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Coding for a Binary Independent Piecewise-Identically-Distributed Source

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Abstract—Two weighting procedures are presented for compaction of output sequences generated by binary independent sources whose unknown parameter may occasionally change. The resulting codes need no knowledge of the sequence length, i.e., they are strongly sequential, and also the number of parameter changes is unrestricted. The additional-storage and computational complexity of this method are less than \( 1/2 \log T \) bits per transition. For the second method we could prove that additional-transition redundancy is not more than \( 1/2 \log T \) bits per transition, which is more than the Merhav bound; however, the storage and computational complexity of this method are also more interesting than those of the first method. Simulations show that the difference in redundancy performance between the two methods is negligible.

Index Terms—Universal noiseless source coding, independent piecewise-identically-distributed source, memoryless source, transition pattern, weighting, arithmetic coding, redundancy bounds.

I. INTRODUCTION

The Elias Algorithm (described in, e.g., Jelinek [1]) produces for any coding distribution \( P_e(x_t^T) \) over all binary sequences \( x_t^T = x_1x_2\cdots x_T \in \{0,1\}^T \), a binary prefix code with codeword lengths \( L(x_t^T) \) such that

\[
L(x_t^T) < - \frac{1}{P_e(x_t^T)} + 2 \quad \text{for all possible } x_t^T. \tag{1}
\]

Possible sequences are sequences that can actually occur, i.e., sequences \( x_t^T \) with actual probability \( P_a(x_t^T) > 0 \). It is required that the coding distribution \( P_e(\cdot) \) satisfies

\[
P_e(\emptyset) = 1, \quad P_e(x_t^T-1) = P_a(x_t^{T-1}, X_t = 0) + P_e(x_t^{T-1}, X_t = 1), \quad \text{for all } x_t^{T-1} \in \{0,1\}^{T-1}, t = 1, \ldots, T, \text{ and } P_e(x_T^T) > 0 \quad \text{for all possible } x_T^T \in \{0,1\}^T. \tag{2}
\]

Note that \( \emptyset \) stands for the empty sequence \( (x_2) \). If the marginals \( P_a(x_t^T), t = 1, \ldots, T \) are sequentially available the arithmetic code can be implemented sequentially. After having accepted a coding redundancy of at most 2 bits, we are left with the problem of finding good coding distributions \( P_e(\cdot) \) for the cases we are interested in.

For binary independent and identically distributed (i.i.d.) sources with unknown parameter \( \theta \) (i.e., the probability of generating a 1), we can assign the block probability \( P_e(x_t^T) = P_e(a, b) \) to a sequence \( x_t^T \) containing \( a \) zeros and \( b \) ones where

\[
P_e(a, b) \triangleq \frac{1}{\pi \sqrt{(1-\theta)^\theta (1-\theta)^{1-\theta}}} \int_0^1 \frac{1}{\sqrt{(1-\theta)\theta}} (1-\theta)^a \theta^b \, d\theta. \tag{3}
\]

This distribution allows sequential updating, i.e., \( P_e(0, 0) = 1 \), and for \( a \geq 0 \) and \( b \geq 0 \)

\[
P_e(a+1, b) = \frac{a+1/2}{a+b+1} \cdot P_e(a, b)
\]

and

\[
P_e(a, b+1) = \frac{b+1/2}{a+b+1} \cdot P_e(a, b). \tag{4}
\]

It was suggested by Krichevsky and Trofimov [2] and satisfies (see [10]), for \( a + b \geq 1 \), the inequality

\[
P_e(a, b) \leq \frac{1}{2} \left( \frac{1}{\sqrt{a+b}} \right)^a \left( \frac{b}{a+b} \right)^b. \tag{5}
\]

Using this inequality we can easily upper-bound the parameter redundancy resulting from this estimator. If \( x_t^T \) is a (nonempty) sequence containing \( a \) zeros and \( b \) ones, with actual probability \( P_a(x_t^T|\theta) \), we obtain for the parameter redundancy relative to the actual source that

\[
\log \frac{P_e(x_t^T|\theta)}{P_e(x_t^T)} = \log \frac{(1-\theta)^a \theta^b}{P_e(a, b)} \leq \log \frac{1}{2} \cdot \frac{1}{\sqrt{a+b}} \left( \frac{a}{a+b} \right)^a \left( \frac{b}{a+b} \right)^b \leq \frac{1}{2} \log T + 1, \quad \text{for all } \theta \in [0,1]. \tag{6}
\]

In other words, the parameter redundancy is uniformly bounded. The inequality in (6) follows from applying the lower bound in (5) for the denominator and observing that the maximum value of the enumerator is achieved for \( \theta = b/(a+b) \). In a more general setting the memoryless source is not identically distributed. We assume in this correspondence that the source parameter \( \theta \) changes occasionally.

Definition 1: An independent piecewise-identically-distributed (i.p.i.d.) binary source generates a sequence \( x_t^T \in \{0,1\}^T \) of independent symbols. During this generation process, the source parameter \( \theta \) makes, say \( t \) transitions and assumes \( C+1 \) values. The transition pattern \( T = (t_1, t_2, \ldots, t_C) \) for sequence \( x_t^T \) indicates that the symbols \( x_{t+1}, x_{t+2}, \ldots, x_{t+t_C-1} \) are generated according to parameter \( \theta_c \), for \( c = 0, C \). We assume here that \( t_0 \geq 1 \) and \( t_{C+1} \geq T+1 \). Note that \( t_{C+1} > t_0 \) for \( c = 0, C \) and, consequently, \( C \leq T-1 \). We can now write for the actual probability

\[
P_e(x_t^T|T, \Theta) = \prod_{i=1}^T P_e(X_t = x_i)
\]

where

\[
P_e(X_t = 1) = 1 - P_e(X_t = 0) = \theta_c, \quad \text{if } t_0 \leq t < t_{c+1}, \quad \text{for } c = 0, 1, \ldots, C. \tag{7}
\]

The vector consisting of the parameters \( \theta_0, \theta_1, \ldots, \theta_C \) is denoted by \( \Theta \) and called the parameter vector. We write \( C(T) \) for the number of transitions \( C \) in pattern \( T \).

Example: Our source generates 1000 binary digits. The starting value of the parameter of our example source is 0.9. This parameter changes into 0.0 at \( t = 301 \). At \( t = 401 \) the parameter value becomes 0.5. Finally, at \( t = 501 \) there is a transition again and
the new value becomes 0.2. The parameter now remains constant until the end of the sequence. This parameter behavior is plotted in Fig. 1. The number of transitions is three, the transition pattern is (301, 401, 601), the parameter vector (0.9, 0.0, 0.5, 0.2).

If we know the actual transition pattern \( T \), we can partition the sequence \( x^T \) into \( C(T) + 1 \) memoryless subsequences and use

\[
P_c(x^T) = \prod_{c=0}^{C(T)} P_c(a(x_{t}^{e+1}), b(x_{t}^{e+1}))
\]

as a coding distribution, where \( a(x_t) \) (resp., \( b(x_t) \)) is the number of zeros (resp., ones) that occur in \( x_t = x_t x_{t+1} \cdots x_j \). Again this coding distribution allows sequential updating. For any sequence \( x^T \in \{0,1\}^T \), using (6) and the convexity of the \( \log(\cdot) \), the parameter redundancy with respect to the actual transition pattern \( T \) and the actual parameter vector \( \Theta \), can be upper bounded as

\[
\log \frac{P_c(x^T | T)}{P_c(x^T)} \leq \frac{C(T) + 1}{2} \log \left( \frac{T}{C(T) + 1} \right) + C(T) + 1,
\]

for all \( \Theta \in [0,1]^{C(T)+1} \). (9)

Note that we do not need a function \( \gamma(\cdot) \) as in [10] here since always \( t_{e+1} > t_e \) for \( c = 0, C(T) \).

If the transition pattern is unknown we can weight the coding distributions corresponding to all transition patterns \( T \) of sequences of length \( T \) and obtain the weighted coding distribution

\[
P_c(x^T) = \sum_T P_T(T) P_c(x^T | T).
\]

Here \( P_T(T) \) is the \textit{a priori} probability that is assigned to transition pattern \( T \). Of course

\[
\sum_T P_T(T) = 1.
\]

For any sequence \( x^T \in \{0,1\}^T \) the transition redundancy with respect to pattern \( T \) can be upper-bounded as

\[
\log \frac{P_c(x^T | T)}{P_c(x^T)} \leq \log \frac{1}{P_T(T)} \quad \text{for all } T
\]

simply by noting that \( P_c(x^T) \geq P_T(T) P_c(x^T | T) \).

The total cumulative redundancy relative to the actual pattern \( T \) and actual vector \( \Theta \) is equal to the sum of the (cumulative) transition, parameter and coding redundancies. Using (1), (9), and (10) we can upper-bound this total redundancy for any sequence \( x^T \) in the following way:

\[
L(x^T) = \log \frac{1}{P_c(x^T | T, \Theta)} \leq \log \frac{P_c(x^T | T)}{P_c(x^T)} + \log \frac{P_T(x^T | T)}{P_T(x^T)} + L(x^T) - \log \frac{1}{P_T(x^T)}
\]

\[
< \log \frac{1}{P_T(T)} + C(T) + 1 \log \left( \frac{T}{C(T) + 1} \right) + C(T) + 3. \quad (11)
\]

This holds for all transition patterns \( T \) of \( x^T \) and parameters \( \Theta \in [0,1]^{C(T)+1} \). Rewriting this bound, and taking the minimum over all transition patterns \( T \) of \( x^T \) and all parameter vectors \( \Theta \), we obtain for all \( x^T \in \{0,1\}^T \) (not necessarily being generated by the source with transition pattern \( T \) and parameter vector \( \Theta \) that

\[
L(x^T) < \min_{\Theta} \left\{ \log \frac{1}{P_c(x^T | T, \Theta)} + \log \frac{1}{P_T(T)} + C(T) + 1 \log \left( \frac{T}{C(T) + 1} \right) + C(T) + 3 \right\}. \quad (12)
\]

From this, we may conclude that weighting methods try to minimize the \textit{total description length} of a sequence. In the following sections we will describe certain \textit{a priori} distributions over the transition patterns, and investigate the coding methods that are based on these distributions.

There is a vast amount of publications in the image, audio, and speech processing literature on segmentation problems. Although at first sight the problem treated here is similar to all these segmentation problems, the approach taken here is entirely different. Our objective is not to determine the boundary points of the (stationary) segments but simply to compress the entire sequence. It is a bit surprising to realize that also in the image, audio, and speech processing literature compression is often the primary goal. However, it seems advantageous to replace this primary objective by a secondary objective, namely, to determine segments of constant statistics.

II. A QUADRATICAL-COMPLEXITY CODING METHOD

We will first define a coding distribution over source sequences and transition patterns that have infinite length in principle. We will see later that the storage complexity resulting from this distribution increases quadratically with the sequence length \( T \). Therefore, we refer to this method as the \textit{quadratical-complexity coding method}.
We can define the distribution after assuming that the source follows a path in a quadratic transition diagram (see Fig. 2).

**Definition 2.** The source starts in state $(1, 0, 1)$ of the quadratic transition diagram. It generates the first output $x_1$ according to parameter $\theta_0$. When the source produces output $x_1$, it is in a state $(t, c, t_e)$, which indicates that the number of parameter changes encountered so far is $c$, that the current parameter is $\theta_0$, and that the last time index before which a change occurred is $t_e$. To obtain Krichevsky-Trofimov estimators we assign, in state $(t, c, t_e)$, the following probabilities to the values 0 and 1 that can be generated:

\[
P_c(X_t = 0 | (t, c, t_e)) = \frac{a(x_t^{-1} - 1)}{t - t_e + 1},
\]

\[
P_c(X_t = 1 | (t, c, t_e)) = \frac{b(x_t^{-1} + 1)}{t - t_e + 1},
\]

(13)

Again, $a(x_t^{-1})$ (resp., $b(x_t^{-1})$) is the number of zeros (resp., ones) that occur in $x_t = x_t x_{t+1} \cdots x_j$. After having generated $x_t$, the parameter can change and the source will go to state $(t+1, c+1, t+1)$. If the parameter remains the same, the next state of the source will be $(t+1, c, t_e)$. We assign the following probabilities to these two alternatives to obtain Krichevsky-Trofimov estimators:

\[
P_c((t + 1, c, t_e) | (t, c, t_e)) = \frac{t - c - 1}{t},
\]

\[
P_c((t + 1, c + 1, t + 1) | (t, c, t_e)) = \frac{c + 1}{t + 1}.
\]

(14)

We should note that state $(t, c, t_e)$ only exists if $0 \leq c \leq t - 1$ and if $c + 1 \leq t_e \leq t$ for $c \neq 0$ and $t_0 = 1$. The source output $x_t$ and the next state $(t+1, p, q)$, where $(p, q) = (c, t_e)$ or $(c+1, t+1)$ are independent of each other given the current state $(t, c, t_e)$. Note that the conditional coding probabilities in (13) do not depend on $c$ while the conditional transition probabilities in (14) do not depend on $t_e$.

We can now specify the quadratic-complexity coding distribution over all source sequences.

**Definition 3.** For the quadratic method, for each sequence $x_T \in \{0, 1\}^T$ we compute for all $x_1 \in \{0, 1\}^T$, $t = 1, T$, the probabilities $P_c(x_1 | (t, c, t_e))$ recursively, i.e.,

\[
P_c(x_1, (t, p, q)) \triangleq \sum_{s, r} P_c(x_1^{-1}, (t-1, r, s)) \cdot P_c((t, p, q) | (t-1, r, s)) P_c(x_1(t, p, q))
\]

(15)

for $t > 1$, and $P_c(x_1(1, 0, 1)) = 1/2$ for both $x_1 = 0$ and 1. The summation is only over $r, s$ such that $(t - 1, r, s)$ is an existing state in the diagram (connected to state $(t, p, q)$).

The coding probabilities $P_c(x_1^T)$, $x_1^T \in \{0, 1\}^T$, $t = 1, T$, are now defined as follows:

\[
P_c(x_1^T) \triangleq \sum_{p, q} P_c(x_1^T, (t, p, q)).
\]

(16)

The summation is only over $p, q$ such that $(t, p, q)$ is an existing state in the transition diagram. Of course, $P_c(\emptyset) = 1$.

It is easy to show that the coding probabilities $P_c(x_1^T)$, $t = 0, T$, satisfy the restrictions (2). After having discussed the implementation of this method it will be clear that this coding distribution is sequentially available.

It is possible to demonstrate that this coding distribution performs a weighting of all coding distributions $P_c(x_1^T | T)$ over all transition patterns $T$ over $T$ source symbols, i.e.,

\[
P_c(x_1^T) = \sum_{p, q} \sum_{T \rightarrow (T,p,q)} P_c(T) P_c(x_1^T | T)
\]

(17)

where

\[
\sum_{T \rightarrow (T,p,q)} P_c(T) P_c(x_1^T | T)
\]

means a summation over all $T$ that lead to state $(T, p, q)$. Furthermore, for a transition pattern $T$ such that $T \rightarrow (T, p, q)$, i.e., a pattern $T$ that leads to $(T, p, q)$

\[
P_{\text{tr}}(T) = P_c(T - 1 - p, p).
\]

(18)

Probability $P_c(x_1^T | T)$ is as defined in (8). This result is a direct consequence of the following lemma.

**Lemma 1.** For $t = 1$, $T$ and any existing state $(t, p, q)$ in the quadratic transition diagram and all $x_1 \in \{0, 1\}^T$

\[
P_c(x_1^T (t, p, q)) = \sum_{T \rightarrow (T,p,q)} P_{\text{tr}}(T) P_c(x_1^T | T)
\]

(19)

with

\[
P_{\text{tr}}(T) = P_c(t - 1 - p, p)
\]

(20)

and

\[
P_c(x_1^T | T) = \left( \prod_{c=0}^{p-1} P_c(a(x_{t+1}^{-1}), b(x_{t+1}^{-1})) \right) \cdot P_c(a(x_{t+1}), b(x_{t+1})).
\]

(21)

**Proof:** We will prove the lemma by induction. First note that the hypothesis holds for $t = 1$. For $t = 1$ only the state $(0, 0, 1)$ exists and $P_{\text{tr}}(1, 0, 1) = P_c(0, 0) = 1$ and $P_c(x_1(1, 0, 1)) = 1/2$ for both $x_1 = 0$ and 1. Next we assume that the hypothesis holds for $t - 1$ when $t = 2, T$. For $t$ we then obtain using (13) and the hypothesis that

\[
P_c(x_1^T (t, p, q)) = \sum_{r,s} \sum_{S \rightarrow (t-1, r, s)} P_{\text{tr}}(S) P_c(x_1^T | S) \cdot P_c((t, p, q) | (t-1, r, s)) P_c(x_1(t, p, q))
\]

(22)

where we have used that

\[
P_{\text{tr}}(S) P_c((t, p, q) | (t-1, r, s)) = P_{\text{tr}}(S, (t, p, q)) = P_{\text{tr}}(T)
\]

and that

\[
P_c(x_1^T | S) P_c(x_1(t, p, q)) = P_c(x_1^{t-1}, x_1 | S, (t, p, q)) = P_c(x_1^T | T)
\]

for

\[
S = (S, (t, p, q)) \quad \text{and} \quad S \rightarrow (t-1, r, s).
\]

To see that $P_{\text{tr}}(T) = P_c(t - 1 - p, p)$ as stated in (20), note that from the induction hypothesis we know that $P_c(S) = P_c(t-2-r, r)$. 

---

Fig. 2. Quadratical transition diagram for $t = 1, 2, 3, 4$. State $(t, c, t_e)$ is denoted as $tct_e$. 

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Now if \( p = r \) then no new parameter change occurred going from \((t - 1, r, s)\) to \((t, p, q)\) and \(P_x(S)\) is multiplied by \(
abla(t - 1 - r - 1/2)/(t - 1)\) and we obtain

\[
P_x(T) = P_x(t - 2 - r, r) \cdot (t - 1 - r - 1/2)/(t - 1) \\
= P_x(t - 1 - r, r) \\
= P_x(t - 1 - p, p).
\]

Alternatively, if \( p = r + 1 \) then a transition occurred. The multiplication factor is now \((r + 1)/2)/(t - 1)\) and we get

\[
P_x(T) = P_x(t - 2 - r, r) \cdot (r + 1)/2)/(t - 1) \\
= P_x(t - 2 - r, r + 1) \\
= P_x(t - 1 - p, p).
\]

Furthermore, (21) can be understood by noting first that by the induction hypothesis

\[
P_x(x_i^{-1} \mid S) = \left( \prod_{c=0 \atop x_i \neq 0}^{t-r+1} P_x(a(x_i^{t+1}), b(x_i^{t+1})) \right) \\
\cdot P_x(a(x_i^{t+1}), b(x_i^{t+1})).
\]

We assume that \( x_i = 0 \), but similar remarks apply for \( x_i = 1 \).

If \( p = r \) then no new parameter transition occurred going from \((t - 1, r, s)\) to \((t, p, q)\) and \(P_x(x_i^{-1} \mid S)\) is multiplied by

\[
(a(x_i^{t+1}) + 1/2)/(t - t + 1)
\]

and we obtain

\[
P_x(x_i \mid T) = \left( \prod_{c=0 \atop x_i \neq 0}^{t-r+1} P_x(a(x_i^{t+1}), b(x_i^{t+1})) \right) \\
\cdot P_x(a(x_i^{t+1}), b(x_i^{t+1})) \\
\cdot (a(x_i^{t+1}) + 1/2)/(t - t + 1) \\
= \left( \prod_{c=0 \atop x_i \neq 0}^{t-r+1} P_x(a(x_i^{t+1}), b(x_i^{t+1})) \right) \\
\cdot P_x(a(x_i^{t+1}) + 1, b(x_i^{t+1})) \\
= \left( \prod_{c=0 \atop x_i \neq 0}^{t-r+1} P_x(a(x_i^{t+1}), b(x_i^{t+1})) \right) \\
\cdot P_x(a(x_i^{t+1}) + 1, b(x_i^{t+1})) \\
\cdot P_x(a(x_i^{t+1}) + 1, b(x_i^{t+1})) \\
\cdot P_x(a(x_i^{t+1}) + 1, b(x_i^{t+1})).
\]

On the other hand, when a transition occurred \( p = r + 1 \) and \( t_{r+1} = t \).

We rewrite

\[
P_x(x_i^{-1} \mid S) = \prod_{c=0 \atop x_i \neq 0}^{t-r} P_x(a(x_i^{t+1}), b(x_i^{t+1})).
\]

The multiplication factor \(P_x(X_t = 0 \mid (t, r + 1, t))\) is 1/2 and we find for

\[
P_x(x_i \mid T) = \left( \prod_{c=0 \atop x_i \neq 0}^{t-r} P_x(a(x_i^{t+1}), b(x_i^{t+1})) \right) \\
\cdot P_x(a(x_i^{t+1}), b(x_i^{t+1})) \\
\cdot (a(x_i^{t+1}) + 1, b(x_i^{t+1})) \\
= \left( \prod_{c=0 \atop x_i \neq 0}^{t-r} P_x(a(x_i^{t+1}), b(x_i^{t+1})) \right) \\
\cdot P_x(a(x_i^{t+1}), b(x_i^{t+1})) \\
\cdot P_x(a(x_i^{t+1}), b(x_i^{t+1})).
\]

\textbf{Theorem 1:} Let \( T \geq 2 \). If we use the quadratical-complexity coding distribution defined in (16), the transition redundancy for any \( x_i \) with respect to transition pattern \( T \) satisfies the upper bound

\[
\frac{\log P_x(x_i \mid T)}{\log P_x(x_i \mid S)} \leq C(T) \log \left( \frac{(T - 1)e}{C(T)} + \frac{1}{2} \log (T - 1) + 1 \right)
\]

for all transition patterns \( T \) over \( T \) symbols. The bound holds also for \( C(T) = 0 \) if we follow the convention \( z \log z = 0 \) for \( z = 0 \).

\textbf{Proof:} From (18) we know that

\[
P_x(T) = P_x(T - 1 - C(T), C(T))
\]

if \( T \) is a transition pattern when the source generates \( T \) symbols. In order to find a useful bound for \( \log P_x(T) \) observe first that for nonnegative integers \( u \) and \( v \) with \( u + v \geq 1 \) by the bound in (5) we get

\[
\log P_x(u, v) \geq u \log \frac{u}{u + v} + v \log \frac{v}{u + v} - \frac{1}{2} \log (u + v) - 1.
\]

For the first term on the right-hand side of this inequality, we can write, using the basic inequality \( \log x \leq 1 - x^{-1} \), that

\[
u \log \frac{u}{u + v} = \frac{-u}{2} \log \frac{u + v}{u} = \frac{-u}{2} \log e.
\]

Assuming that \( u \log u = 0 \) for \( u = 0 \), we can see that this inequality also holds for \( u = 0 \). Combining all this leads to

\[
\log P_x(u, v) \geq -v \log \left( \frac{u + v}{v} \right) - \frac{1}{2} \log (u + v) - 1.
\]

For any sequence \( x_i \in \{0, 1\}^T \) the transition redundancy can now be upper-bounded as in (10), i.e.,

\[
\log \frac{P_x(x_i \mid T)}{P_x(x_i \mid S)} \leq \log \frac{1}{P_x(T)}, \text{ for all } T.
\]

Substitution of \( u = T - 1 - C(T) \) and \( v = C(T) \) in (27) yields the theorem. Note that \( u + v = T - 1 \geq 1 \).

The theorem says that the transition redundancy is bounded by a bias term which is roughly \( \frac{1}{2} \log T \) bits and then for each transition roughly \log T additional bits. The bias term can be considered a consequence of not knowing in advance the number of transitions in the sequence. The additional terms seem to be necessary since the decoder needs (approximations of) the coordinates of the transitions.

\textbf{Example (Continued):} The transition redundancy for our example source generating the sequence \( x_i \), with \( T = 1000 \) is, according to Theorem 1, upper-bounded by

\[
3 \log \frac{999e}{3} + \frac{1}{2} \log 999 + 1 = 35.45 \text{ bits.}
\]

For the generated sequence we have also \textit{computed} the transition redundancy

\[
\log \left( \frac{P_x(x_i \mid T)}{P_x(x_i)} \right), \text{ for } t = 1, T.
\]

This redundancy is plotted in Fig. 3. It satisfies the bound 35.45 bits as we can see. Moreover, we see that for each transition we get a sharp increase of this redundancy. We can explain this by noting that the decoder must be informed almost immediately about these transitions and this requires roughly \log t bits for a transition that occurred at time \( t \). It may not be necessary, however, to transmit the exact time to the decoder, especially if the parameter change is small.
After this the records twice the number of records that are stored for processing between the terms corresponding to $s$ are determined using the counts $a(x_{t-1}^0)$ and stored in nodes $(t,p,q)$. Summing over all these nodes as in (16) gives $P_c(x_t)$. The counts $a(x_{t-1}^0)$ and $b(x_{t-1}^0)$ in record $(t,p,q)$ are determined using the counts $a(x_{t-1}^0)$ and $b(x_{t-1}^0)$ in record $(t-1,p,q)$. For example, if $x_t = 1$ count $a(x_t^0) = a(x_{t-1}^0)$ and $b(x_t^0) = b(x_{t-1}^0) + 1$ for all $q < t$ and $a(x_t^0) = 0$ and $b(x_t^0) = 1$ for $q = t$. If $x_t = 0$, we increase the $a$-counts instead of the $b$-counts. After this the records $(t-1,r,s)$ can be deleted.

Storage Complexity: Since for processing symbol $x_t$ there are $(t^2 - t + 2)/2$ nodes that need to be stored (while the nodes corresponding to $x_{t-1}$ are deleted), the instantaneous storage complexity grows quadratically in $T$.

Computational Complexity: To determine the total number of computations (multiplications) to process $x_T^n$, note that there are two ways to leave the record $(t-1,r,s)$. Each of them involves a multiplication. If a transition is assumed, the next record is $(t,r+1,s)$, otherwise, the next record is $(t,r,s)$. The number of computations related to processing $x_t$ for $t > 1$ is, therefore, twice the number of records that are stored for processing $x_{t-1}$, i.e., $(t - 1)^2 - (t - 1) + 2 = t^2 - 3t + 4$. The total number of computations for processing $x_T^n$ follows from summing the terms corresponding to $x_t$ for $t = 2,T$, which appears to be $(T^3 - 3T^2 + 8T - 6)/3$. Therefore, there is a cubic dependency between the total computational complexity and the source sequence length $T$.

III. A LINEAR-COMPLEXITY CODING METHOD

The quadratical-complexity coding method gives a nice performance as we have seen in the previous section, but the complexity of this method is also quite large. Therefore, we will discuss a method here that has a storage complexity which is linear in the sequence length $T$. Although in theory the (transition) redundancy resulting from this method is higher than the redundancy for the quadratical method, the differences turn out to be small in practise. We will now assume that the source follows a path in a linear transition diagram (see Fig. 4).

Definition 4: The source now starts in state $(1,1)$ of the linear transition diagram and generates the first output $x_1$ according to parameter $\theta_0$. When the source produces output $x_t$, it is in a state $(t,c)$, which indicates that the last time index before which a change occurred is $t_c$. The current parameter is $\theta_c$. As in (13), we assign the following probabilities to the values 0 and 1 that can be generated:

\[
P_c(X_t = 0|t, t_c) = \frac{a(x_{t-1}^0) + 1/2}{t - t_c + 1}, \quad P_c(X_t = 1|t, t_c) = \frac{b(x_{t-1}^0) + 1/2}{t - t_c + 1}.\]

(29)

After having generated $x_t$, the parameter can change and the source will go to state $(t + 1,t_c)$. If the parameter remains the same, the next state of the source will be $(t + 1, t_c)$. We now assign the following probabilities to these two alternatives to obtain the Krichevsky–Trofimov estimators:

\[
P_k((t + 1, t_c)|(t, t_c)) = \frac{t - t_c + 1/2}{t - t_c + 1}, \quad P_k((t + 1, t + 1)|(t, t_c)) = \frac{1/2}{t - t_c + 1}.
\]

(30)

where we should note that always $1 \leq t_c \leq t$. The source output $x_t$ and the next state $(t + 1,q)$ where $q = t_c$ or $t + 1$ are independent of each other given the current state $(t,c)$. Note that now the conditional coding probabilities in (29) and the conditional transition probabilities in (30) depend solely on $t_c$ and not on $c$.

We will now specify the linear-complexity coding distribution over the source sequences.

Definition 5: For each sequence $x_T^n \in \{0,1\}^T$ we compute for all $x_t^i \in \{0,1\}^i, t = 1,T$, the probabilities $P_c(x_t^i, (t,c))$ recursively,
i.e.,
\[ P_c(x^T_1, (t, q)) \triangleq \sum_q P_c(x^T_1, (t, s)) \cdot P_c((t, q)|(t - 1, s))P_c(x_t|(t, q)) \]
(31)
for \( t > 1 \), and \( P_c(x_1, (1, 1)) = 1/2 \) for both \( x_1 = 0 \) and 1. The summation is only over \( s \) such that \( 1 \leq s \leq t - 1 \).

The coding probabilities
\[ P_c(x^T_1, x^t_1 \in \{0, 1\}^t, t = 1, T \]
are now defined as follows:
\[ P_c(x^T_1) \triangleq \sum Q P_c(x^T_1, (t, q)). \]
(32)

Here we assume that the summation is only over \( q \) such that \( 1 \leq q \leq t \). Of course, \( P_c(\phi) = 1 \).

Just like for the quadratical method, we can demonstrate that also the linear-complexity coding distribution is a weighting of all coding distributions \( P_c(x^T_1|T) \) over all transition patterns \( T \) over \( T \) source symbols, i.e.,
\[ P_c(x^T_1) = \sum Q \sum_{T \rightarrow (T, q)} P_c(T)P_c(x^T_1|T) \]
(33)
where
\[ \sum_{T \rightarrow (T, q)} \]
means a summation over all \( T \) that lead to state \( (T, q) \). The probability \( P_c(x^T_1|T) \) is as defined in (8).

Furthermore, for a transition pattern \( T \) such that \( T \rightarrow (T, q) \) for some \( q \), it follows from inspection that
\[ P_c(T) = \left( \prod_{t = 0}^{C(T) - 1} P_c(t_{c+1} - t_c - 1, 1) \right) P_c(T - t_{C(T)}, 0) \]
(34)
where the \( t_c \) for \( c = 1, C(T) \) are the transition times in \( T \). This probability is composed out of \( C(T) + 1 \) Krichevsky-Trofimov estimators, the first \( C(T) \) of them "end" after a transition occurs, the last one "continues" until the end of the sequence.

The next theorem gives an upper bound on the transition redundancy for the linear method.

**Theorem 2:** Let \( T \geq 2 \). If we use the linear coding distribution defined in (32), the transition redundancy for any \( x^T_1 \) with respect to pattern \( T \) satisfies the upper bound
\[ \log \frac{P_c(x^T_1|T)}{P_c(x^T_1)} \leq \frac{3C(T)}{2} \log \frac{T - 1}{C(T)} + \frac{1}{2} \log (T - 1) + C(T) \log (2e) + 1 \]
(35)
for all transition patterns \( T \) over \( T \) symbols. The bound holds also for \( C(T) = 0 \) if we follow the convention \( \log 0 = 0 \) for \( z = 0 \).

**Proof:** The probability \( P_c(T) \) assigned to pattern \( T \) is given by (34). By the lower bound in (5) and by the basic inequality \( \ln z \leq z - 1 \) we obtain that
\[ \log P_c(u - 1, 1) \geq (u - 1) \log \frac{u - 1}{u} + \log \frac{1}{u} \geq \frac{1}{2} \log u - 1 \]
\[ = -\frac{u - 1}{2} \ln 2 + \frac{1}{2} \log u - 1 \]
\[ \geq -\frac{u - 1}{2} + \frac{1}{2} \log u - 1 \]
\[ = -\log e + \frac{3}{2} \log u - 1 \]
\[ = -\frac{3}{2} \log u - \log (2e) \]
(36)
for integers \( u \geq 1 \). For integers \( v \geq 1 \) the lower bound in (5) yields
\[ \log P_c(v, 0) \geq -\frac{1}{2} \log v - 1 \]
(37)
while for \( v = 0 \) we obtain \( \log P_c(v, 0) = 0 \). Substituting these bounds in \( \log (1/P_c(x_1, (t - 1, s))) \), where \( P_c(T) \) is as in (34), yields two terms. The first term corresponds to the \( C(T) \) transitions. Assuming that \( C(T) > 0 \) and using (36) we get for this first term
\[ \sum_{c = 0, C(T) - 1} \frac{3}{2} \log (t_{c+1} - t_c) + C(T) \log (2e) \]
\[ \leq \frac{3C(T)}{2} \log \frac{1}{C(T)} \sum_{c = 0, C(T) - 1} (t_{c+1} - t_c) \]
\[ + C(T) \log (2e) \]
\[ = \frac{3C(T)}{2} (\log (t_{c+1} - t_c) + C(T) \log (2e) \]
\[ \leq \frac{3C(T)}{2} \log \frac{T - 1}{C(T)} + C(T) \log (2e) \]
(38)
where we used the \cap-convexity of the log in the first inequality and the fact that \( t_{c+1} \geq t_c \), for \( c = 0, C(T) - 1 \). Note also that since \( z \log z = 0 \) for \( z = 0 \), the bound on the first term holds for \( C(T) = 0 \).

For the second term in the bound for \( \log (1/P_c(x_1, (t - 1, s))) \) we get using (37) for \( T > t_{C(T)} \) that
\[ \frac{1}{2} \log (T - t_{C(T)}) + 1 \leq \frac{1}{2} \log (T - 1) + 1 \]
(39)
since \( t_{C(T)} \geq t_0 = 1 \). If \( T = t_{C(T)} \), the second term is zero, which is upper-bounded by \( \frac{1}{2} \log (T - 1) + 1 \).

Combining all this we can conclude that
\[ \log \frac{1}{P_c(T)} \leq \frac{3C(T)}{2} \log \frac{T - 1}{C(T)} + \frac{1}{2} \log (T - 1) + C(T) \log (2e) + 1 \]
(40)
which yields the theorem.

The theorem says that the transition redundancy is now bounded by a bias term which is roughly \( \frac{3}{2} \log T \) bits and then for each transition roughly \( \frac{3}{2} \log T \) additional bits. We see that each transition costs roughly 50% more than for the quadratical method.

**Example (Continued):** The transition redundancy for the linear method for our example source is upper-bounded by
\[ \frac{9}{2} \log 999 + \frac{1}{2} \log 999 + 3 \log (2e) + 1 = 51.02 \text{ bits} \]
This transition redundancy is plotted in Fig. 5. As we can observe the bound 51.02 bit is satisfied. Moreover, it appears that the measured transition redundancy is not as close to the upper bound as for the quadratical method.

**Implementation:** The coding distribution suggested in Definition 5 leads to an implementation in which in all existing nodes (records) \( (t - 1, s) \) the probabilities \( P_c(x_{t-1}^s, (t - 1, s)) \) and the counts \( a(x_{t-1}^s) \) and \( b(x_{t-1}^s) \) are stored. The probabilities \( P_c(x_1^T, (t, q)) \) are now determined using (31) and stored in nodes \( (t, q) \). Summing over all these nodes as in (32) gives \( P_c(x_1^T) \). The counts \( a(x_1^T) \) and \( b(x_1^T) \) in record \( (t, q) \) are determined using the counts \( a(x_{t-1}^s) \) and \( b(x_{t-1}^s) \) in record \( (t - 1, q) \) just as in the quadratical method. After updating, the records \( (t - 1, s) \) can be deleted.
Fig. 5. Redundancy for the linear method as a function of $t$ for $t = 1, 2, \ldots, 1000$.

Storage Complexity: Since for symbol $x_t$, there are $t$ nodes that need to be stored (while the nodes corresponding to $x_{t-1}$ are deleted), the instantaneous storage complexity grows linearly in $T$.

Computational Complexity: To compute the total number of computations (multiplications) to process $x_t^T$, note that there are again two ways to leave the record $(t - 1, s)$. Each of them involves a multiplication. If a transition is assumed, the next record is $(t, t)$, otherwise, the next record is $(t, s)$. The number of computations related to processing $x_t$ for $t > 1$ is, therefore, twice the number of records that are stored for processing $x_{t-1}$, i.e., $2(t - 1)$. The total number of computations for processing $x_t^T$ follows from summing the terms corresponding to $x_t$ for $t = 2, T$, which appears to be $T^2 - T$. Therefore, there is a quadratic dependency between the total computational complexity and the source sequence length $T$ for the linear-complexity coding method.

IV. CONCLUSION

Merhav [3] investigated compression techniques for sources whose parameters can change abruptly at unknown points. He showed that for any uniquely decodable code the redundancy is lower-bounded by $\frac{1}{2} \log T$ bits for each parameter (parameter redundancy) plus $\log T$ bits for each transition (transition redundancy). Moreover, it is outlined by Merhav how to achieve this lower bound with a strongly sequential universal code, i.e., a code that does not need knowledge of $T$.

Actually, Merhav treated the achievability case where there is only one transition. The extension to more transitions is straightforward, he claims. However, if the number of transitions is not known in advance, the Merhav procedure can become quite complex since it involves weighting over the coding distributions $P_e(x_t^T | T)$ corresponding to all transition patterns $T$ of sequence $x_t^T$. Note that there are $2T - 1$ such patterns, hence this would lead to a complexity which is exponential in $T$.

In contrast to Merhav, we concentrated on finding explicit coding methods for the case where the number of transitions is unrestricted and that do not have exponential complexity behavior. We suggested two coding methods, the quadratical-complexity method and the linear-complexity method, both based on the concept of weighting (see, e.g., Ryabko [7], [6] but more recently also [10]) and Weinberger et al. [9]). Both methods are analyzed and it was shown that the quadratical method achieves the Merhav lower bound in the sense that each additional transition gives a redundancy increase of roughly $\log T$ bits. The linear method has a slightly larger transition redundancy ($\frac{1}{2} \log T$ bits per additional transition) but also a more acceptable complexity.

Both methods have a bias term contributing to the transition redundancy which is roughly $\frac{1}{2} \log T$ bits. This bias term is the result of not knowing the number of transitions in advance. From simulations we could see that in practice the redundancies of both methods are very close.

Just like Merhav, we would like to point out that this source coding situation should be treated as a Minimum Description Length (MDL) problem (see Rissanen [4], [5]). What the transmitter has to convey to the receiver is a specification (code) of the transition pattern (the model in terms of [10]) plus the code for the source sequence given the pattern. The last code contains a term coming from the fact that we do not know the parameters (parameter redundancy). The pattern is chosen in such a way that the cost of specifying the pattern plus the code length of the source sequence given the pattern, is minimized (see also [12]). We are not making a definite choice here, instead we are weighting over all transition patterns such that only the best pattern survives.

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REFERENCES

Fault-Tolerant Cube Graphs and Coding Theory

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Abstract—Hypercubes, meshes, tori, and Omega networks are well-known interconnection networks for parallel computers. The structure of those graphs can be described in a more general framework called cube graphs. The idea is to assume that every node in a graph with $q$ nodes is represented by a unique string of $q$ symbols over $GF(q)$. The edges are specified by a set of offsets, those are vectors of length $q$ over $GF(q)$, where the two endpoints of an edge are an offset apart. We study techniques for tolerating edge faults in cube graphs that are based on adding redundant edges. The redundant graph has the property that the structure of the original graph can be maintained in the presence of edge faults. Our main contribution is a technique for adding the redundant edges that utilizes constructions of error-correcting codes and generalizes existing ad hoc techniques.

Index Terms—Fault tolerance, parallel computing, interconnection networks, hypercubes, omega networks, error-correcting codes.

I. INTRODUCTION

The mesh, torus, hypercube, and Omega networks are some of the most important interconnection networks for parallel computers. One of the most important issues in the design of a system which contains many components is the system’s performance in the presence of faults. Hence, it is of major practical importance to develop efficient techniques (in terms of the cost of the redundancy) to handle faults in those architectures.

Our approach for handling faults is based on a graph model. In this model the architecture is viewed as a graph, where the nodes represent the processors and the edges represent communication links between the nodes. A target graph is selected and the required amount of fault tolerance $f$ is determined. Then a fault-tolerant graph, that has the same number of nodes as the target graph, is defined with the property that given any set of $f$ or fewer faulty edges, the remaining graph (after removal of the faulty edges) is guaranteed to contain the target graph as a subgraph. Note that this approach guarantees that any algorithm designed for the target graph will run with no slowdown in the presence of $f$ or fewer edge faults in the fault-tolerant graph, regardless of their distribution. Minimizing the cost in this model amounts to constructing a fault-tolerant graph with minimum degree. (See [4] for constructions that address node faults.)

The key to our fault-tolerant constructions is a technique based on error-correcting codes for adding redundant edges using so-called wildcard dimensions. Our technique is capable of handling an arbitrary number of faults over arbitrary alphabets. This is a generalization of the construction in [5] that is limited to the case of MDS (Maximum Distance Separable) codes. The key to our generalization is proving the equivalence between the constructions of fault-tolerant cube graphs and constructions of generator matrices of error-correcting codes (while in [5] the construction is based on parity check matrices of error-correcting codes.) First, we will describe the technique for constructing fault-tolerant hypercubes, then we will explain how it can be extended to Omega networks, tori, meshes, and cube-connected cycles (CCC).

Let $Q_i$ denote the $i$-dimensional hypercube. It consists of $2^i$ nodes, where node $i, 0 \leq i \leq 2^i - 1$, is represented by the $i$-bit binary representation of $i$. Two nodes, say $X = (x_{i-1}x_{i-2}...x_0)$ and $Y = (y_{i-1}y_{i-2}...y_0)$, are connected by an edge iff there is a single $j$ for which $x_j \neq y_j$, and this edge is called a dimension-$j$ edge. Hence, the set of edges in the hypercube can be partitioned into $i$ dimensions.

In [9], it was suggested to add another set of edges to the hypercube, called the wildcard dimension, resulting in a folded hypercube. The folded hypercube topology was also defined independently in [7], but with a different name (the bisectional interconnection network) and with a different addressing scheme for the nodes. A formal definition of the folded hypercube is given next.

Definition 1: An $i$-dimensional folded hypercube, denoted by $F_i$, is an $i$-dimensional hypercube to which extra links are added connecting every pair of nodes that are bit-wise complements of each other.

A wildcard edge in $F_i$ is an edge that connects node $X = (x_{i-1}x_{i-2}...x_0)$ and node $\overline{X} = (\overline{x}_{i-1}\overline{x}_{i-2}...\overline{x}_0)$. For example, $F_3$ is depicted in Fig. 1; notice that the neighbors of node (000) are (001), (010), (100) and (111). The structure of $Q_i$ and $F_i$ can be described using a more general framework, see also [8].

Definition 2: Let $\ell$ be a positive integer and let $q$ be a prime number. Let $S \subseteq \{0,1,...,q-1\}^\ell$. The graph $N(q,\ell,S)$ is a graph with $q^\ell$ nodes and with the edges specified by the set of vectors in $S$. Any two nodes, say $X$ and $Y$, are connected by an edge iff there exists a vector $V \in S$ such that $X + V = Y$ where addition is vector addition performed over $GF(q)$.