_k} \sum_{t=1}^{T_k-1} \alpha_t^k(i) a_{ij} b_j(O_{t+1}^k) \beta_{t+1}^k(j)}{\sum_{k=1}^{K} \frac{1}{P_k} \sum_{t=1}^{T_k-1} \alpha_t^k(i) \beta_t^k(i)} \tag{76}$$

$$\hat{b}_{ij}(\eta) = \frac{\sum_{k=1}^{K} \frac{1}{P_k} \sum_{\substack{t=1 \\ O_t = w_\eta}}^{T_k-1} \alpha_t^k(i) \beta_t^k(i)}{\sum_{k=1}^{K} \frac{1}{P_k} \sum_{t=1}^{T_k-1} \alpha_t^k(i) \beta_t^k(i)} \tag{77}$$

The same procedure of normalisation could be used in the case of continuous density distribution to find the parameters of the model.

9-3. Initial Model Parameters Estimate factor:

When we initially try to build an HMM model we normally have nothing but streams of observations. If we are fortunate then we have parameters from old models, which is not normally the case. To put the initial model parameters, we have to be careful as one might easily slip into divergence with bad model initialisation. The problem with discrete observations HMM is less effective as we can initialise the model parameters with random values, but taking into consideration the constraints in (9), (10), and (11). In continuous density HMM (CHMM) the problem is more serious and the parameters should be judiciously selected to get rid of the divergence fate. Let us take the problem in Left-Right HMM Topology and suggest a safe way to follow.

The parameters that constitute any model λ are π, A, and B. For π it is straight forward and known to be always $\pi = [1\ 0\ 0\ 0\ 0 \ldots 0]$, of course this is with Left-Right topology models. For the states' transition parameters $A=[a_{ij}]$ the choice is also flexible and if we have the topology of Fig(2) then A will be the following matrix for seven states model:

$$A = \begin{bmatrix} a_{11} & a_{12} & a_{13} & 0 & 0 & 0 & 0 \\ 0 & a_{22} & a_{23} & a_{24} & 0 & 0 & 0 \\ 0 & 0 & a_{33} & a_{34} & a_{35} & 0 & 0 \\ 0 & 0 & 0 & a_{44} & a_{45} & a_{46} & 0 \\ 0 & 0 & 0 & 0 & a_{55} & a_{56} & a_{57} \\ 0 & 0 & 0 & 0 & 0 & a_{66} & a_{67} \\ 0 & 0 & 0 & 0 & 0 & 0 & a_{77} \end{bmatrix} \quad (78)$$

The values of a_{ij} can be selected as:

$a_{ii} = 0.94$, $a_{i,i+1} = 0.04$, and $a_{i,i+2} = 0.02$ for $1 \le i \le 5$

$a_{ii} = 0.97$, $a_{i,i+1} = 0.03$ for $i = 6$

$a_{ii} = 1$ for $i = 7$

These values are deduced from the fact that the observations tend to stay in their current state and have less tendency to move to the next state and more less tendency to jump the next state. After optimisation we can see that the observations wanted to stay in their current state is true, this imply that $a_{ii} > a_{i,i+1}$ and $a_{ii} > a_{i,i+2}$. However, the observations might prefer to stay in the next state or jump it, i.e. $a_{i,i+1} > a_{i,i+2}$ or $a_{i,i+1} < a_{i,i+2}$.

A more precise way is by using initial uniform segmentation of each utterance into the proposed number of states and apply the following algorithm [11]:

The suitable a_{ij} (in programming we use the notation a(i,j)) initialisation is found to be dependent on states' duration. In this case the duration D of each state is estimated by:

$$D = \frac{1}{NxK} \sum_{n=1}^{K} | x_n | \quad (79)$$

where: K is the number of versions of an utterance in the training set.

N is the number of states

$|X_n|$ is the length of the *n*th utterance in the training observations.

Then, the transition matrix elements a(i,j) are estimated by:

For $1 \le i \le N-2$

$$a(i,i) = \frac{D-1}{D} \tag{80}$$

$$a(i,i+1) = .8 \times (1 - a(i,i)) \tag{80a}$$

$$a(i,i+2) = 1 - a(i,i) - a(i,i+1) \tag{80b}$$

For $i = N-1$

$$a(i,i) = \frac{D-1}{D} \tag{80c}$$

$$a(i,i+1) = 1 - a(i,i) \tag{80d}$$

For $i = N$

$$a(i,i) = 1 \tag{80e}$$

The formula is inferred from the fact that if there are K elements of duration D in each state then there will be only one transition to the next state.

What is left now is the most problematic parameters B = $\{b_i(O_t)\}$, they have to be very well initialised. In our case we suggest the following steps:

1. Uniformly segment the utterances of each spoken entity by N states.
2. Take the mean and the covariance of each segment.
3. Consider the observations follow Gaussian density probability distribution.

After the previous suggestions for initialising the model parameters we can apply Viterbi Algorithm to extract the optimum parameters.

The technique described in this section is for unimodal (single mixture) distributions. To extend it to multimodal (multimixture) distributions the following are suggested:

1. Apply the same procedures for π, A, B used in unimodal distributions.
2. After uncovering the real state sequence from Viterbi Algorithm, aggregate the observations, of all the versions of the spoken entity, belonging to each state in separate cells.
3. Use Vector quantization technique to cluster each cell into several mixtures.

4. Optimise the clustering by any known statistical technique; such as maximum likelihood or expectation maximisation.
5. Find the mean and the covariance of each cluster (mixture)

The model is now complete.

9-4. Number of States Factor

There is one thing left which has to be decided from the initial instant of designing the model. It is the optimum number of states needed to model the problem. There is no straightforward answer to this requirement. The number of states is decided empirically depending on the nature of the problem. Sometimes previous experience about the problem is necessary or one has to suggest different numbers of states then select the one that gives the best results. Also, if we could define the physical meaning of the states we can limit the number of states. In an isolated words recogniser the number of states are suggested to be between 4 and 12. This is justified by assuming that the states are representing the phonemes or the phones of the utterances. In phonemes modelling the number of states are mostly assumed to be 3, as the phonemes could be segmented into initial, stable, and final states.

9–5. State Duration Incorporation

The basic HMM does not take into consideration the state duration factor in its modelling procedure. This is considered a major weakness in the model since the duration carries important information about the temporal structure of the speech signal. Our duty now is to find some useful way to include the duration within the conventional model. Let us first ask this question: What is the probability of being in a state for τ instants?

The answer lies in finding the probability density function $p_i(\tau)$ which has the definition of:

$$p_i(\tau) = P(q_1=S_i, q_2=S_i, q_3=S_i, \ldots, q_\tau=S_i, q_{\tau+1}=S_j, \ldots) \quad (81)$$

$$= \pi_i (a_{ii})^{\tau-1}(1-a_{ii}) \quad (81a)$$

Now we can calculate the expected duration in state i by the following equation:

$$\bar{\tau}_i = \sum_{\tau=1}^{\infty} \tau p_i(\tau) \tag{82}$$

Using (81a) and considering $\pi_i = 1$ we get:

$$\begin{aligned}
\bar{\tau}_i &= \sum_{\tau=1}^{\infty} \tau (a_{ii})^{\tau-1}(1-a_{ii}) \\
&= (1-a_{ii}) \sum_{\tau=1}^{\infty} \tau (a_{ii})^{\tau-1} \\
&= (1-a_{ii}) \frac{\partial}{\partial a_{ii}} \sum_{\tau=1}^{\infty} (a_{ii})^{\tau} \\
&= (1-a_{ii}) \frac{\partial}{\partial a_{ii}} \left(\frac{a_{ii}}{1-a_{ii}}\right) \\
&= \frac{1}{1-a_{ii}}
\end{aligned} \tag{83}$$

Now let us return to our first example about weather forecasting and ask the question: What is the average number of consecutive sunny, cloudy, and rainy days?

The answer is by applying (83) using the values of a_{ii} from Table-1 to get:

$$Sunny\ days = \frac{1}{1-a_{11}} = \frac{1}{1-0.7} \approx 3$$

$$Cloudy\ days = \frac{1}{1-a_{22}} = \frac{1}{1-0.8} \approx 5$$

$$Rainy\ days = \frac{1}{1-a_{33}} = \frac{1}{1-0.6} \approx 3$$

Unfortunately this duration distribution is meaningless when we try to apply it to speech recognition problems. Therefore, another way to incorporate the duration has to be considered. One option is to include the state duration in the model formulas, this requires reformulating the whole model parameters [4]. The model works perfectly in this case but the problem now is with the vast increase in computational cost that makes the use of this new model impractical.

The other option is to use heuristic techniques to include the duration to obtain comparable performance as the correct theoretical duration inclusion with very low computational and storage costs. The state duration probability function $p_j(\tau)$ is estimated during the model training case and defined as:

$p_j(\tau)$: is the probability of being in state j for τ duration.

The duration probability density function is considered to be Gaussian with 3 to 5 mixtures.

During recognition the state duration are calculated from the backtracking procedure in Viterbi Algorithm. Then, the log likelihood value is incremented by the log of the duration probability value as below:

$$\log[\hat{P}(q,O \mid \lambda)] = \log[P(q,O \mid \lambda)] + \eta \sum_{j=1}^{N} \log[p_j(\tau_j)] \tag{84}$$

where η is a scaling factor.

τ_j is the normalised duration of being in state j, as detected by Viterbi Algorithm.

9-6. Data Representation Factor

The training and testing speech data are taken from:

http://Kel.otago.ac.nz/hyspeech/corpus

The initial sets of data are the digits 0-9 spoken by 21 speakers (11 males and 10 females) and each digit is spoken three times by each speaker. Among those words 42 uttered digits are used for training and 15 for testing. The speech data in the Otago Speech Corpus are sampled at 22050 Hz with short silences before and after each utterance.

The next step is to transform the time signal into Mel scale coefficients. The number of coefficients are selected to be 26 (12 mels and 12 delta mels with one power and its delta). Also experiments have been done on 13 coefficients without considering the dynamic behaviour of the signal. The Mel scale coefficients as extracted features are selected because they imitate to some extent the feature selection in human ears. The Mel scale method considers the spectrum to be linearly distributed below 1000Hz and logarithmically above that. This makes the filter banks move on linear centres below 1000Hz (i.e. 100, 200, 300, …, 1000) and on logarithmic centres over that (i.e. 1149, 1320,1516,....). A very good characteristic of the Mel scale coefficients is that they allow the use of Euclidean distance measures in finding the distance between two examples. This greatly reduces the computational cost of procedures that depend on distance measures like those VQ.

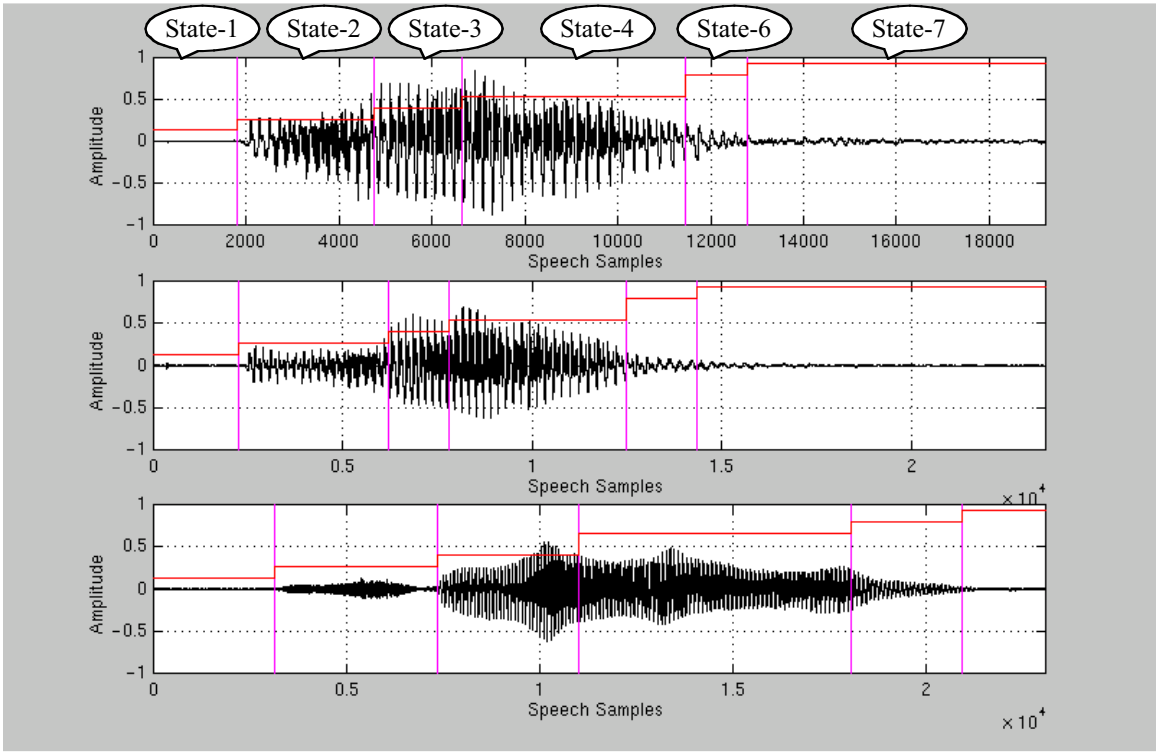
10. Results and Conclusions

In this section we are going to show some experimental results and discuss some useful conclusions.

10-1. Experiment One

The first experiment dealt with the segmentation of the spoken words into states. Fig.(8) shows different versions of the spoken word "zero" by three different speakers. We can see clearly how the time signal varied even for the same word. The states are found by Viterbi Algorithm and assigned clearly to their corresponding segments. Also we can see that the observations are not always passing through all the states that the model designed on. In this case state 5 was jumped by digit zero observations when they were submitted to the digit zero model.

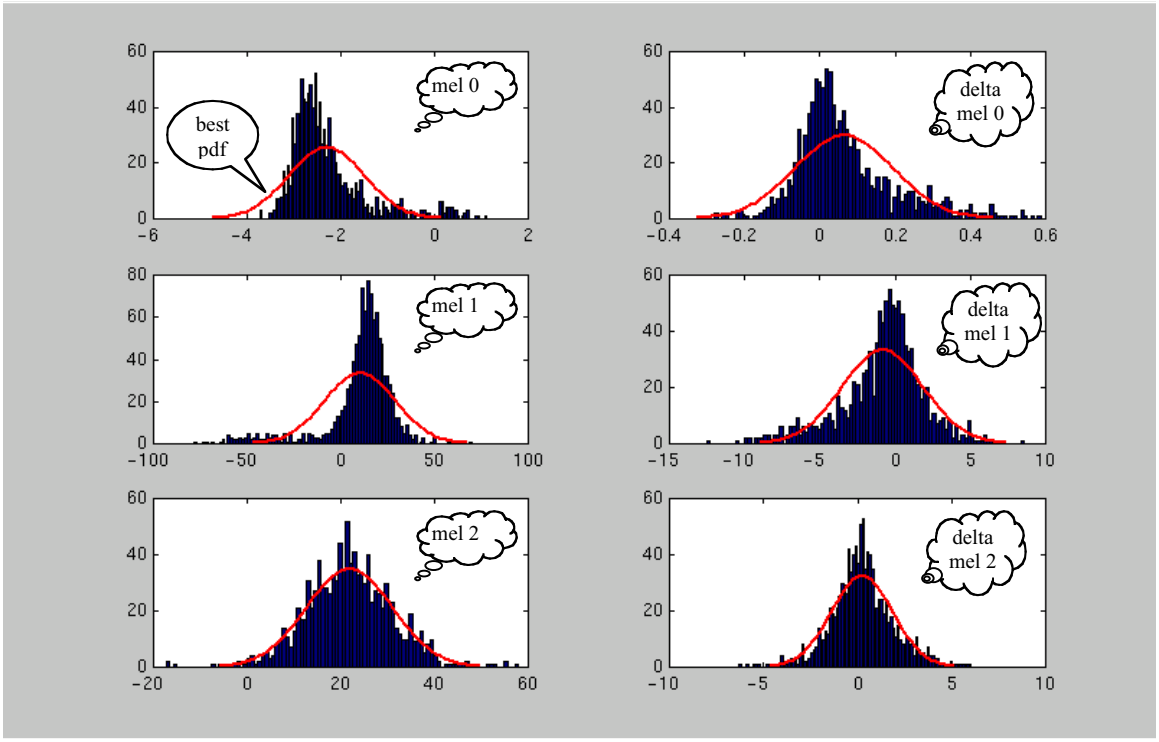
Fig.(8) States' Assignment of Digit ZERO Presented to the ZERO State Model.
The analysis uses 13 Mel scale coefficients without taking the dynamic coefficients into consideration. There are 6 states detected in all the three versions of the spoken digit ZERO, i.e., 1, 2, 3, 4, 6, and 7.



10-2. Experiment Two

The second experiment dealt with the Mel scales coefficients distribution. Fig.(9) Shows some distributions of Mel scale coefficients and their deltas of state one in spoken digit zero. The power of the signal and its delta are represented by mel 0 and delta mel 0. The Mel coefficients capture the stable signal characteristics, while the deltas capture the dynamic characteristics. Also, we can see from this figure the best fit probability distribution function (pdf) for each coefficient. It is clear that some coefficients like mel 0, mel 1 and their deltas are far from being represented by single pdf. This consolidates the need of multimodal (multimixture) representation of the coefficients. In our model we approximate the Mel scale coefficients distribution by 5 to 9 mixtures.

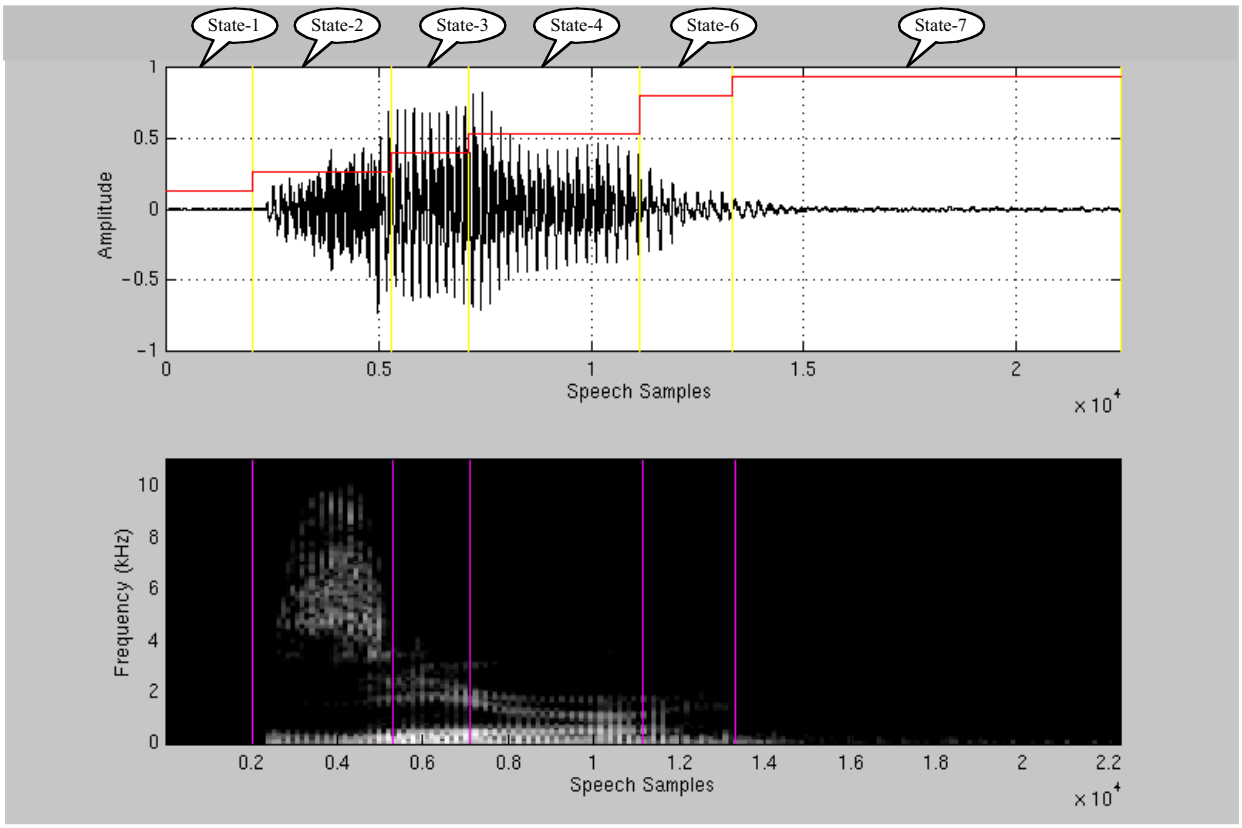
Fig.(9) Mel Scale Coefficients Distribution
The histogram and the best fit normal pdf of mel0,mel1, and mel2 with their deltas.



10-3. Experiment Three

The third experiment was carried out to show the correspondence between the speech signal, states, and the spectra. This relation gave us more understanding and confidence on the behaviour of the observation vectors within each state. Fig.(10) shows this clearly and we can notice the difference in spectral behaviour of different states.

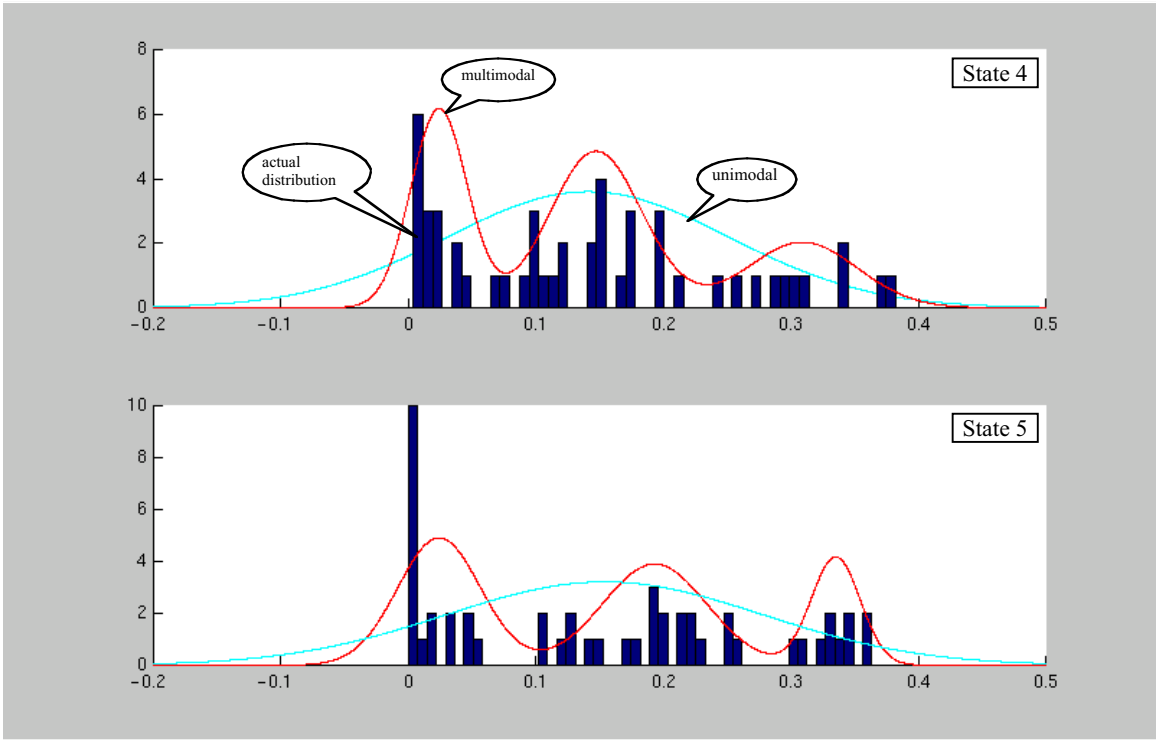
Fig.(10) Correspondence between the time signal samples, states, and spectrum of spoken digit zero.



10-4. Experiment Four

The fourth experiment was about the representation of the state duration distribution using multimodal (multimixture) probability distribution. Fig.(11) shows the normalised duration probability distribution for unimodal and multimodal (with 3 mixtures) representation. The multimodal pdf shows superiority in representing the distribution.

Fig.(11) Multimodal Representation of States' Duration.
The unimodal poorly represents the states' duration, while the multimodal smoothly follows the duration distribution even with only three mixtures used.



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