ON NON-LINEAR FINITE STATE DIGITAL CHANNEL CODING

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ABSTRACT

A discrete-time digital channel coding, in which the encoder and decoder are both time-invariant finite state machines, is investigated. The channel is assumed to be memoryless. Two measures of performance are considered: the probability of error and the minimum free distance. Greater emphasis is placed on the minimum free distance since it is easier to evaluate than the probability of error criterion, and for small channel crossover probability it is a very good indicator of system performance; it is defined as the Hamming distance between two possible output sequences that correspond to distinct state sequences of the same length that are identical at both the start and finish. An algorithm to calculate the distance between such possible output sequences was developed; it also identifies all such pairs of output sequence. Another algorithm, based on the finite state machine properties of the encoder, was found to generate the finite state decoder. Finally, after having found the pairs of encoder-decoder finite state machines, we simulated the communication system.

SARI

Suatu kode saluran digital dengan waktu tercacah yang mempunyai pasangan encoder-decoder berupa mesin berkeadaan terhingga dan invarian waktu akan diteliti. Saluran dianggap tanpa daya ingat. Digunakan dua buah ukuran kinerja yaitu probabilitas kesalahan dan jarak bebas minimum. Penggunaan jarak bebas minimum lebih ditekankan karena lebih mudah dievaluasi dibandingkan probabilitas kesalahan. Untuk probabilitas kesalahan saluran yang kecil, jarak itu merupakan indikator yang sangat baik bagi kinerja sistem; ia didefinisikan sebagai jarak Hamming minimum antara dua buah kemungkinan sekuen keluaran yang berupa sekuen keadaan yang unik dengan panjang dan keadaan yang sama, baik di awal maupun di akhir sekuen. Sebuah algoritma untuk menghitung jarak antara sekuen keluaran yang sedemikian itu telah pula dikembangkan; algoritma tersebut juga mampu mengidentifikasi semua pasangan sekuen keluaran itu. Sebuah algoritma lainnya, didasarkan pada sifat-sifat encoder sebagai mesin dengan keadaan berhingga, yang mampu menyusun decoder dengan keadaan berhingga, telah pula ditemukan. Akhirnya, setelah menemukan berbagai pasangan encoder-decoder yang berupa mesin dengan keadaan berhingga, sistem komunikasi digital disimulasikan.

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1 INTRODUCTION

A particular kind of digital communication system has, as its objective, a channel coding. The main idea behind any channel coding scheme is to add some redundant bits to the information sequence so that it has the capability to combat channel noise; the cost of implementing such a system and the rate of information transmission should also be considered. The channel encoder transforms the information sequence \underline{X} into a discrete encoded sequence \underline{D} which is called a codeword, i.e. the information sequence with the additional redundant bits. This discrete encoded sequence \underline{D} is input to the communication channel which outputs a sequence \underline{C} . The channel decoder then transforms the received sequence \underline{C} into a binary sequence \underline{Y} called the estimate information sequence. Ideally, \underline{Y} will be a delayed replica of the information sequence \underline{X} although the noise may cause some decoding errors. This is the type of digital communication system we are concerned with in this paper.

Most of the research on the channel codes has been on classes of linear codes. The basic purpose of codes is to correct errors caused by noisy communication channels, and for this purpose linear codes have many practical advantages. However, the class of non-linear codes is larger than that of linear codes. Consequently, if we want to obtain the largest possible number of codewords with a given property, we should consider non-linear codes. The intent of this paper is to investigate the class of non-linear finite state codes, i.e. codes generated by non-linear finite state encoders, by analyzing the state transition diagram of the codes, i.e. the topological, rather than algebraic, structure. There are two types of channel codes in common use today: block codes and convolutional codes. Block codes were the earliest type of codes to be investigated, and remain the subject of the overwhelming bulk of the coding literature. On the other hand, the performance of convolutional codes has proved to be equal, or superior, to that of comparable block codes in nearly every type of practical application. This anomaly, wherein convolutional codes are easy to implement but relatively unstudied, is due largely to the difficulty of analyzing convolutional codes. Note that every convolutional code is a finite state code; however, not every finite state code is a convolutional code. Here we examine the structure of relatively good codes to understand what makes them better than others, i.e. in terms of closed paths that diverge from and remerge with the same state and the input-output labelling. The results we obtain are far from solving the general problem; however, they do contribute to a better understanding of such codes.

Convolutional codes were first introduced by Elias [1] in 1955 as an alternative to block codes. The Viterbi algorithm [2] was proposed in 1967 as an alternative to the then existing sequential and threshold decoding techniques. This algorithm is a recursive optimal solution to the problem of estimating the

state sequence of discrete time finite-state Markov process in memoryless noise. Many problems in digital communications can be cast in this form [4].

A complete treatment of convolutional codes, and one of the most enlightening, first appeared in a series of papers by Forney [5,6]. In [5] Forney defined a convolutional encoder as any constant linear sequential circuit. The associated code is the set of all output sequences resulting from any set of all input sequences. He emphasized the algebraic structure of the convolutional codes in his first paper. In the second one [6], the topological structure was emphasized. Here, convolutional codes were characterized by a trellis structure. Maximum likelihood decoding was characterized as finding the shortest path through the code trellis; an efficient solution for which is the Viterbi algorithm.

The key concept in the performance analysis of the Viterbi algorithm is the concept of an error event. A first time error event occurs when the true state sequence \underline{S} is first excluded by the state sequence actually chosen by the Viterbi algorithm \widehat{S} . By using this concept, Viterbi [3] derived a bound on the probability of decoding error using the Viterbi algorithm for convolutional codes and memoryless channels. A simple function, the minimum free distance, was found to be a good indicator of the bit error probability performance. The minimum free distance became the most important distance measure for convolutional codes. The best achievable minimum free distance for a convolutional code with a given rate and encoder memory has not been determined exactly. Most code constructions for convolutional codes have been done by computer search. This has prevented the construction of good long codes, since most computer search techniques are time consuming and limited to relatively short constraint lengths. Larsen [7], Paaske [8], and Johanesson [9] published results of a computer construction technique for non-catastrophic codes with maximal minimum free distance for rates 1/4, 1/3, 1/2, 2/3, and 3/4. Upper and lower bounds on the minimum free distance for the best convolutional code have also been obtained by using a random coding approach by Paaske [8] and Costello [10]. An efficient algorithm for finding the minimum free distance based on the state transition diagram of the code has been developed by Bahlet al. [11] and modified by Larsen [12]. This algorithm searches through the state transition diagram of the encoder for the minimum free distance; the diagram is considered as a weighted directed graph whose nodes are the states of the encoder and whose branches are the allowable state transitions. The use of a bidirectional search made this algorithm considerably more efficient than any previously known algorithm.

Most of the results stated above (except [6,11,12,]) were derived by using the algebraic structure; in addition, all these results are restricted to the class of linear codes. Methods to design the best non-linear time invariant finite state codes, by using the topological structure, have not been reported to our knowledge.

To deal with this difficult problem, in section 2 we adopt the model developed by Gaarder and Slepian [13]. They developed a finite state model for a digital transmission system. Their objective was to find a pair of transmitter-receiver finite state machines, which would optimize a given criterion function for source coding. However, we use this model for channel coding rather than source coding. In our context the objective would be to find a pair of encoder-decoder finite state machines, which would guarantee a similar result to that of maximum likelihood decoding. An advantage of this model lies in the fact that this model is powerful for analyzing both linear and non-linear finite state codes.

To judge the system performance we will use two measures: probability of error and minimum free distance. To deal with the first measure, in section 3 we derive a general bound on the bit error probability based upon the error event probability at a particular node; the bit error probability can be approximated by a relatively simple function of a generalized minimum free distance. The bound on bit error probability for convolutional codes is a simplified form of this general bound. This bound leads us to investigate the second performance measure, minimum free distance, in section 4; in particular, we investigate what factors make a finite state encoder have a large minimum free distance. In section 5 we will discuss the catastrophic property; how the catastrophic property can be identified from the state transition diagram of the encoder will be our concern. For simplicity we restrict ourselves to finite state encoders with one closed communicating class. In section 6 we will use a general bidirectional search algorithm to deal with closed paths identification and the minimum free distance calculation. In section 7 the problem treated is the approximation of the Viterbi algorithm by a finite state machine. A general treatment of this problem did not exist previously. Based on this approximation we will construct our decoder machine. However, the results cannot be extended to general channels because the path metrics are not integers. Finally, two computer simulation results will be shown in section 8; each has rate 1/2. Since the number of decoder states grown rapidly only the decoder states of the first example is shown.

2 THE SYSTEM MODEL

Figure 1 shows the general finite state digital communication system. There is a digital source which has statistically independent, identically distributed outputs and a channel encoder and decoder, both of which are time invariant finite state machines. The channel linking them is assumed to be memoryless and time invariant. Furthermore, the entire system is discrete and operates in discrete time. That is, all the system variables take values in finite or countably infinite sets and changes occur only at fixed instants in time.

Thus, at time n the digital source generates an output sequence, say X_1, X_2, \ldots Here, each X_i is drawn from a set of source symbols $X = (1, 2, \ldots, N_X)$. The channel encoder accepts the sequence X_1, X_2, X_3, \ldots as an input and produces as an output the sequence D_1, D_2, D_3, \ldots For $n = 1, 2, 3, \ldots$ the output D_n at time n takes values from $\not D = (1, 2, \ldots, N_D)$. We represent the channel encoder as a time invariant finite state machine with N_S states, i.e. $\not S = (1, 2, \ldots, N_S)$. Let $S_n \in \not S$ be the channel encoder state at time n. We write

$$S_{n+1} = \sigma(X_n, S_n) \tag{1}$$

$$D_{n} = \delta(X_{n}, S_{n}) \tag{2}$$

where the functions $\sigma(.,.)$ and $\delta(.,.)$ constitute a description of the encoder. Obviously these functions can also be described by a transition table or transition diagram.

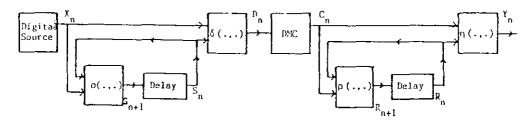


Figure 1. The general finite state digital communication system

The output of the channel encoder, i.e. the encoded message sequence, D_1 , D_2 , . . . is input to the communication channel which outputs the sequence C_1 , C_2 , . . . at a fixed rate. Each C_n is a noisy version of D_n and takes values in a given set $\not C = (1, 2, ..., N_C)$.

Similarly, we represent the decoder as a time invariant finite state machine with $N_{\rm R}$ states, i.e. $R = (1, 2, 3, ..., N_{\rm R})$. The decoder accepts the sequence $C_1, C_2, ...$ as an input and produces as its output the sequence $Y_1, Y_2, ...$ Each Y_n for n = 1, 2, ... takes values from the set X.

Let R_n be the decoder state at time n. We write

$$R_{n+1} = \rho \left(C_n, R_n \right) \tag{3}$$

$$Y_{n} = \eta (C_{n}, R_{n}). \tag{4}$$

By defining the channel encoder-decoder as time invariant finite state machines one can prove that:

1. When the X_i is a sequence of an independent identically distributed random variable, then (S_n, R_n) is a first order Markov chain for a noiseless channel $(C_n = D_n)$ or an additive channel: $C_n = D_n + N_n$ where + denotes a mod. N_C addition and N_n is an independent identically distributed random variable.

2. (D_n, S_n, R_n, Y_n) is a first order Markov chain for a noiseless channel $(C_n = D_n)$ or an additive channel: $C_n = D_n + N_n$ where + denotes a mod. N_C addition and N_n is a sequence of independent identically distributed random variables. These are shown in Appendix A and B.

We now pose the problem we want to investigate:

Given the channel output sequence C_1 , C_2 , C_3 , ..., the channel matrix, the finite state digital channel encoder, find the finite state digital channel decoder such that the output sequence of this machine, i. e. Y_1 , Y_2 , Y_3 , ... is an estimate of X_1 , X_2 , X_3 , ... which is close to that of a maximum-likelihood decoder.

Maximum-likelihood decoding for finite state codes can be realized with the Viterbi algorithm. However, since the number of decoder states have to be finite, a truncated version of the Viterbi algorithm will be used to solve the above problem. In general, the best way to approximate the Viterbi algorithm with a finite number of the decoder states remains an unresolved problem. We restrict ourselves to a specific case. These difficulties will be more evident in the next section.

3 PROBABILITY OF ERROR

Since both the complexity and the operation of the finite state encoder and decoder depend only on the number of states, code rate, and channel parameters, the block length of the code is irrelevant. Furthermore, the bit error performance is primarily a function of relative distances among signals, which may be determined from the code state diagram, whose structure and complexity depend strongly on the constraint length but not on the block length. In addition, block error probability is not a reasonable performance measure, particularly when, as is often the case, an entire message is encoded by the finite state encoder as a single block, whereas in block coding the same message would be encoded into many smaller blocks. Ultimately, the most useful measure is bit error probability $P_{\rm b}$ which is the expected number of bit errors per bit decoded.

While our ultimate goal is to upper bound P_b , we initially consider a more readily determined performance measure, a first error event probability at time i, which we denote $P_f(i)$. Let us recall that first error event at time i occurs when the true state sequence, \underline{S} , is first excluded at time i by the state sequence actually chosen by the Viterbi algorithm \underline{S} ; equivalently one can say that a first error event occurs at time i if the Viterbi path and the correct path are different for the first time at time i+1. For further analysis it is convenient to expand the state transition diagram of the encoder in the time domain, that

is, to represent each time with a separate state diagram; the resulting structure is called a trellis diagram. In the trellis we can now define that the error event occurs at time i if the Viterbi and the correct path are the same at time i but they are different at time i + 1. Figure 2 shows a situation when a first error event occurs at time i in the trellis. By definition we get:

 $P_{\rm f}(i)$ = Pr[the Viterbi path and the correct path, in the trellis, are the same at time i but different at time i + 1]. (5)

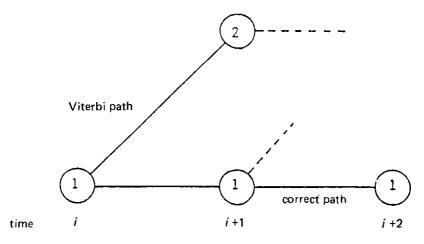


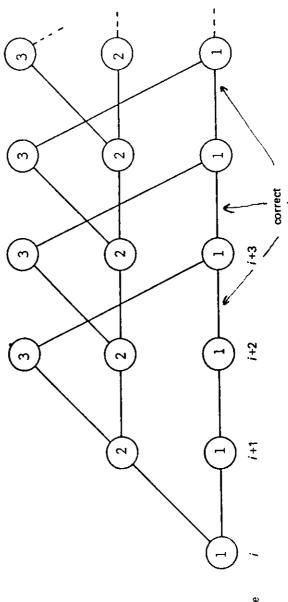
Figure 2 An example of a first error event at time i

Although the Viterbi path and the correct path are different at time i+1, they remerge at some time, at least when the decoder is forced to arrive at a known final state to make the decision. Let these two paths remerge at time i+m. Since all possible Viterbi paths of length i+m or more can cause a first error event at time i, the first error event probability at time i, $P_{\rm f}(i)$, can be overbounded, by the union bound, by the sum of the error probabilities of each of these paths. Obviously Eq. (5) can be rewritten:

 $P_{\rm f}(i) = Pr[\bigcup_{m} \{ \text{ the Viterbi path and the correct path, in the trellis, are the same at time } i \text{ but different at time } i+1 \text{ and remerge at time } i+m \} \}.$ (6) $P_{\rm f}(i) \leq \sum_{m} Pr$ [the Viterbi path and the correct path, in the trellis, are the same at time i but different at time i+1 and remerge at time i+m].

Figure 3 shows some possible Viterbi paths that diverge from the correct path at time i and remerge at time i + m for some value of m, known as the incorrect subset for time i. For this to occur, the correct path metric increments over the unmerged segment, that is over the time span between i and i + m, must be less





time

than those of the incorrect path. Let \underline{S} and $\underline{\widehat{S}}$ denote the correct and the Viterbi paths, respectively; given that $\underline{S} = \underline{S}_k$ is the correct path then Eq. (6) can now be rewritten:

$$P_{\mathbf{f}}\left(i \mid \underline{S} = \underline{S}_{\mathbf{k}}\right) \leq \sum_{S_{\mathbf{k}}} \sum_{\epsilon} \sum_{A_{\mathbf{k}i}} Pr\left\{M_{i+m}\left(\widehat{\underline{S}}_{\mathbf{k}}\right) - M_{i}\left(\widehat{\underline{S}}_{\mathbf{k}}\right) > M_{i+m}\left(\underline{S}_{\mathbf{k}}\right) - M_{i}\left(\underline{S}_{\mathbf{k}}\right)\right\} \tag{7}$$

where $M_j(\underline{S})$ is the metric of the \underline{S} path at time j and A_{ki} is the incorrect subset of $\underline{S} = \underline{S}_k$ for time i, that is, all possible incorrect paths that diverge from the correct path $\underline{S} = \underline{S}_k$ at time i. By noticing that this bound is independent of i, one finds that it holds for all i. Since $M_{i+m}(\underline{S}_k) - M_i(\underline{S}_k)$ is simply the metric increment of \underline{S}_k over the time span between i and i+m then the first error event probability at any time, given that $\underline{S} = \underline{S}_k$ is the correct path, is simply upper-bounded by:

$$P_{\mathbf{f}}(\underline{S} = \underline{S}_{\mathbf{k}}) \leq \hat{\underline{S}}_{\mathbf{k}} \sum_{\epsilon} \sum_{A_{\mathbf{k}}} Pr[\Delta M(\hat{\underline{S}}_{\mathbf{k}}, \underline{S}_{\mathbf{k}})]$$
 (8)

where $\Delta M(\hat{S}_k, \underline{S}_k)$ is the difference between the metric increment of $\underline{\hat{S}}_k$ and \underline{S}_k over the unmerged segment, and A_k is the incorrect subset of $\underline{S} = \underline{S}_k$. Clearly the first error event at any time is upper-bounded by:

$$P_{\mathbf{f}} \leq \sum_{\mathbf{k}} Pr[\underline{S} = \underline{S}_{\mathbf{k}}] \sum_{\widehat{\underline{S}}_{\mathbf{k}}} \sum_{\epsilon A_{\mathbf{k}}} Pr[\Delta M(\widehat{\underline{S}}_{\mathbf{k}}, \underline{S}_{\mathbf{k}})]. \tag{9}$$

The final Viterbi path can diverge and remerge with the correct path any number of times, that is, it can contain any number of error events. Figure 4 shows a situation where multiple error events occur. After the first error event has occurred, the two paths to be compared will both be incorrect paths, one of which will be a previous error event. This is illustrated in Figure 5, where it is assumed that some segments of the correct path \underline{S} have been excluded for the first time at time i by an incorrect path \underline{S}_1 , and assumed that there is another incorrect path \underline{S}_2 that diverges from \underline{S} at time j. However, since at time j they both have the same metric, then whether some segments of \underline{S} will be excluded

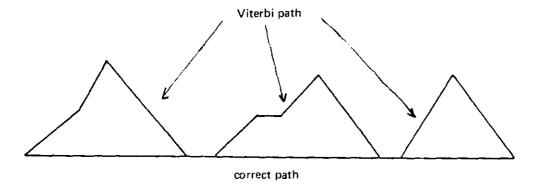


Figure 4 A situation where multiple error events occur

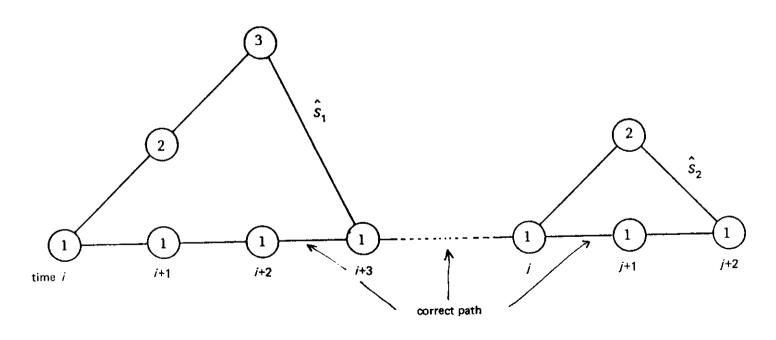


Figure 5 A comparison of two error events

again at time j by \widehat{S}_2 depends only on their metric increment over the time span between j and j+2. If the metric increment over the unmerged segment, that is over the time span between j and j+2, for \widehat{S}_2 exceeds that of some segments of \underline{S} then \underline{S} will be excluded by \widehat{S}_2 at time j, and we say the second error event has occurred at time j. Let $P_2(j)$ denote the probability that some segments of \underline{S} over the time span between j and j+2 is excluded by \widehat{S}_2 at time j. Now, after the second error event has occurred, the Viterbi path contains \widehat{S}_1 , some segments of \underline{S} over the time span between i+3 and j, and \widehat{S}_2 . The probability for this to occur, that is, the error event probability at time j, can be approximated by $P(E_j) = P_f \times P_2(j)$. Obviously, if \underline{S} has not been excluded by \widehat{S}_1 at time i then $P_2(j)$ is simply equal to $P_f(j)$, a first error event probability at time j. We conclude from this exposition that the error event probability at time j, $P(E_j)$, is upper-bounded by:

$$P(E_i) \le P_i(j) \tag{10}$$

However, since it holds for all time, we will use the error event probability at any time P(E) rather than for a particular time $P(E_j)$. In addition, we restrict ourselves to finite state codes with one closed communicating class. Let us now evaluate the upper-bound on the error event probability at any time, P(E).

From Eqs. (9) and (10) one can find:

$$P(E) \leq \sum_{\mathbf{k}} Pr\left[\underline{S} = \underline{S}_{\mathbf{k}}\right] \sum_{\underline{S}_{\mathbf{k}}} \sum_{\epsilon : A_{\mathbf{k}}} Pr\left[\Delta M(\underline{S}_{\mathbf{k}}, \underline{S}_{\mathbf{k}})\right]$$
 (11)

Let $\widehat{\underline{D}}_k$ and \underline{D}_k be the corresponding code sequences of the Viterbi path $\widehat{\underline{S}}_k$ and the correct path \underline{S}_k , respectively. Each term of the summation over A_k in (11) is the pairwise error probability for two code sequences over the unmerged segment. For a binary input channel this is readily upper-bounded by a function of the distance between code sequences over this segment. For, if the total Hamming distance between code sequences $\widehat{\underline{D}}_k$ and \underline{D}_k (over their unmerged segment) is $d(\widehat{\underline{D}}_k,\underline{D}_k)=d$, for the binary symmetric channel, the pairwise error probability is bounded by [3]

$$P_{\rm d} \le \exp\left[d \ln \sum_{\ell} (p_0(c) p_{\rm i}(c))^{0.5}\right]$$
 (12)

where $P_i(c)$ is the conditional (channel transition) probability of output c given that the input was i(i = 0,1). Thus, given that there are a(d) incorrect paths which are at Hamming distance d from the correct path over the unmerged segment, we obtain

$$P(E) \leq \sum_{k} Pr\left[\underline{S} = \underline{S}_{k}\right] \sum_{d=d_{\min}}^{\infty} Pr\left[\text{error caused by any one of } a(d(\underline{\hat{D}}_{k}, \underline{D}_{k}))\right]$$
incorrect path at distance $d(\underline{\hat{D}}_{k}, \underline{D}_{k})$

$$P(E) \leq \sum_{k} Pr[\underline{S} = \underline{S}_{k}] \sum_{d=d_{\min}}^{\infty} a(d(\widehat{\underline{D}}_{k}, \underline{D}_{k})) Z^{d(\widehat{\underline{D}}_{k}, \underline{D}_{k})}$$
(13)

$$a(d(\hat{\underline{D}}_k, \underline{D}_k)) = \sum_{k}, \Delta_{k k'}$$

$$\Delta_{k k} = \begin{cases} 1; \underline{D}_{k} - \widehat{\underline{D}}_{k} \neq \underline{D}_{k'} - \widehat{\underline{D}}_{k'}, k' < k \text{ for all } k \\ 0; \text{ otherwise} \end{cases}$$

where $Z = \sum_{\phi} (p_0(c)p_1(c))^{0.5}$, d_{\min} is the minimum distance of any path from

the correct path and $\Delta_{\mathbf{k}|\mathbf{k}'}$ guarantees that the same configuration on the unmerged segment will not be counted twice by a(d), with this $P(E | \underline{S} = \underline{0}) = P(E)$. Henceforth now we will write $d(\widehat{\underline{D}}_{\mathbf{k}}, \underline{D}_{\mathbf{k}})$ as d.

If the channel transition probability is small, Z will be very small; thus one can

approximate $\sum_{d=d_{\min}}^{\infty} Z^{d}$ by $Z^{d_{\min}}$. Consequently the summation over all k is

approximately the summation over k such that $d = d_{min}$. Hence we get

$$P(E) \sim \sum_{\mathbf{k}} Pr[\underline{S} = \underline{S}_{\mathbf{k}}] \ a(d_{\min}) Z^{\dim_{\mathbb{N}}} : \widehat{\underline{S}}_{\mathbf{k}}, \underline{S}_{\mathbf{k}} \in A_{\min}$$
 (14)

where A_{\min} is the set of all possible $\widehat{\underline{S}}_k$ and \underline{S}_k for all k such that $d = d_{\min}$. Assume that all \underline{S}_k are equally probable and have length ℓ over the unmerged

segment, then $Pr[\underline{S} = \underline{S}_k] = \frac{1}{2} \ell$. Let ℓ_{\min} be the minimum length of all unmer-

ged segments which produces d_{\min} . We now find:

$$P(E) \le (1/(2^{\ell_{\min}})) a(d_{\min}) Z^{\ell_{\min}}$$
 (15)

The Viterbi algorithm requires that a final decision on the most probable code path be made by forcing the coder into a known state. However, one can choose any state as the final state. For simplicity we choose the final state equal to the initial state. The distance between the correct and incorrect paths is then simply equal to the distance between two distinct equal length closed paths which we call the free distance and d_{\min} is the minimum free distance, d_{free} . Now we get the final form by rewriting the Eq. (15):

$$P(E) \le (1/(2^{\ell_{\min}})) a(d_{\text{free}}) Z^{\text{d} \text{free}}$$
(16)

where $a(d_{\text{free}})$ can be evaluated from $L_{1, \mathcal{R}'}$ which is the set of all possible closed paths of length \mathcal{R}' that begin with and end with I if the minimum distance between closed paths in $L_{1,\mathcal{R}'}$ is equal to d_{free} .

For an illustrative example let the encoder be as shown in Figure 6. Let us choose the initial and final state I equal to state 1. By using the computational results shown in Table 4 we get:

 $L_{1,6} = \{\underline{S}_1, \underline{S}_2, \ldots, \underline{S}_{16}\} = \{1333321, 1333221, \ldots, 1222212\}$. From Table 5 we find that there are pairs of correct and incorrect paths with the distance over the unmerged segment equal to 3. Clearly the unmerged segment with minimum length will be contained in the following pairs of closed paths:

$$\begin{array}{l} (\underline{S}_{1},\underline{S}_{2}), \ (\underline{S}_{2},\underline{S}_{4}), (\underline{S}_{3},\underline{S}_{4}), (\underline{S}_{3},\underline{S}_{7}), \ (\underline{S}_{4},\underline{S}_{8}), (\underline{S}_{5},\underline{S}_{6}), \ (\underline{S}_{5},\underline{S}_{13}), \ (\underline{S}_{6},\underline{S}_{8}), (\underline{S}_{6},\underline{S}_{14}), \\ (\underline{S}_{7},\underline{S}_{8}), \ \ (\underline{S}_{7},\underline{S}_{15}), \ (\underline{S}_{8},\underline{S}_{16}), \ (\underline{S}_{9},\underline{S}_{10}), \ (\underline{S}_{10},\underline{S}_{12}), \ (\underline{S}_{11},\underline{S}_{12}), \ (\underline{S}_{11},\underline{S}_{15}), \\ (\underline{S}_{12},\underline{S}_{16}), \ (\underline{S}_{13},\underline{S}_{14}), \ (\underline{S}_{14},\underline{S}_{16}), \ (\underline{S}_{15},\underline{S}_{16}). \end{array}$$

But when we look at the ummerged segment it is apparent that there are only three distinct unmerged segments with minimum length $\ell_{\min} = 2$; (132, 122), (212, 222), and (332, 322); obviously $a(d_{\text{free}}) = 3$. If the channel crossover probability is p, one finds

$$P(E) \le (1/4) (3) ((p^{0.5} (1-p)^{0.5})^3).$$

For $p = 10^{-2}$ one can easily get $P(1) \le 3.0 \times 10^{-4}$.

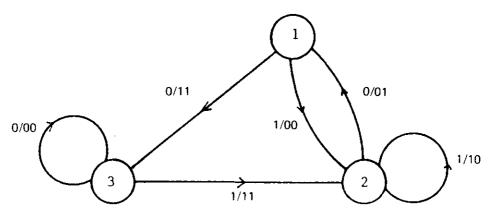


Figure 6 An encoder with three states and $d_{\text{tree}} = 3$

Turning now to the bit error probability, we note that the expected number of bit errors, caused by any incorrect path which diverges from the correct path, can be bounded by weighting each term of the union bound by the number of bit errors which occurs on that incorrect path. This number is the number of $\widehat{X}_n \neq X_n$ over the unmerged segments where \widehat{X}_n and X_n are the information digits corresponding to the incorrect and correct path, respectively. Thus the bound on the expected number of bit errors caused by an incorrect path is:

$$P_{b} = E[n_{b}] \le \sum_{k} Pr[\underline{S} = \underline{S}_{k}] \sum_{i=d}^{\infty} i \ a(d, i) \ Z^{d}$$

$$\tag{17}$$

where a(d, i) is the number of paths diverging from the correct path at distance d and with the number of $\hat{X}_n \neq X_n$ over the unmerged segments is equal to i. For the same condition we can substitute (16) into (17) and yield:

$$P_{\rm b} \le (1/(2^{\ell_{\rm min}})) \sum_{i} i \, a(d_{\rm free}, i) \, Z^{\rm d_{\rm free}} \tag{18}$$

For an illustrative example let us use the previous example. It is easy to see that i = 1 for all the unmerged segments and hence $P_b = P(E) \le 3.0 \times 10^{-4}$. It will be shown in the computational result that for the same encoder and p our simulation get the relative frequency of the error bit which is equal to 1.0×10^{-4}

4 MINIMUM FREE DISTANCE

The most important measure for finite state codes is the minimum free distance d_{free} . This is largely due to the relation between this distance and the bit error probability according to Eq. (18). In general, the minimum free distance is defined as the minimum Hamming distance between distinct output paths corresponding to state paths of the same length that begin and end with the same state:

$$d_{\text{free}} = \min \left\{ d(\underline{D}', \underline{D}''); \underline{X}' \neq \underline{X}'' \text{ and } \underline{X}', \underline{X}'' \in L \right\}$$

$$L = \bigcup_{\mathbf{I}, \mathbf{\hat{\chi}}} L_{\mathbf{I}, \mathbf{\hat{\chi}}}, d(\underline{D}', \underline{D}'') = \sum_{i} |D_{i}' - D_{i}''|$$
(19)

where \underline{D}' and \underline{D}'' are the codewords that correspond to the information sequences \underline{X}' and \underline{X}'' , respectively, i runs over ℓ branches; and $L_{i,\ell}$ is the set of information sequences produced by traversing closed state paths of length ℓ that diverge from and remerge with the state I.

For a linear code the distance between any two codewords is equal to the weight of another codeword. Hence, for a convolutional code, the minimum free distance is the minimum weight of a codeword produced by a non-zero information sequence. Then for the class of linear codes, Eq. (19) is simplified to:

$$d_{\text{free}} = \min \left(d(\underline{0}, \underline{D}''); \underline{X}'' \neq 0 \text{ and } \underline{X}'' \in L \right)$$

$$L = \bigcup_{\varrho} L_{0, \varrho}, \text{ and } d(0, D'') = \sum_{i} |0 - D_{i}''|$$
(20)

where \underline{D}'' is the codeword corresponding to the information sequence \underline{X}'' ; and $L_{0,\ell}$ is the set of information sequences produced by traversing closed state path of length ℓ that diverge from and remerge with the all-zero state.

Since i in Eq. (19) and (20) runs over ℓ branches, then in terms of graph structure, it seems that an encoder with a greater minimum ℓ should have a bigger minimum free distance. Figure 7 gives an illustrative example of this structural observation. We expect that M_1 will have a bigger minimum free distance than M_2 since the minimum ℓ in M_1 is 3 while in M_2 it is only 2. Those closed state

paths with minimum ℓ in M_1 are $L_{1,3}$ = (1111, 1241), $L_{2,3}$ = (2412, 2342), $L_{3,3}$ = (3333, 3423), and $L_{4,3}$ = (4124, 4234) while those closed state paths with minimum ℓ in M_2 are $L_{1,2}$ = (111, 121) and $L_{2,2}$ = (222, 212).

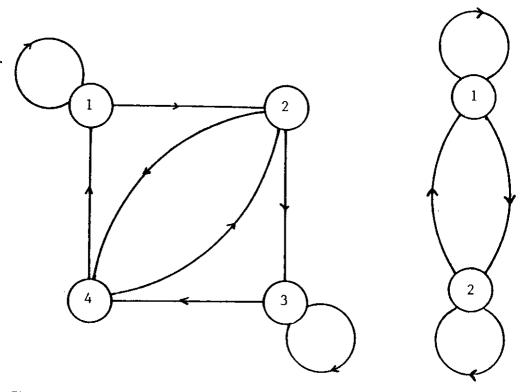
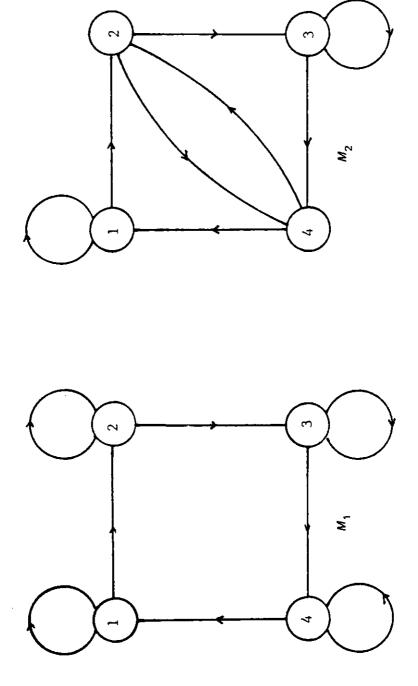


Figure 7 Two encoders with different minimum lengths of closed paths

However, maximum minimum ℓ is not the only condition to ensure a good encoder in terms of its graph. Even an encoder, say M_1 , with its minimum ℓ is greater than that of M_2 , say, may result in a smaller minimum free distance if there are more branches incommon to the closed state paths in $L_{1,\ell}$ for M_1 than for M_2 . Figure 8 will clarify this immediately.

The encoder M_1 has the minimum ℓ equal to 4 while in M_2 it is only 3. Those sets of closed state paths with ℓ equal to 4 in M_1 are $L_{1,4}=(11111,12341)$, $L_{2,4}=(22222,23412), L_{3,4}=(33333,34123),$ and $L_{4,4}=(44444,41234);$ and for M_2 those sets of closed state paths with minimum $\ell=3$ are $L_{1,3}=(1111,1241), L_{2,3}=(2412,2342), L_{3,3}=(3333,3423),$ and $L_{4,3}=(4124,4234).$ But there are many branches incommon for the longer pairs of closed state paths in M_1 . Let us examine a set of closed state paths, with length 5, that diverges from and remerges with state 1, that is $L_{1,5}=(112341,122341).$ It is clear that there are three branches, out of 5, coincide in $L_{1,5}$ of M_1 and this certainly





reduces the minimum free distance. This circumstance does not occur in M_2 so that one can expect that M_2 will have a bigger minimum free distance than M_1 . However, the largest value of ℓ needed to ensure that one encoder is better than another is still unresolved.

A good labelling should produce maximum minimum free distance while not allowing catastrophic properties; in addition, it also could lead back to the initial state for a fixed input sequence. For a linear finite state encoder (with the all zero state as the initial state) the all zero input leads back to the initial state. So obviously given a good graph with a large minimum ℓ and a minimum number of branches coinciding in $L_{1,\ell}$ another problem remains: the labelling of the branches.

The task of finding methods to design optimal and practical encoders in terms of the state transition diagram appears to be inherently very difficult. Here we provide a computer program to enumerate and identify all closed paths, and having found these closed paths the distances will be calculated for a given encoder. All details of this graph enumerator will be discussed in the next section.

5 CATASTROPHIC PROPERTY

A finite state encoder is catastrophic if a finite number of channel errors cause an infinite number of decoding errors. This undesirable circumstance may happen if there are at least two infinite information sequences \underline{X} and \underline{X}' that can be encoded into \underline{D} and \underline{D}' , respectively, such that

$$d(\underline{D}, \underline{D}') \le \infty \text{ an } d(X, X') = \infty$$
 (21)

where $d(\underline{Y}, \underline{Z})$ denotes the Hamming distance between \underline{Y} and \underline{Z} . If the channel noise changes \underline{D}' into \underline{D} , by simply changing a finite number of digits of \underline{D}' , then \underline{X}' will be estimated as \underline{X} that is infinitely distant from \underline{X}' .

For a linear code it is well known that the state transition diagram of a catastrophic code contains a loop of zero weight other than the self-loop around the all zero state. An example of a linear catastrophic encoder in terms of its state transition diagram is shown in Figure 9. The state transition diagram in Figure 9 contains two cycles, i. e. 00 00 and 11 11, that are generated by different input sequences but have the same output sequences. This certainly will produce catastrophic codes. The following example will clarify this catastrophic property. Let $\underline{X} = 000000 \dots$ and so $\underline{D} = 000000000 \dots$. Obviously $\underline{X}' = 01111111 \dots$ with $\underline{D}' = 001101000000000 \dots$ in which $d(D, D') < \infty$ for $d(\underline{X}, \underline{X}') = \infty$. Consequently, if the channel noise changes the three non zero digits of \underline{D}' then \underline{X}' will be estimated as \underline{X} that is infinitely distant from \underline{X}' . Hence this is a catastrophic code.

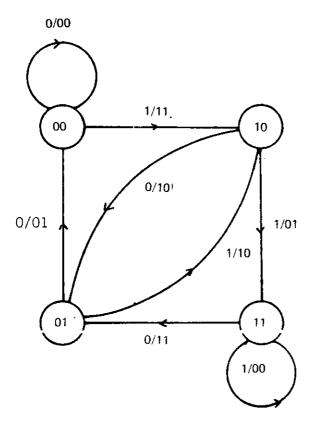


Figure 9 An example of a linear catastrophic finite state encoder with four states

Figure 10 shows an example of a non-linear finite state encoder, in terms of the state transition diagram, that will produce a catastrophic code. The state transition diagram in Figure 10 contains two cycles, i. e. 11 and 232, that are generated by different input sequences but have the same output sequences. This is an obvious catastrophic encoder. The following simple example will clarify this catastrophic property. Let $\underline{X} = 11010101010\dots$ and so $\underline{D} = 111000000\dots$ Clearly there is $\underline{X}' = 0000\dots$ with $\underline{D}' = 000000\dots$ has $d(\underline{D},\underline{D}') < \infty$ for $d(\underline{X},\underline{X}') = \infty$. Hence this is a catastrophic code.

From the above observations one can conclude that in the state transition diagram of a finite state encoder with one closed communicating class Eq. (21) will be satisfied if there are at least two cycles generated by two different input sequences that generate exactly the same output sequence when the cycle repeats forever.

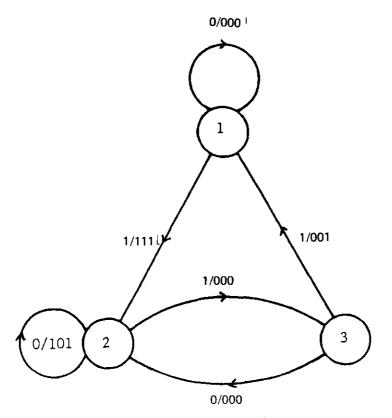


Figure 10 An example of a non-linear catastrophic finite state encoder with three states

6 CLOSED PATH IDENTIFICATION AND dfree CALCULATION

A comparison of the performance of encoders with the same minimum free distance $d_{\rm free}$ can be obtained by calculating the number of pairs of closed paths that produce $d_{\rm free}$; the better encoder has fewer such pairs. To identify these pairs we use a generalized bidirectional search. This search algorithm will be based on the finite state encoder as a directed graph that can be visualized as a directed tree.

Consider a finite state encoder M with N_S states as a weighted directed graph; the nodes are the N_S states of M and the branches are the allowable state transitions. The branches of this graph are labelled with the input/output pairs according to Eq. (3). A closed path of length ℓ that diverges from and remerges with node I can be visualized as a rooted directed tree with ℓ levels and the terminal node equal to the root I, i. e. the node at level 0, If K = (0,1) then this tree is a binary rooted directed tree and there will be 2^{ℓ} paths from the root to the 2^{ℓ} terminal nodes. Henceforth we consider only binary K. To identify all closed paths of length ℓ that diverge from and remerge with state I the search

should be conducted on those 2^{ℓ} paths that have terminal nodes equal to I at level ℓ . Computationally all $\sum_{i=0}^{\ell} 2^i = 2^{\ell+1} - 1$ nodes should be stored in the memory.

To reduce the number of nodes stored the closed paths are searched from two directions; forward (F) and backward (B). This reduces the number of nodes

stored to $\sum_{i=0}^{Q/2} 2^{i+1} = 2(2^{(Q/2)+1}-1)$ if the length of the F path is equal to the

length of the B path. If ℓ is odd, we choose $\ell_1 = \lfloor \ell/2 \rfloor + 1$ as the length of the F path, where $\lfloor y \rfloor$ is the greatest integer smaller than or equal to y, and hence $\ell_2 = \ell - \ell_1$ is the length of the B path. The closed paths can be determined by merging all F paths and B paths.

An $F_{\rm I}$ path is defined as a path starting with node I with all arrows pointing in the forward direction. Since the input is binary, there are 2^i paths of length i branches with the same initial node I. The $F_{\rm I}$ path obviously can be visualized as binary rooted tree with I as the root. Figure 11 shows this observation.

Let $F_{I,i}$ be the set of all forward paths of length i branches whose initial node is I:

$$F_{1,i} = \{ \underline{f}_i; i = 1, 2, \dots, 2^i \}$$
 (22)

where each f_j path is a sequence of states $f_j = f_{j0} = f_{j0}$, f_{j1} , ..., f_{ji} and $f_{jk} = \sigma(x, f_{j(k-1)})$ for some x. Obviously $f_{j0} = I$ for all j.

Let $H_{I,i}$ be the set of terminal nodes of all F paths of length i whose initial node is I. Since each branch represents only allowable state transitions, we can write:

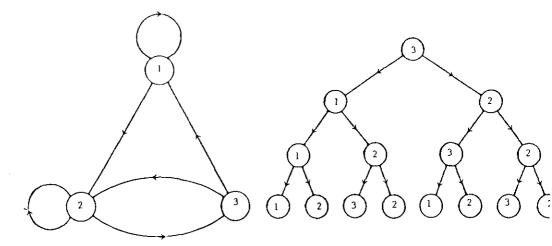


Figure 11. A binary encoder M with its F_3 path that is a binary rooted directed tree

$$H_{1,i} = \{ \sigma(h_{j(i-1)}, k'); k' \in X, h_{j(i-1)} \in \$, \ j = 1, 2, \dots, 2^{(i-1)} \}$$
 (23)

Obviously for i = 1 we can write $H_{I,1} = (\sigma(I,0), \sigma(I,1))$.

Without loss of generality, and in a computationally simple manner, we generate $H_{\mathrm{L},\mathrm{i}}$ as follows:

$$H_{1,i} = \{ \sigma(h_{j(i-1)}, 0), \sigma(h_{j(i-1)}, 1); h_{j(i-1)} \in \}, j = 1, 2, ..., 2^{(i-1)} \}$$
 (24)

This equation implies that the left branches at level i are produced by the input 0 to the nodes at i-1 while the right branches are produced by the input 1. Thus we can generate all the elements of H_{1}, ℓ_{1} recursively from the Eq. (24) for $i = 1, 2, \ldots, \ell_{1}$.

The next problem is to recover $F_{\mathbf{I},\mathfrak{Q}_1}$ from the elements of $H_{\mathbf{I},\mathfrak{Q}_1}$. Recall that $f_j \in F_{\mathbf{I},i}$ has been defined such that $f_{ji} \in H_{\mathbf{I},i}$ for $i = 1,2,\ldots,\mathfrak{Q}_1$. Hence the problem of recovering $F_{\mathbf{I},\mathfrak{Q}_1}$ is the same as the problem of choosing the element of $H_{\mathbf{I},i}$ to which f_{ji} is equal to for all j and i. Since $H_{\mathbf{I},i}$ has been generated according to Eq. (24) then this problem can be solved very easily by defining the counting function that in fact serves as a pointer:

If

$$K(j,m) = [(j+2^{(i-m)}-1)/2^{(i-m)}]$$
(25)

then

$$f_{\rm jm} = h_{\rm Km} \tag{26}$$

where

 $h_{\mathrm{Km}} \in H_{\mathrm{I},\mathrm{m}} \text{ and } H_{\mathrm{I},0} = \{1\}$

[y] = the greatest integer smaller than or equal to y

$$j = 1, 2, \dots, 2^{i}$$

$$m = 0,1,\ldots,i$$

$$i = 1,2,\ldots,\ell_1$$

Hence E_q . (26) defines $F_{I,\ell_1} = \{f_j; j=1,2,\ldots,2^{\ell_1}\}$. Note that F_{I,ℓ_1} can also be generated by considering input sequences to the state I as a binary number. Example: Consider the finite state encoder M of Figure 6. Let $\ell_1 = 3$ and I = 3. Suppose we want to recover $f_1 = f_{10}f_{11}, f_{12}, f_{13}$. From (22) and (23) we get $H_{3,1} = \{h_{11}, h_{21}\} = \{1, 2\}; H_{3,2} = \{h_{12}, h_{22}, h_{32}, h_{42}\} = \{1, 2, 2, 3\}$ and

 $H_{3,3} = \{h_{13}, h_{23}, h_{33}, h_{43}, h_{53}, h_{63}, h_{73}, h_{83}\} = \{1, 2, 2, 3, 2, 3, 1, 2\}$. From (25) we get K(1,0) = K(1,1) = K(1,2) = K(1,3) = 1 and from (26) yields $f_{10} = h_{10} = 3$; $f_{11} = h_{11} = 1$; $f_{12} = h_{12} = 1$ and $f_{13} = h_{13} = 1$. Hence $f_{13} = 3, 1, 1, 1$. Analogously we can find all $f_{13} = 1, f_{13} = 1$.

Define a B_1 (backward) path as one starting with I and with all arrows pointing

to the backward direction. Unlike an $F_{\rm I}$ path, the $B_{\rm I}$ path cannot be visualized as a binary tree because there may be states that can be reached only through a single branch or through more than two branches. Figure 12 will clarify this observation. State 3 has only one predocessor-state 2; state 2 has all three states as predecessors. Clearly B_3 cannot be visualized as a binary rooted directed tree. The treatment of $B_{\rm I}$ paths is more complicated than that of $F_{\rm I}$ paths. In a convolutional code, the degree of branches in and out of each state is exactly two. Consequently the $F_{\rm I}$ and $B_{\rm I}$ paths in the convolutional codes can be visualized as a binary directed tree; obviously the closed path analysis in a linear class is easier than in a non-linear class of finite state codes.

Let $G_{I,i}$ be the set of terminal nodes of all B_I paths of length i whose initial node is I:

$$G_{I,i} = \{g_{ji}; g_{ji} \in \$, j = 1, 2, \dots, N_{Q}(i)\}$$
 (27)

The elements of $G_{1,i}$ can be searched recursively by noticing that all nodes at level n are the next state of all nodes at level n + 1. Since all arrows are pointing to the backward direction:

$$G_{I,i} = \{k; \sigma(k,k') = g_{j(i-1)}, k' \in X, k \in \S, j = 1,2,..., N_Q(i-1)\}$$
 (28)

Obviously $G_{1,0} = \{I\}$ or $g_{i0} = I$. $N_Q(i)$ is defined by the counting function:

$$N_{Q}(i) = \sum_{j=1}^{N_{Q}} \sum_{k=1}^{(i-1)} \sum_{k'=0}^{N_{S}} \Delta(\sigma(k,k') - g_{j(i-1)})$$
(29)

where

$$\Delta(y) = \begin{cases} 1; y = 0 \\ 0; \text{ otherwise} \end{cases}$$

and
$$N_{\mathcal{O}}(j) = 1$$
 for $j \leq 0$.

 $N_{\mathbf{Q}}(\mathbf{i})$ counts the number of terminal nodes of $B_{\mathbf{I}}$ paths of length \mathbf{i} . This number is obviously equal to the number of $B_{\mathbf{I}}$ paths of length \mathbf{i} itself.

Let $B_{I,i}$ be the set of all backward paths of length i branches whose initial nodes is I:

$$B_{I,j} = \{ \underline{b}_j : j = 1, 2, \dots, N_Q(i) \}$$
 (30)

Associated with each path of b_j is a sequence of states $\underline{b}_j = b_{j0}, bj_1, \ldots, b_{ji}$ arranged in the reverse manner such that $b_{jm} \in G_{\mathbf{i},(\mathbf{i}-m)}$ for $m = 0,1,\ldots,i$.

Since the elements of $G_{I,i}$ for all i can be generated recursively according to Eq. (28), then the problem of recovering all backward paths \underline{b}_{j} is the same as the problem of choosing the elements of $G_{I,(i-m)}$ to which b_{jm} is equal to for all m and all j. Though the problem is similar to that of recovering the \underline{f}_{i}

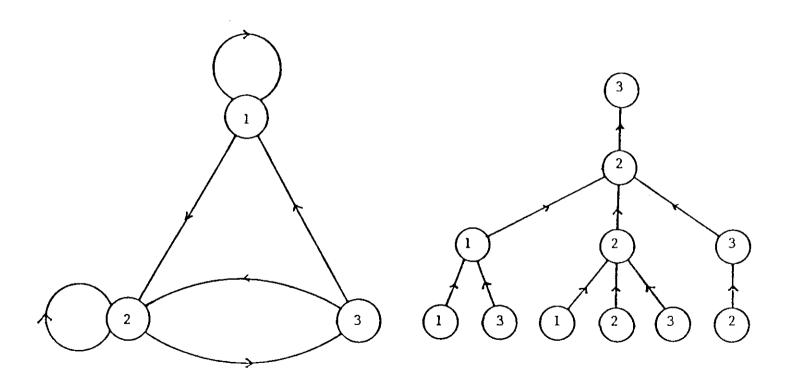


Figure 12 A binary encoder M with its B_3 path that is not a binary rooted directed tree

path, the solution is quite different since the B_1 path cannot be visualized as a binary rooted directed tree. Here we need more counting functions to be used as pointers in $G_{1,(i-m)}$ to which b_{im} is equal to where $i = 1, 2, ..., \ell_2$.

Define the counting function:

$$N_{q}(i,j) = \sum_{k=1}^{N_{S}} \sum_{k'=0}^{\Sigma} \Delta (\sigma(k,k') - g_{j(i-1)})$$
(31)

$$j = 1,2,...,N_{O}(i-1); i = 1,2,...,\ell_{2}.$$

This counts the number of predecessors of $g_{j(i-1)}$ for each j. Note that the predecessors of $g_{j(i-1)}$ are $g_{ni} \in G_{l,i}$ for all j and $n = 1, 2, \ldots, N_Q(\ell_2)$. In other words there are $N_q(i,j)$ elements of $G_{l,i}$ pointing toward $g_{j(i-1)}$.

Define another counting function:

$$N_{\rm A}(k,j(k)) = a_{\rm Nq\ (k+1,j(k))}$$
 (32)

$$N_{\mathbf{A}}(\ell_2, j(\ell_2)) = 1 \tag{33}$$

where

$$a_{i(k,j(k))} = a_{i(k,j(k))-1} + N_{A}(k+1, c_{i(k,j(k)),j(k)})$$

$$c_{i(k,j(k)),j(k)} = \begin{cases} 1; j(k) = 1 \\ \max c_{s(j(k)-1)} + i; \text{ otherwise} \end{cases}$$

$$k = \ell_2 - 1, \ell_2 - 2, ..., 1$$

 $j(k) = 1, 2, ..., N_Q(k)$
 $i(k, j(k)) = 1, 2, ..., N_Q(k+1, j(k))$

This counts the number of occurrences of the elements of $G_{1,k}$ in b_{jk} of $\underline{b}_j \in B_{1,i}$ for all k. So the elements of B_{1,ℓ_2} can be described completely by:

$$b_{\mathbf{k}\mathbf{j}}(\hat{\mathbf{g}}_{2} - \mathbf{k})) = \mathbf{g}_{\mathbf{j}\mathbf{k}} \tag{34}$$

where $k = 1, 2, ..., k_2; j(k) = 1, 2, ..., N_Q(k)$ and $j(k, j(k)) = 1, 2, ..., N_Q(k)$

 $i(k, j(k)) = 1, 2, ..., N_A(k, j(k)).$

Example: Consider the finite state encoder M of Figure 6. Let $\ell_2 = 3$ and I = 3. Suppose we want to recover $\underline{b}_1 = b_{10}, b_{11}, b_{12}, b_{13}$. From (28) and for i = 1 we get $N_Q(1) = 1$ since $g_{10} = I = 3$ for all i and $N_Q(0) = 1$. Then from (26) and (27) we get $G_{3,1} = \{g_{11}\} = \{2.\}$ If we do the same process for i equal to 2 and 3, then $N_Q(2) = 3$; $G_{3,2} = \{g_{12}, g_{22}, g_{32}\} = \{1,2,3\}$ and $N_Q(3) = 6$; $G_{3,3} = \{g_{13}, g_{23}, g_{33}, g_{43}, g_{53}, g_{63}\} = \{1,3,1,2,3,2\}$. From (30) we get $N_Q(1,1) = 1$, $N_Q(2,1) = 3$, $N_Q(3,1) = 2$, $N_Q(3,2) = 3$ and $N_Q(3,3) = 1$. From (3.12) we find

 $N_{\rm A}(3,1)=N_{\rm A}(3,2)=\ldots=N_{\rm A}(3,6)=1, N_{\rm A}(2,1)=2, N_{\rm A}(2,2)=3, N_{\rm A}(2,3)=1$ and $N_{\rm A}(1,1)=6$. Now from (3.14) we find that $b_{12}=b_{22}=b_{32}=\ldots=b_{62}=g_{11}=2,\ b_{11}=b_{21}=g_{12}=1,\ b_{31}=b_{41}=b_{51}=g_{22}=2,\ b_{61}=g_{32}=3,\ b_{10}=g_{13}=1,\ b_{20}=g_{23}=3,\ b_{30}=g_{33}=1,\ b_{40}=g_{43}=2,\ b_{50}=g_{53}=3,\ {\rm and}\ b_{60}=g_{63}=2.$ Since $b_{\rm j}\varrho_2=b_{\rm j3}={\rm I}=3$ for all j, hence $b_1=b_{10}$ b_{11} b_{12} $b_{13}=1123.$

After having found F_{1,ℓ_1} and B_{1,ℓ_2} , one can find all closed paths of length $\ell = \ell_1 + \ell_2$ that diverge from and remerge with state I by merging the elements of F_{1,ℓ_1} and B_{1,ℓ_2} , which is shown in Figure 13, with the following merging rule.

Let $L_{1,\ell}$ be set of all closed paths of length ℓ that diverge from and remerge with state I:

$$L_{I,\ell} = \{ S_i; j = 1, 2, \dots, N_I(\ell) \}$$
 (35)

where $N_1(\ell)$ is defined by

$$N_{\mathbf{I}}(\ell) = \sum_{\mathbf{m} = \mathbf{k}} \sum_{\mathbf{k}} (f_{\mathbf{k} \hat{\ell}_{\mathbf{I}}} - b_{\mathbf{m} \, \mathbf{0}}) \tag{36}$$

Associated with each closed path \underline{S}_j is a sequence of states $\underline{S}_j = S_{j0}, S_{j1}, \ldots, S_j$ where $S_{ji} \in \mathbb{S}$ for all i and all j. Obviously $S_{j0} = S_j = I$. From the derivation of (25) and (33) it follows that the elements of $L_{I,Q}$ are identified completely by:

$$S_{ji} = \begin{cases} f_{ki}, i = 0, 1, \dots, \ell_1 \\ b_{m(i-\ell_1)}, i = \ell_1 + 1, \ell_1 + 2, \dots, \ell \end{cases}$$
(37)

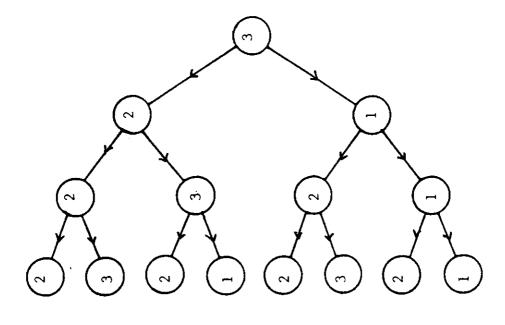
such that

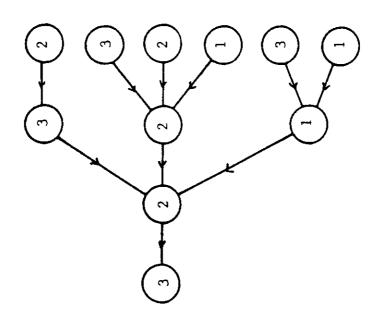
$$f_{\mathbf{k}\,\ell_1} = b_{\mathbf{m}\,0} \tag{38}$$

where $k = 1, 2, \ldots, 2^{\ell_1}$ and $m = 1, 2, \ldots, N_Q(\ell_2)$. Hence $N_I(\ell)$ and $L_{I,\ell}$ enumerate and identify all closed paths of length ℓ that diverge from and remerge with the state I respectively. The number of closed paths of length ℓ that diverge from and remerge with state I may also be calculated by using the adjacency matrix of the encoder.

Let the adjacency matrix $M = [m_{ij}]$ of an encoder M be an $n \times n$ matrix with m_{ij} defined as:

$$m_{ij} = \begin{cases} 1 ; S_j = \sigma(S_i, X_i) \\ 0; \text{ otherwise} \end{cases}$$





Let $m_{ij}(\ell)$ be the (i,j) entry of the ℓ^{th} power of the matrix M, then from [21] we get $N_{I}(\ell) = m_{II}^{(\ell)}$ Now we are ready to calculate the distance between two closed paths.

Consider two closed paths of length ℓ , in terms of state sequences, that diverge from and remerge with state I:

$$\underline{S}_1 = S_{10}, S_{11}, \dots, S_1 \varrho \tag{39}$$

$$\underline{S}_2 = S_{20}, S_{21}, \dots, S_{2\hat{V}} \tag{40}$$

where $S_{j0} = S_{j\ell} = I$, and $S_{ji} \in \underline{S}$ for all i and all j. These two closed paths can be described in terms of output label sequences \underline{D}_1 and \underline{D}_2 , respectively, as follows:

$$\underline{D}_{1} = D_{10}, D_{11}, \dots, D_{1(\hat{X}-1)}$$
(41)

$$\underline{D}_2 = D_{20}, D_{21}, \dots, D_{2(k-1)} \tag{42}$$

where $D_{ji} = \delta(S_{ji}, X_n)$ such that $\sigma(S_{ji}, X_n) = S_{j(i+1)}$ where $X_n \in X = \{0,1\}$.

Hence the distance between \underline{S}_1 and \underline{S}_2 is simply:

$$d_{12} = \sum |D_{1i} - D_{2i}|. (43)$$

7 THE DECODER

Consider a path of length α in a state transition diagram of an encoder of a complete and simple minimal machine (N_S, N_X, N_D) . There are N_X paths of length α entering the N_S states for $\alpha \ge N_S - 1$. There is at least one path of length α entering each state.

If we apply a truncated version of the Viterbialgorithm, then the number of paths of length α entering N_S states can be reduced to exactly N_S paths where each of them minimizes the metric $M(\underline{D})$. Obviously the last symbol in each of these paths is unique.

Let $\underline{S}_{in}^{\alpha}$ be the i^{th} path of length α at time n with metric M_{in} and terminated by the state i. We can write:

$$\underline{S}_{in}^{\alpha} = S_{i(n+\alpha)}, S_{i(n+\alpha-1)}, \dots, S_{in}
S_{ij} \in \$ = (1,2,3,\dots,N_S); S_{i(n+\alpha)} = i
i = 1,2,3,\dots,N_S$$
(44)

 $j = n, n+1, \ldots, n+\alpha$.

This path corresponds to an encoder output sequence of:

$$\underline{D}_{in} = D_{i(n+\alpha-1)}, D_{i(n+\alpha-2)}, \dots, D_{in}$$
(45)

where $D_{ij} = \delta(S_{ij}, X_j)$ such that $\sigma(S_{ij}, X_j) = S_{i(j+1)}$. We can search $S_{j(n+1)}^{\alpha}$ recursively as follows:

$$\underline{S}_{j(n+1)}^{\alpha} = r(\underline{S}_{in}^{\alpha}, C_{n+\alpha})$$
(46)

where r(.,.) is defined such that:

$$M_{j(n+1)} = \min\{M_{in} - \ln Pr(C_{n+\alpha} | D_{i(n+\alpha)}); D_{i(n+\alpha)} = \delta(S_{i(n+\alpha)}, X_k)\}.$$
 (47)

Since $\sigma(S_{i(n+\alpha)}, X_k) = S_{i(n+\alpha+1)}$ we find

$$\underline{S}_{j(n+1)}^{\alpha} = S_{j(n+\alpha+1)}, S_{i(n+\alpha)}, \dots, S_{i(n+1)}$$
where $S_{j(n+\alpha+1)} = j$. (48)

Let $\underline{S}_n = (\underline{S}_{1n}^{\alpha}, \underline{S}_{2n}^{\alpha}, \dots, \underline{S}_{N_S n}^{\alpha})^T$ and the corresponding metric vector

 $\underline{M}_n = (M_{1n}, M_{2n}, \ldots, M_{N_S n})^T$. A transition from \underline{S}_n to \underline{S}_{n+1} is simply a transition of its components according to E_q . (48). Now we can write the state of the channel decoder at time n as follows:

$$R_{n} = (\underline{M}_{n-\alpha}, \mathcal{S}_{n-\alpha})$$
or
$$(49)$$

$$R_{n} = \begin{bmatrix} M_{1(n-\alpha)}, 1, S_{1(n-1)}, \dots, S_{1(n-\alpha)} \\ M_{2(n-\alpha)}, 2, S_{2(n-1)}, \dots, S_{2(n-\alpha)} \\ \vdots & \vdots & \vdots \\ \vdots & \vdots & \vdots \\ M_{N_{S}^{1}(n-\alpha)}, N_{S}, S_{N_{S}(n-1)}, \dots, S_{N_{S}(n-\alpha)} \end{bmatrix}$$

where $n \ge \alpha$. The next state R_{n+1} is just

$$R_{n+1} = (\underline{M}_{n-\alpha+1}, \underline{S}_{n-\alpha+1})$$
 (51)

Since the decoder states are derived from a truncated version of the Viterbi algorithm, the output of this machine at time n can be computed from the fact that S_n consists of all survivors with the best metrics. So we first choose $\underline{S}_{in}^{\alpha}$ so that M_{in} is the smallest component of \underline{M}_n , then the output is the input to the encoder machine that causes the transition from $S_{i(n-\alpha)}$ to $S_{i(n-\alpha+1)}$:

$$Y_{n} = X_{n-\alpha} \tag{52}$$

where $M_{\rm in}={\rm min.}\ M_{\rm n}$ such that $S_{i(n-\alpha+1)}=\sigma(X_{n-\alpha},S_{i(n-\alpha)}); M_{\rm n}$ stands for the components of $\underline{M}_{\rm n}$. From Eq. (3.32) one can easily notice that the decoding decisions always lag the progress of the decoder by an amount equal to the path length α .

It is obvious from Eq. (1) and Eq. (49) that Eq. (50) will generate more decoder states as n increases; consequently the decoder does not have a finite number of states. This is certainly an undesirable circumstance.

Consider a finite state encoder and a path of length α in its state transition diagram. Let this encoder have N_S states. Since the elements of $\underline{S}_{jn}^{\alpha}$ are drawn from a fixed finite set, the number of possible combinations of the elements of \underline{S}_n , i.e. all paths of length α , is also finite; however, the \underline{M}_n components make the number of decoder states grow to infinity since each M_{in} is nondecreasing. Fortunately, the absolute value of the \underline{M}_n components are not important. The Viterbi algorithm requires only the differences between components of \underline{M}_n . This condition allows us to keep \underline{M}_n in a fixed range. The finiteness of the decoder state will be based on this property of the Viterbi algorithm.

To maintain the differences between the \underline{M}_n components fixed while their absolute values grow is achieved by subtracting the smallest component of M_n from all components of \underline{M}_n . Hence at all times there is at least one component equal to zero. Now, if the metrics between states of the encoder only take integer values, \underline{M}_n will take only a finite number of values. For the BSC this expectation is fulfilled. In applying the Viterbi algorithm to the BSC, one finds that the algorithm essentially finds the path closest to the received word in Hamming distance.

We can choose any state as the initial state of the encoder and label it as state 1. R_0 is generated by assuming that the decoder receives a sequence of 0's of length α where $\alpha \ge N_{\rm S}-1$. So we have $S_{\rm i0}=1$ for all $i=1,2,\ldots,N_{\rm S}$. Hence we can write

$$R_{0} = \begin{bmatrix} M_{10}, 1, S_{1(\alpha-1)}, S_{1(\alpha-2)}, \dots, 1 \\ M_{20}, 1, S_{2(\alpha-1)}, S_{2(\alpha-2)}, \dots, 1 \\ \dots, \dots, \dots, \dots, 1 \\ M_{N_{S}}, N_{S}, S_{N_{S}}(\alpha-1), \dots, 1 \end{bmatrix}$$
(53)

where $S_{ij} \in \$ = (1,2,...,N_S)$ for $i = 1,1,...,N_S$ and $j = 1,2,...,\alpha-1$.

Let M_{j0} be the smallest component of the set $M_n = \{M_{i0}; i = 1, 2, ..., N_S\}$. Then the Eq. (53); can be rewritten:

$$R_{0} = \begin{bmatrix} M_{10} - M_{j0}, 1, S_{1(\alpha-1)}, S_{1(\alpha-2)}, \dots, 1 \\ M_{20} - M_{j0}, 2, S_{2(\alpha-1)}, S_{2(\alpha-2)}, \dots, 1 \\ \vdots, \dots, \dots, \dots, \dots, 1 \\ \vdots, \dots, \dots, \dots, \dots, 1 \\ 0, \dots, S_{j(\alpha-1)}, S_{j(\alpha-2)}, \dots, 1 \\ \vdots, \dots, \dots, \dots, \dots, \dots, 1 \\ M_{N_{S}0} - M_{j0}, N_{S}, S_{N_{S}}(\alpha-1), \dots, \dots, 1 \end{bmatrix}$$

$$(54)$$

For a BSC with transition probability p < 1/2, the received sequence C is binary and the log-likelihood function becomes

$$\ln Pr(C/D) = d(C, \underline{D}) \ln(p/(1-p)) + N \ln(1-p)$$
(55)

where $d(\underline{C},\underline{D})$ is the Hamming distance between \underline{C} and \underline{D} . Since when p is less than 1/2, $\ln (p/(1-p)) < 0$ and $N \ln (1-p)$ is constant for all \underline{D} , an equivalent maximum likelihood decoder for the BSC chooses \underline{D} as the codeword that minimizes the Hamming distance

$$d(\underline{C},\underline{D}) = \sum_{i} |C_{i} - D_{i}|. \tag{56}$$

Hence in applying the Viterbi algorithm to the BSC, the metric becomes an integer.

For many of DMC's positive integers can be the metric. The metric $M(C_i/D_i) = \ln Pr(C_i/D_i)$ can be replaced by b_2 ($\ln Pr(C_i/D_i) + b_1$) where b_1 is any real number and C_2 is any positive real number. A path \underline{D} that maximizes $\ln Pr(C_i/D_i)$ in fact also maximizes b_2 ($\ln Pr(C_i/D_i) + b_1$) and hence the modified metrics can be used without affecting the performance of the Viterbi algorithm.

If b_1 is chosen to make the smallest metric equal to 0, b_2 can then be chosen so that all metrics can be approximated by integers. There are many sets of integer metrics possible for a given DMC, depending on the choice of b_2 . The performance of the Viterbi algorithm is now suboptimum due to the modified metrics.

For some transition probabilities such as 1/3, 1/7, etc., even using the modified metrics, we may never have a set of integers as our metrics. From now on we restrict ourselves to the BSC only.

8 COMPUTATIONAL RESULT

Two programs, written in FORTRAN 77, were run on an TSO-IBM 3081 to obtain the results in this section. One program enumerated and identified the closed paths, the minimum free distance, and simulated the communication system while the other generated the finite state decoder.

Two examples will be shown; each has rate 1/2. Since the number of decoder states grows rapidly, only the decoder state of the first example is shown.

EXAMPLE 1

The linear finite state encoder M_x with two states is chosen because the number of decoder states is small; this makes it possible to show the state transition diagram on one page. Figure 14 shows the encoder while Figure 15 shows the corresponding finite state decoder; state 1 is chosen as the initial state. The decoder states are computationally generated in tabular form; they are shown in Table 1. The enumeration and identification of the closed paths, and the

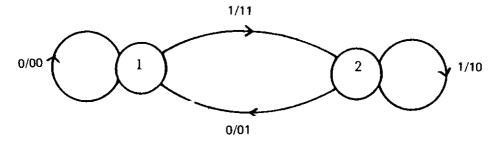


Figure 14 A linear encoder $M_{_{\rm X}}$ with two states

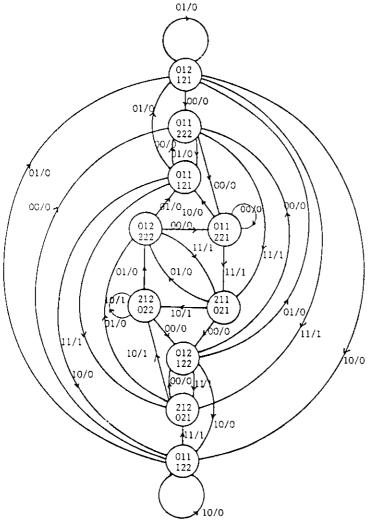


Figure 15. The state transition diagram of a finite state decoder of $M_{_{\mathrm{X}}}$

distance, are not shown here because the main purpose in this linear example is to give an idea about generating the finite state decoder.

Table 1 The next-state and the output table of the decoder M_{ν}

T:	Present		Output						
Time	State	IN = 00	IN = 01	IN = 10	IN = 11	00	01	10	11
0	011 221	011 221	011 221	011 121	211 021	0	0	0	1
1	011 121	011 222	012 121	011 122	212 021	0	0	0	1
2	211 021	012 122	012 222	212 022	012 122	0	0	1	0
3	011 222	011 221	012 121	011 121	212 021	0	O	0	1
4	012 121	011 222	012 121	011 122	212 021	0	0	0	1
5	011 122	011 222	012 121	011 122	212 021	0 0		0	1
6	212 021	012 122	012 222	212 022	012 122	0 0		1	0
7	012 122	011 222	012 121	011 122	212 021	0 0		0	1
8	012 222	011 221	011 121	011 121	211 021	0 0		0	1
9	212 022	012 122	012 222	212 022	012 122	0	0	1	O

EXAMPLE 2

Figure 16 shows an example of a finite state encoder with three states. The number of decoder states with truncated path length 2 is 79. There are too many to be shown here. The closed path enumeration and identification is computed for $\alpha = 6$. Table 2 shows the F_1 paths, the B_1 paths and the corresponding closed paths of length 6 in terms of the encoder state sequence.

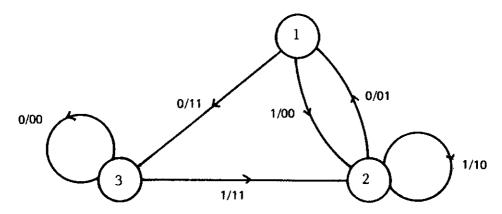


Figure 16 A finite state encoder M_{γ} with three states

Table 2 The F₁ and the B₁ paths and their corresponding closed paths of length 6

Length	1	F ₁ Paths	B ₁ Paths	Closed Paths	No
6	1	13333	121	1333321	1
	,	13332	221	1333221	2
		13321	321	1332121	3
		13322		1332221	4
		13213		1321321	5
		13212		1321221	6
		13221		1322121	7
		13222		1322221	8
1	,	12133		1213321	9
		12132		1213221	10
		12121		1212121	11
]		12122		1212221	12
		12213		1221321	13
ļ i		12212		1221221	14
		12221		1222121	15
		12222		1222221	16

Table 3 shows the distances among the closed paths of length 6 that diverge and remerge with state 1. These distances are presented in an $n \times n$ matrix form where n is the corresponding closed path; the entry (k, j) is the distance between closed path i and closed path j.

Table 3 The Hamming distances between closed paths of length 6 that diverge from and remerge with state 1

Cd. Pth.	DISTANCE															
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
1	0	3	5	4	5	4	6	5	5	8	6	5	6	5	7	6
2	3	0	4	3	4	5	5	4	8	5	5	4	5	6	6	5
3	5	4	0	3	6	5	3	6	6	5	5	8	7	6	4	7
4	4	3	3	0	5	4	6	3	5	4	8	5	6	5	7	4
5	5	4	6	5	0	3	5	4	6	5	7	6	3	6	8	7
6	4	5	5	4	3	0	4	3	5	6	6	5	6	3	7	6
7	6	5	3	6	5	4	0	3	7	6	4	7	8	7	3	6
8	5	4	6	3	4	3	3	0	6	5	7	4	7	6	6	3
9	5	8	6	5	6	5	7	6	0	3	5	4	5	4	6	5
10	8	5	5	4	5	6	6	5	3	0	4	3	4	5	5	4
11	6	5	5	8	7	6	4	7	5	4	0	3	6	5	3	6
12	5	4	8	5	6	5	7	4	4	3	3	0	5	4	6	3
13	6	5	7	6	3	6	8	7	5	4	6	5	0	3	5	4
14	5	6	6	5	6	3	7	6	4	5	5	4	3	0	4	3
15	7	6	4	7	8	7	3	6	6	5	3	6	5	4	0	3
16	6	5	7	4	7	6	6	3	5	4	6	3	4	3	3	0

The results of the simulation study are shown in Table 4; it shows the number of bit errors for a particular channel crossover probability and truncated length α given 10^4 uses of the channel.

Table 4 Number of bit errors for a particular crossover probability P and truncated length α given 10^4 uses of the channel

Р					
-,. <u></u>	α = 2	α = 3	α = 6	α = 9	
1. 0x10 ⁻²	0	1	1	1	
2. 0×10 ⁻²	3	2	2	2	
4. 0x10 ⁻²	29	29	22	22	
6. 0x10 ⁻²	79	75	56	54	
8. 0x10 ⁻²	117	104	98	101	
1. 0x10 ⁻¹	209	229	187	185	
2. 0x10 ⁻¹	831	835	754	747	

9 CONCLUSIONS

This paper studied a non-linear class of finite state digital channel coding with the aim of finding some properties of the code which improve the performance. In this case, the performance is measured by the minimum free distance since it is easier to evaluate than the probability of error criterion, and for small channel crossover probability it is a very good indicator of system performance. The analysis has indicated that the structures of the finite state machines are important to performance. In particular, the pairs of closed paths that diverge from and remerge with the same state and how we identify and improve them, in order to find a bigger minimum free distance, are very important; in addition, the catastrophic property has also been investigated, based on the machine structure. Thus, a new approach has been initiated which emphasizes on the structure imposed by modelling our encoder-decoder as a finite state machine; consequently it is not limited to only linear codes. This is in stark contrast to most existing methods of analysis which consider only the class of linear codes.

The general problem itself remains unsolved and there are many aspects of it that may be investigated. Even for the binary case much remains to be done. Although we adopted a truncated version of the Viterbi algorithm for the decoder, we assumed that the system performance could be approximated by the ideal Viterbi algorithm. The analysis of the loss in performance caused by the truncation strategy remains untouched. Consequently, there is a room to improve our bound on bit error probability; a random coding approach could also be used to deal with the error probability measure. Another assumption in our present analysis is that of perfect synchronization. Finding a pair of finite state encoder-decoder that always synchronize is another problem. We mentioned and proved some Markov chain properties, yet we did not exploit these powerful properties. These properties, along with renewal theory, may be useful for the synchronization analysis. The use of a bidirectional search to identify the critical closed paths seems to be helpful. Of course, by the very nature of our algorithm, we can not use this search algorithm if the number of the encoder states is too large. This is a classical problem that remains unresolved when one deals with a graph. In addition, we have not considered the case when the metrics are real numbers. This problem seems much more difficult than the loss of performance due to truncation and code synchronization. We do not know how to deal with this general problem. We leave this as an open problem for further investigations.

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APPENDIX A

With the notation of section 2 we want to show that (S_n, R_n) is a first order Markov Chain for both the noiseless channel and the additive channel: $C_n = D_n + N_n$ where + denotes a mod. N_C addition and the (N_n) are a sequence of independent, identically distributed, random variables.

Proof:

a. Noiseless channel case:

$$\begin{split} & Pr\left[\left. S_{\mathbf{n}} = s_{\mathbf{n}} \right. \left. \left. \left. \left. \left. \left. \left. \left. \left. \left(S_{i}, R_{i} \right) = (s_{i}, r_{i}), i < n - 1 \right. \right] \right. \right. \right. \\ & = Pr\left[\left. \left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho(D_{\mathbf{n}-1}, r_{\mathbf{n}-1}) = r_{\mathbf{n}} \right. \left(\left. \left. \left(S_{i}, R_{i} \right) = (s_{i}, r_{i}), i \leq n - 1 \right. \right] \right. \\ & = Pr\left[\left. \left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) = r_{\mathbf{n}} \right| \left. \left(S_{i}, R_{i} \right) = (s_{i}, r_{i}), i \leq n - 1 \right. \right] \right. \\ & = Pr\left[\left. \left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) = r_{\mathbf{n}} \right. \right] \right. \\ & = Pr\left[\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) = r_{\mathbf{n}} \right. \right] \right. \\ & = Pr\left[\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) = r_{\mathbf{n}} \right. \right] \right. \\ & = Pr\left[\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) = r_{\mathbf{n}} \right. \right] \right. \\ & = Pr\left[\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) = r_{\mathbf{n}} \right. \right] \right] \right. \\ & = Pr\left[\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) = r_{\mathbf{n}} \right. \right] \right] \right] \right] \\ & = Pr\left[\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) \right. \right] \right] \right] \right] \right] \\ & = Pr\left[\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) = s_{\mathbf{n}}, \rho\left(\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right), r_{\mathbf{n}-1} \right) \right. \right] \right] \right] \right] \right] \\ & = Pr\left[\left. \left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \right] \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \\ & = Pr\left[\left. \left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1} \right) \right] \\ & = Pr\left[\left(X_{\mathbf{n}-1}, s_{\mathbf{n}-1}$$

Hence:

$$Pr[S_n = s_n \& R_n = r_n | S_i = s_i, R_i = r_i, i \le n - 1]$$

= $Pr[S_n = s_n \& R_n = r_n | S_{n-1} = s_{n-1}, R_{n-1} = r_{n-1}]$
which completes the proof.

b. Additive channel:

 $C_n = D_n + N_n$ where + denotes a mod. N_C addition and the N_n are independent, identically distributed random variables;

$$Pr[S_n = S_n \& R_n = r_n \mid (S_i, R_i) = (s_i, r_i), i \le n - 1]$$

By using our finite state decoder model one can easily get:

$$= Pr \left[\sigma \left(X_{n-1}, s_{n-1} \right) = s_n, \rho \left(C_{n-1}, r_{n-1} \right) = r_n + (s_i, r_i) = (s_i, m_i), i \le n-1 \right]$$

$$= Pr \left[\sigma \left(X_{n-1}, s_{n-1} \right) = s_n, \rho \left(\delta \left(X_{n-1}, s_{n-1} \right) + N_{n-1}, r_{n-1} \right) = r_n \right]$$

$$\left(S_i, R_i \right) = (s_i, r_i), i \le n-1 \right]$$

$$= Pr \left[\sigma \left(X_{n-1}, s_{n-1} \right) = s_n, \rho \left(\delta \left(X_{n-1}, s_{n-1} \right) + N_{n-1}, r_{n-1} \right) = r_n \right]$$

$$since S_i \text{ and } R_i \text{ are functions of only } X_1, \dots, X_{i-1} \text{ and } N_1, \dots, N_{i-1}. \text{ Hence:}$$

$$= Pr \left[S_n = s_n \& R_n = r_n + S_{n-1} = s_{n-1}, R_{n-1} = r_{n-1} \right]$$

which completes the proof.

APPENDIX B

With the notation of section 2 we want to show that (D_n, S_n, Y_n, R_n) is a first order Markov Chain for both a noiseless channel: $C_n = D_n$, and an additive channel: $C_n = D_n + N_n$ where + denotes a mod. N_C addition and the (N_n) are independent identically distributed random variables.

Proof:

a. For the noiseless channel: $C_n = D_n$,

$$\begin{split} & Pr[D_{\mathbf{n}} = d_{\mathbf{n}}, S_{\mathbf{n}} = s_{\mathbf{n}}, Y_{\mathbf{n}} = y_{\mathbf{n}} \ R_{\mathbf{n}} = r_{\mathbf{n}} | (D_{\mathbf{i}}, S_{\mathbf{i}}, Y_{\mathbf{i}}, R_{\mathbf{i}}) = (d_{\mathbf{i}}, s_{\mathbf{i}}, y_{\mathbf{i}}, r_{\mathbf{i}}), i \leq n-1] \\ & = Pr[D_{\mathbf{n}} = d_{\mathbf{n}}, \ S_{\mathbf{n}} = s_{\mathbf{n}}, \ Y_{\mathbf{n}} = y_{\mathbf{n}}, \ R_{\mathbf{n}} = r_{\mathbf{n}} | D_{\mathbf{n}-1} = d_{\mathbf{n}-1}, \ Y_{\mathbf{n}-1} = y_{\mathbf{n}-1}, \\ & (D_{\mathbf{j}}, \ Y_{\mathbf{j}}) = (d_{\mathbf{j}}, y_{\mathbf{j}}), j \leq n-2, (S_{\mathbf{i}}, \ R_{\mathbf{i}}) = (s_{\mathbf{i}}, \ r_{\mathbf{i}}), i \leq n-1] \end{split}$$

By using the finite state model one finds:

Hence:

$$\begin{aligned} & Pr[D_n = d_n, \, S_n = s_n, \, Y_n = y_n, \, R_n = r_n \, | D_i = d_i, \, S_i = s_i, \, Y_i = y_i, \, R_i = r_i, i \leq n-1] \\ & = Pr[D_n = d_n, \, S_n = s_n, \, Y_n = y_n, \, R_n = r_n \, | D_{n-1} = d_{n-1}, \, S_{n-1} = s_{n-1}, \\ & Y_{n-1} = y_{n-1}, \, R_{n-1} = r_{n-1}] \\ & \text{which completes the proof.} \end{aligned}$$

b. For additive channel:

 $C_n = D_n + N_n$ where + denotes a mod. N_C addition and the N_n are independent identically distributed random variables.

$$\begin{split} & Pr[D_{n} = d_{n}, S_{n} = s_{n}, Y_{n} = y_{n}, R_{n} = r_{n} | (D_{i}, S_{i}, Y_{i}, R_{i}) = (d_{i}, s_{i}, y_{i}, r_{i}), i < n - 1] \\ & = Pr[D_{n} = d_{n}, S_{n} = s_{n}, Y_{n} = y_{n}, R_{n} = r_{n} | D_{n-1} = d_{n-1}, Y_{n-1} = y_{n-1}, D_{j} = d_{j}, Y_{j} = y_{j}, j \le n - 2, S_{i} = s_{i}, R_{i} = r_{i}, i \le n - 1] \end{split}$$

By using the finite state model one finds:

=
$$Pr[\delta(X_n, \sigma(X_{n-1}, s_{n-1})) = d_n, \sigma(X_{n-1}, s_{n-1}) = s_n,$$

$$\begin{array}{l} \eta \; (\; \delta \; (X_{\rm n}, \sigma (X_{\rm n-1}, s_{\rm n-1})) + N_{\rm n}, \rho (d_{\rm n-1}, r_{\rm n-1})) = y_{\rm n}, \\ \rho (d_{\rm n-1} + N_{\rm n-1}, r_{\rm n-1}) = r_{\rm n} \; | \; D_{\rm n-1} = d_{\rm n-1}, \; Y_{\rm n-1} = y_{\rm n-1}] \; , \\ {\rm since} \; X_{\rm n} \; {\rm and} \; X_{\rm n-1} \; {\rm are} \; {\rm independent} \; {\rm of} \; X_{\rm k} \; {\rm and} \; N_{\rm k} \; {\rm for} \; k < n-2 \; {\rm and} \; {\rm also} \; \\ {\rm independent} \; {\rm of} \; S_{\rm m} \; {\rm and} \; R_{\rm m} \; {\rm for} \; {\rm all} \; m. \end{array}$$

Hence:

$$Pr[D_n = d_n, S_n = s_n, Y_n = y_n, R_n = r_n \mid D_{n-1} = d_{n-1}, S_{n-1} = s_{n-1}, Y_{n-1} = y_{n-1}, R_{n-1} = r_{n-1}]$$
 which completes the proof.