

# An Efficient Joint Source-Channel Decoder with Dynamical Block Priors

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An efficient joint source-channel (s/c) decoder based on the side information of the source and on the MN-Gallager algorithm over Galois fields is presented. The dynamical block priors (DBP) are derived either from a statistical mechanical approach via calculation of the entropy for the correlated sequences, or from the Markovian transition matrix. The Markovian joint s/c decoder has many advantages over the statistical mechanical approach. In particular, there is no need for the construction and the diagonalization of a  $q \times q$  matrix and for a solution to saddle point equations in  $q$  dimensions. Using parametric estimation, an efficient joint s/c decoder with the lack of side information is discussed. Besides the variant joint s/c decoders presented, we also show that the available sets of autocorrelations consist of a convex volume, and its structure can be found using the Simplex algorithm.

## §1. Introduction

Source coding is a process for removing redundant information from the source information symbol stream. Channel coding is a procedure for adding redundancy as protection into the information stream which is to be transmitted. During the last decade engineers and also physicists have designed efficient error correction techniques such as Low-Density-Parity-Check-Codes (LDPC)<sup>1),2)</sup> or Turbo codes, that nearly saturate Shannon's limit.

The Shannon separation theorem states that source coding (compression) and channel coding (error protection) can be performed separately and sequentially, while maintaining optimality <sup>1),3)</sup>. However, this is true only in the case of asymptotically long block lengths of data and point-to-point transmission. In many practical applications, the conditions of the Shannon's separation theorem neither hold, even approximately. Thus, considerable interest has developed in various schemes of joint source-channel (s/c) coding, where compression and error correction are combined into one mechanism.

The paper is organized as follows. In Section 2, Statistical Mechanical (SM) methods are used to explore properties of correlated sequences. In section 3, the space of possible sets of autocorrelations is investigated. In sections 4 Mackay and Neal's (MN) algorithm for error correction is briefly introduced, and the results of section 2 are used to extend this algorithm to a joint s/c scheme. The estimation of the threshold of the code using the scaling argument for the convergence time is presented in section 5. Section 6 presents simulation results, and compares joint to separation schemes. In Sections 7 and 8, the joint s/c problem is attacked from a different point of view, a Markovian approach, and an efficient joint scheme *with the lack of side information* is demonstrated.

## §2. Joint s/c coding - Statistical Mechanical approach

In our recent papers<sup>4),5)</sup> a particular scheme based on a SM approach for the implementation of the joint s/c coding was presented and the main steps are briefly summarized below. The original boolean source is first mapped to a binary source  $\{x_i = \pm 1\}$   $i = 1, \dots, L$ , and is characterized by a finite set of autocorrelations bounded by the length  $k_0$

$$C_{k_1, \dots, k_m} = \frac{1}{L} \sum_{i=1}^L x_i \prod_{j=0}^m x_{(i+k_j) \bmod L} \quad (2.1)$$

where  $k_m \leq k_0$  is the highest length autocorrelation taken and the total number of possible different autocorrelations is  $2^{k_0}$ . For  $k_0 = 2$ , for instance, there are only 4 possible correlations,  $C_0, C_1, C_2$  and  $C_{12}$ , and for  $k_0 = 3$  there are 8 possible different correlations;  $C_0, C_1, C_2, C_3, C_{12}, C_{13}, C_{23}, C_{123}$ , where we do not assume left-right symmetry for the source. Note that for general  $k_0$  and  $m = 1$ , Eq. (2.1) reduces to the two-point autocorrelation function <sup>7)</sup>. The number of sequences obeying these  $2^{k_0}$  constraints is given by

$$\Omega = Tr_{\{x_i = \pm 1\}} \prod_{\{k_1, k_2, \dots, k_m\}} \delta\left(\sum_{i=1}^L x_i \prod_{j=0}^m x_{i+k_j} - LC_{k_1, \dots, k_m}\right) \quad (2.2)$$

where  $m = 0$  stands for  $C_0$ . Using the integral representation of the delta functions, Eq. (2.2) can be written as

$$\Omega = \int \prod_{\{k_1, \dots, k_m\}} dy_{\{k_1, \dots, k_m\}} \exp\left(\sum_{k_1, \dots, k_m} -y_{k_1, \dots, k_m} C_{k_1, \dots, k_m}\right) Tr \exp\left(\sum_{k_1, \dots, k_m} y_{k_1, \dots, k_m} \sum_i x_i \prod_{j=0}^m x_{i+k_j}\right)$$

Since  $k_j \leq k_0$ , the last term of this equation indicates that the trace can be performed using the standard transfer matrix (of size  $2^{k_0} \times 2^{k_0}$ ) method<sup>8)</sup>. More precisely, assume two successive blocks of  $k_0$  binary variables denoted by  $(x_1, \dots, x_{k_0})$  and  $(x_{k_0+1}, \dots, x_{2k_0})$ . The element  $(i, j)$  of the transfer matrix is equal to the value of the last exponential term (on the r.h.s of the trace), where the first block is in state  $i$  (among  $2^{k_0}$  possible states) and the second block is in state  $j$ . The transfer matrix is a non-negative matrix (as long as the  $y_{k_1, \dots, k_m}$  are real numbers), and the leading eigenvalue is positive and non-degenerate<sup>8)</sup>. In the leading order one finds

$$\Omega = \int dy_k \exp\left\{-L\left[\sum_{k_1, \dots, k_m} y_{k_1, \dots, k_m} C_{k_1, \dots, k_m} - \ln \lambda_{max}(\{y_{k_1, \dots, k_m}\})\right]\right\} \quad (2.3)$$

where  $\lambda_{max}$  is the maximal eigenvalue of the corresponding transfer matrix. For large  $L$  and using the saddle point method, the entropy,  $H_2(\{C_{k_1, \dots, k_m}\})$ , is given in the leading order by

$$H_2(\{C_{k_1, \dots, k_m}\}) = \frac{1}{\ln 2} \left[ \frac{1}{k_0} \ln \lambda_{max}(\{y_{k_1, \dots, k_m}\}) - \sum_{k_1, \dots, k_m}^{k_0} y_{k_1, \dots, k_m} C_{k_1, \dots, k_m} \right] \quad (2.4)$$

where  $\{y_{k_1, \dots, k_m}\}$  are determined from the saddle point equations of  $\Omega$  (4), (5). Assuming a Binary Symmetric Channel (BSC) and using Shannon's lower bound, the channel capacity of sequences with a given set of autocorrelations bounded by a distance  $k_0$  is given by

$$C = \frac{1 - H_2(f)}{H_2(\{C_{k_1, \dots, k_m}\}) - H_2(P_b)} \quad (2.5)$$

where  $f$  is the channel bit error rate and  $p_b$  is a bit error rate. The saddle point solutions derived from Eq. (2.4) indicate that the equilibrium properties of the one-dimensional Ising spin system ( $x_i = \pm 1$ ) with up to order  $k_0$  multi-spin interactions

$$H = - \sum_i \sum_{k=1}^{k_0} \frac{y_{k_1, \dots, k_m}}{\beta} x_i \prod_{j=0}^m x_{i+k_j} \quad (2.6)$$

obey in the leading order the autocorrelation constraints, Eq. (2.1). This property of the effective Hamiltonian, Eq. (2.6), is used in simulations to generate an ensemble of signals (source messages) with the desired set of autocorrelations. *Note that in the following we choose  $\beta = 1$ , and hence we denote  $\{y_{k_1, \dots, k_m}\}$  as interactions.*

### §3. Possible sets of autocorrelations and the Simplex algorithm

The entropy of correlated sequences can be calculated from Eq. (2.4). For the simplest case of sequences obeying only  $C_1$  and  $C_2$  the numerical solution of the saddle point equations indicate that the entropy is positive only in the regime

$$-(1 + C_2)/2 \leq C_1 \leq (1 + C_2)/2 \quad (3.1)$$

where outside of this regime the entropy is zero. At the boundaries,  $C_1 = |(1+C_2)/2|$ , two phenomena are observed: (a) the entropy falls abruptly to zero at the boundary, and (b)  $y_1$  and  $-y_2$  diverge at the boundary (the one-dimensional Hamiltonian, Eq. (2.6) consists of frustrated loops).

These limited results obtained from the numerical solutions of the saddle point equations suffer from the following limitations: (a) finding the boundaries of the region in the space of  $\{C_{k_1, \dots, k_m}\}$  with a finite entropy is very sensitive to the numerical precision since on the boundary the  $\{|y_i|\}$  diverge; (b) it is unclear whether the available space consists of a connected regime; (c) the question of whether out of the space with a finite entropy, there are a finite or infinite number of sequences (for instance  $e^{\sqrt{L}}$ ) obeying the set of autocorrelations cannot be answered using the saddle point method; (d) extension of the saddle point solutions to identify the boundaries of the finite entropy regime to many dimensions is a very heavy numerical task.

To overcome these difficulties, we show below how the possible sets of autocorrelations can be identified using the Simplex algorithm.

For the case of only two constraints  $C_1$  and  $C_2$ , for instance, following the methodology of the transfer matrix, let us concentrate on four successive binary variables  $S_i, S_{i+1}, S_{i+2}, S_{i+3}$ , where  $S_i = \pm 1$ . Since the Hamiltonian, Eq. (2.6),

obeys in this case an inversion symmetry, let us examine only the 8 configurations out of 16 where  $S_3 = -$ ,  $(\pm \pm -\pm)$ . For these 8 configurations one can assign the following marginal probabilities,  $P_{\pm\pm-\pm}$ , where each probability stands for the fraction of sequences obeying  $C_1$  and  $C_2$  with a given state for these four successive binary variables. In the SM language we measure the probabilities of these four states in thermal equilibrium of the micro-canonical ensemble obeying Eq. (2.1). It is clear that the Hamiltonian, Eq. (2.6), is translationally invariant,  $P(S_i, S_{i+1}, S_{i+2}, S_{i+3})$  is independent on  $i$  after averaging over all sequences obeying constraints (2.1).

For these 8 marginal probabilities one can write the following 14 equations:

$$\begin{aligned}
P_{----+} + P_{----} + P_{+--+} + P_{+---} - P_{-+++} - P_{-+++} - P_{+----} - P_{+----} &= C_1/2 \\
P_{----+} + P_{----} + P_{+---} + P_{+---} - P_{-+++} - P_{-+++} - P_{+----} - P_{+----} &= C_1/2 \\
P_{-+---} + P_{----} + P_{+---} + P_{+---} - P_{-+++} - P_{-+++} - P_{+----} - P_{+----} &= C_1/2 \\
P_{----+} + P_{----} + P_{-+---} + P_{-+---} - P_{-+++} - P_{-+++} - P_{+----} - P_{+----} &= C_2/2 \\
P_{----} + P_{-+++} + P_{+---} + P_{+---} - P_{-+++} - P_{-+++} - P_{+----} - P_{+----} &= C_2/2 \\
P_{----} + P_{-+++} + P_{-+---} + P_{-+---} + P_{+----} + P_{+----} + P_{+---} + P_{+---} &= 1/2 \\
0 \leq P_{\pm\pm-\pm} \leq 1 & \tag{3.2}
\end{aligned}$$

For a given  $C_1$ , these 14 equations can be solved for the minimum and the maximum available  $C_2$  using the Simplex method. Running over values of  $-1 \leq C_1 \leq 1$ , we indeed recover the result of Eq. (3.1). However, the *Simplex solution indicates the lack of even a finite number of sequences beyond the regime with finite entropy*. Hence, a simple geometrical calculation obeying constraint 3.1 indicates that the fraction of the 2D space  $(-1:1, -1:1)$  of  $(C_1, C_2)$  with available sequences is  $1/2$ .

For the case of three constraints,  $C_1, C_2$  and  $C_3$ , one can similarly write 45 equalities and inequalities for the 32 probabilities of 6 successive binary variables  $P_{\pm\pm\pm\pm-\pm}$ . For a given  $C_1$  and  $C_2$ , these 45 equations and inequalities can be solved for the minimum and the maximum available  $C_3$  using the Simplex method. The Simplex solution indicates: (a) the solution space in the three-dimensional box  $(-1 : 1, -1 : 1, -1 : 1)$  for  $(C_1, C_2, C_3)$  is a connected region bounded by a few planes. This result is consistent with the solution obtained from Eqs. (2.3,2.4); (b) the fraction of the volume of the box obeying the three constants is  $\sim 0.222$ . Preliminary results indicate that for 4 ( $C_i, i = 1, 2, 3, 4$ ) and 5 ( $C_i, i = 1, 2, 3, 4, 5$ ) constraints the available volume is  $\sim 0.085, 0.034$ , respectively.

The fraction of possible sets of autocorrelations appears to decrease as the number of constraints increases. However, the question of whether the fraction of available autocorrelations drops exponentially with the number of constraints as well as its detailed spatial shape is the subject of our current research.

We conclude the discussion in this section with the following general result. The available volume for the general case of  $q$  constraints  $\{C_{k_1, \dots, k_m}\}$ ,  $k_m < \log_2(q)$ , is convex. The main idea is that one can verify that the set of equalities can be written in a matrix representation in the following form

$$\mathbf{MP} = \mathbf{C} \tag{3.3}$$

where  $\mathbf{M}$  is a matrix with elements  $\pm 1$ ;  $P$  represents the marginal probabilities  $P(\pm, \pm, \dots)$  and  $C$  represents the desired correlations or a normalization constant (for instance  $C_1/2$ ,  $C_2/2$  and  $1/2$ , for the case of Eq. (3·2)). The inequalities force the probabilities into the range  $[0 : 1]$ . Clearly if  $P_1(\pm, \pm, \dots)$  and  $P_2(\pm, \pm, \dots)$  are two sets of probabilities obeying Eq. (3·3) then

$$\lambda P_1 + (1 - \lambda) P_2 \tag{3·4}$$

is also a solution of the set of the equalities ( $0 \leq \lambda \leq 1$ ). Hence, the available volume is convex.

#### §4. Joint s/c decoder: Statistical Mechanical approach

The transfer matrix method indicates that the relevant scale of the correlated source message is  $k_0$ . Hence, our encoding/decoding procedure is based on the MN code for a finite field  $q = 2^{k_0} - 10$ , which is based on the construction of two sparse matrices  $A$  and  $B$  of dimensionalities  $(L_0/R) \times L_0$  and  $(L_0/R) \times (L_0/R)$  respectively, where  $R$  is the code-rate and the number of symbols in the source is  $L_0 = L/k_0$ . The matrix  $B^{-1}A$  is then used for encoding the message

$$t = B^{-1}Ax \pmod{q} \tag{4·1}$$

The finite field message vector  $t$  is mapped to a binary vector and then transmitted. The received message,  $r$ , is corrupted by the channel bit error rate,  $f$ .

The decoding of symbols of  $k_0$  successive bits (named in the following as a *block* of bits or binary variables) is based on the solution of the syndrome

$$Z = Br = Ax + Bn \pmod{q} \tag{4·2}$$

where  $n$  stands for the corresponding noise of  $k_0$  successive bits. The solution of the  $L_0/R$  equations with  $L_0(1/R + 1)$  variables is based on the standard message passing algorithm introduced for the MN decoder over Galois fields with  $q = 2^{k_0} - 10$  and with the following modification. The horizontal pass is left unchanged, *but a dynamical set of probabilities assigned for each block is used in the vertical pass*. The Dynamical Block Probabilities (DBP),  $\{P_n^c\}$ , are determined following the current belief regarding the neighboring blocks and are given by

$$\gamma_n^c = S_I(c) \left( \sum_{l=1}^q q_L^l S_L(l, c) \right) \left( \sum_{r=1}^q q_R^r S_R(c, r) \right) \quad ; \quad P_n^c = \frac{\gamma_n^c}{\sum_{j=1}^q \gamma_n^j} \tag{4·3}$$

where  $l/r/c$  denotes the state of the left/right/center ( $n-1 / n+1 / n$ ) block respectively and  $q_L^l/q_R^r$  are their posterior probabilities.  $S_I(c) = e^{-\beta H_I}$  stands for the Gibbs factor of the inner energy of a block ( $k_0$  successive binary variables spins), characterized by an energy  $H_I$  at a state  $c$ , see Eq. (2·6). Similarly  $S_L(l, c)$  ( $S_R(c, r)$ ) stands for the Gibbs factor of consecutive Left/Center (Center/Right) blocks at a state  $l, c$  ( $c, r$ ) (4), (5). The complexity of the calculation of the block prior probabilities is  $O(Lq^2/\log q)$  where  $L/\log q$  is the number of blocks. The decoder complexity per

iteration of the MN codes over a finite field  $q$  can be reduced to order  $O(Lqu)$ , where  $u$  stands for the average number of checks per block. Hence the total complexity of the DBP decoder is of the order of  $O(Lqu + Lq^2/\log q)$ .

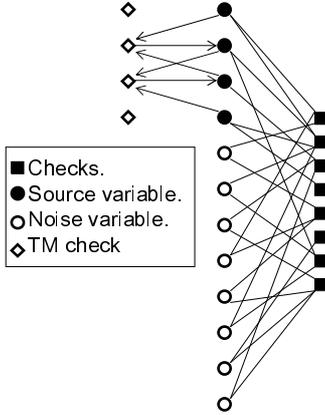


Fig. 1. A message passing in the joint s/c decoder is represented by a bipartite graph with an additional layer.

Another way to represent the dynamical behavior of the SM joint s/c decoder is in the framework of message passing on a graph (Fig. 1). Typically for LDPC, the graph is bipartite and consists of variable nodes (circles) and check nodes (squares). For the MN algorithm, there are two types of variable nodes: source nodes (filled circles) and noise nodes (open circles). A message from variables to checks is a horizontal pass, and a message from checks to variables is a vertical pass. In the joint s/c decoder there is a *third* layer (diamonds): each element in this layer sends a message (outbound arrow) to a single source variable (namely, the dynamical block prior), and receives two messages (inbound arrows) from the neighboring source variables (namely their a-posteriori probabilities).

For simplification of the discussion below, in almost all of the simulation results we concentrate on rate  $1/3$  and the construction of the matrices  $A$  and  $B$  follow reference 12) which is sketched in Fig. 2. The advantage of this construction is that the matrices  $A$  and  $B$  are very sparse, but the threshold of the code for large blocks is only  $1 - 3\%$  of the channel capacity (12), (13). Furthermore, since  $B$  has a systematic structure, the complexity of the encoder scales linearly with  $L$  although  $B^{-1}$  is dense (14), (15). Of course, codes with higher thresholds exist, hence the performance of the joint s/c algorithm reported below should be interpreted as a lower bound. (Results for a limited example with rate greater than one,  $R > 1$ , are briefly discussed in reference 16))

We conclude this section with the comment that the possibility of the SM joint s/c algorithm in the framework of the MN-Gallager decoder to the Gallager decoder (2) is in question. In the Gallager decoder we first solve  $L_0(1/R - 1)$  equations for the noise variables, and only in the final step is the message recovered. Since the

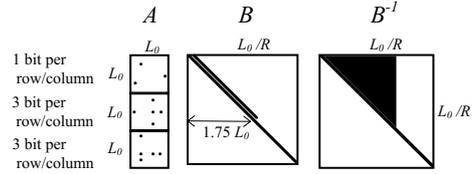


Fig. 2. The structure of the matrices  $A$  and  $B$  for the MN decoder taken from reference 12), for rate  $1/3$ . The black dots (area) denote the non-zero elements of the matrices  $A$ ,  $B$ ,  $B^{-1}$ .

noise is not spatially correlated, we do not see a simple way to incorporate in the Gallager case the side information about the spatial correlations among the message variables. The equivalence between these two (MN-Gallager and Gallager) similar decoders therefor also in doubt.

For illustration, in Fig. 3 we present results for rate  $R = 1/3$ ,  $L = 10,000$ ,  $q = 4$  and 8 where the decoding is based on the dynamical block posterior probabilities, Eq. (4.3), and with the following parameters. For  $q = 4$  (open circles)  $C_1 = 0.55$ ,  $C_2 = 0.5$ ,  $C_{12} = 0.4$  ( $y_1 = 0.275$ ,  $y_2 = 0.291$ ,  $y_{12} = 0.149$ ) and  $H_2 = 0.683$ . Shannon's lower bound, Eq. (2.5), is denoted by the double dotted line, where for  $p_b = 0$  the channel noise level is  $f_c = 0.227$ . For  $q = 8$  (open diamonds)  $C_1 = 0.77$ ,  $C_2 = 0.69$ ,  $C_3 = 0.56$ ,  $C_{123} = 0.7$  ( $y_1 = 0.349$ ,  $y_2 = 0.36$ ,  $y_3 = -0.211$ ,  $y_{123} = 0.443$ ) and  $H_2 = 0.453$ . Shannon's lower bound is denoted by the dashed line, where for  $p_b = 0$  the channel noise level is  $f_c = 0.275$ . Each point was averaged over at least 1,000 messages. These results for both  $q = 4$  and 8 indicate that the threshold of the presented decoder with  $L = 10,000$  is  $\sim 15\% - 20\%$  below the channel capacity for infinite source messages. It is worth mentioning, that without using the dynamical block priors, Eq. (4.3), the decoder fails to decode at  $f_c \approx 0.13 - 0.14$ .

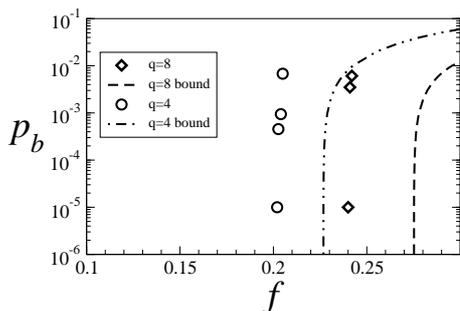


Fig. 3. Simulation results for rate  $R = 1/3$ ,  $L = 10,000$ ,  $q = 4$  and 8. Each point was averaged over at least 1,000 source messages with the desired set of autocorrelations. (refer to text for description)

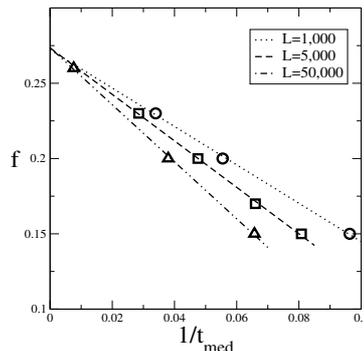


Fig. 4. The flip rate  $f$  as a function of  $1/t_{med}$  for  $GF(4)$  with  $C_1 = C_2 = 0.8$  and  $L = 1,000, 5,000, 50,000$ . The lines are a result of a linear fit. The threshold,  $f_\infty \sim 0.272$ , extrapolated from the scaling behavior Eq. (5.1), is independent of  $L$ .

## §5. The threshold of the code

The threshold  $f_c$  for  $L \rightarrow \infty$  is estimated from the scaling argument of the convergence time, which was previously observed for  $q = (212, 13)$ . The median convergence time, measured in iterations of the MN algorithm, is assumed to diverge as the level of noise approaches  $f_c$  from below. More precisely, we found that the scaling for the divergence of  $t_{med}$  is *independent of  $q$*  and is consistent with:

$$t_{med} = \frac{A}{f_c - f} \quad (5.1)$$

where for a given set of autocorrelations and  $q$ ,  $A$  is a constant. Moreover, for a given set of autocorrelations and a finite field  $q$ , the extrapolated threshold  $f_c$  is independent of  $L$ , as demonstrated in Fig. 4. This observation is essential to determine the threshold of a code based on the above scaling behavior. Note that the estimation of  $t_{med}$  is a simple computational task in comparison with the estimation of low bit error probabilities for large  $L$ , especially close to the threshold. We also note that the analysis is based on  $t_{med}$  instead of the *average* number of iterations,  $t_{av}$ ,<sup>12)</sup> since we wish to prevent the dramatic effect of a small fraction of finite samples with slow convergence or no convergence.<sup>17)</sup> We note that preliminary results indicates that for a given  $\{C_k\}$ ,  $f_c(q)$  appears asymptotically to be consistent with  $f_c(q) \sim f_c - const/q$ .

### §6. Comparison between joint and separation schemes

Results of simulations for  $q = 4, 8, 16$  and  $32$  and selected sets of autocorrelations are summarized in Table I (Fig. 5) and the definition of the symbols is:  $\{C_k\}$  denotes the imposed values of two-point autocorrelations as defined in eqs. 2·1 and 2·2;  $\{y_k\}$  are the interactions strengths, Eq. (2·6);  $H$  represents the entropy of sequences with the given set of autocorrelations, Eq. (2·4);  $f_c$  is the estimated threshold of the MN decoder with the DBP derived from the scaling behavior of  $t_{med}$ , Eq. (5·1);  $f_{Sh}$  is Shannon's lower bound, Eq. (2·5); Ratio is the efficiency of our code,  $f_c/f_{Sh}$ ;  $Z_R$  indicates the gzip compression rate averaged over files of the sizes  $10^5 - 10^6$  bits with the desired set of autocorrelations. We assume that the compression rate with  $L = 10^6$  achieves its asymptotic ratio, as was indeed confirmed in the compression of files with different  $L$ ;  $1/R^*$  indicates the ideal (minimal) ratio between the transmitted message and the source signal after implementing the following two steps: compression of the file using gzip and then using an *ideal optimal encoder/decoder*, for a given BSC with  $f_c$ . A number greater than (less than) 3 in this column indicates that the MN joint s/c decoder is more efficient (less efficient) in comparison to the channel separation method using the standard gzip compression. The last four columns of Table I (Fig. 5) are devoted to the comparison of the presented joint s/c decoder with advanced compression methods.  $PPM_R$  and  $AC_R$  represent the compression rate of files of the size  $10^5 - 10^6$  bits with the desired autocorrelations using the Prediction by Partial Match<sup>18)</sup> and for the Arithmetic Coder<sup>19)</sup>, respectively. Similarly to the gzip case,  $1/R_{PPM}$  and  $1/R_{AC}$  denote the optimal (minimal) rate required for the separation process (first a compression and then an ideal optimal encoder/decoder) assuming a BSC with  $f_c$ .

Table I indicates the following main results: (a) a degradation in the performance is observed as the correlations are enhanced. (b) The efficiency of our joint s/c coding technique is superior to the alternative standard gzip compression in the separation technique. The gain disappears as the entropy decreases. These results are farther discussed in 5) .

The DBP decoder based on the SM approach suffers one major disadvantage: computing the  $y_k$  interactions, which involves finding the eigenvalues of a  $q \times q$  matrix ( $O(q^3)$ ), and solving the saddle-point equations, 2·4, which is also a heavy numerical

$q$	$c_1$	$c_2$	$c_3$	$c_4$	$c_5$	$y_1$	$y_2$	$y_3$	$y_4$	$y_5$	$H$	$f_c$	$f_{sh}$	Ratio	$Z_R$	$\frac{1}{R^*}$	$PPM_R$	$\frac{1}{R_{PPM}}$	$AC_R$	$\frac{1}{R_{AC}}$
4	0.65	0.65	-	-	-	0.29	0.53	-	-	-	0.58	0.239	0.247	0.97	0.69	3.32	0.64	3.1	0.65	3.12
4	0.72	0.72	-	-	-	0.3	0.61	-	-	-	0.49	0.253	0.266	0.95	0.61	3.3	0.55	3.001	0.58	3.13
4	0.8	0.8	-	-	-	0.32	0.71	-	-	-	0.38	0.273	0.294	0.93	0.5	3.23	0.43	2.76	0.48	3.1
4	0.86	0.86	-	-	-	0.37	0.81	-	-	-	0.29	0.293	0.321	0.91	0.41	3.16	0.32	2.52	0.38	2.9
8	-0.65	0.6	-0.55	-	-	-0.31	0.23	-0.23	-	-	0.59	0.236	0.244	0.97	0.71	3.34	0.66	3.14	0.67	3.16
16	0.6	0.6	0.58	0.6	-	0.10	0.17	0.164	0.33	-	0.57	0.229	0.249	0.92	0.7	3.12	0.66	2.95	0.67	2.98
32	0.62	0.7	0.55	0.55	0.6	0.32	0.82	-0.44	-0.22	0.51	0.49	0.242	0.266	0.91	0.62	3.06	0.56	2.77	0.6	2.96

Fig. 5. Results for  $q = 4, 8, 16, 32$  and selected sets of two-point autocorrelations  $\{C_k\}$ 

task, even for  $k_0 > 4$ .

### §7. Markovian joint s/c decoder

In order to overcome the abovementioned drawbacks of the SM approach, we now treat the source sequence from a different point of view, by assuming that these sequences were generated by a Markov process. Hence, the sequence can be described by an alphabet  $GF(q)$ , a transition matrix  $T_{kj} = P(x_i = j | x_{i-1} = k)$ , and the stationary solutions of the process,  $P(j)$ . The key point is that for large messages,  $T$  and  $P(j)$  can be (approximately) measured by the sender (for each chunk of data) with  $O(L_0)$  operations.  $T$  and  $P(j)$  may then be transmitted reliably to the decoder and used as side information in the decoding process (similar to the transmission of the  $y_1, \dots, y_{k_0}$  interactions in the SM approach).

Consider three successive symbols  $x_{i-1}, x_i, x_{i+1}$  in such a sequence. The probability of the triplet  $a, b, c$  is given by:

$$P(a, b, c) = P(a, b) \cdot P(c|a, b) = P(a, b) \cdot P(c|b) = \frac{P(a, b)P(b, c)}{P(b)} \quad (7.1)$$

where use has been made of the Bayes Rule:  $P(x, y) = P(x) \cdot P(y|x)$ , and fact that the process is memoryless. Now, given the a-posteriori probabilities for the first and last symbols in the triplet:  $q_{i-1}^a = Pr(x_{i-1} = a)$  and  $q_{i+1}^c = Pr(x_{i+1} = c)$ , one can calculate a dynamical block prior (corresponding to the prior in (4.3) for the probability that  $x_i = b$ :

$$\begin{aligned} Pr(x_i = b) &= \frac{1}{Z} \cdot \sum_{a,c=1}^q P(a, b, c) \cdot q(a) \cdot q(c) = \\ &= \frac{1}{Z} P(b)^{-1} \left( \sum_{a=1}^q P(a, b) q_{i-1}^a \right) \cdot \left( \sum_{c=1}^q P(b, c) q_{i+1}^c \right), \end{aligned} \quad (7.2)$$

where  $Z$  is a normalization constant such that:  $\sum_{b=1}^q Pr(x_i = b) = 1$ .

The extension of the MN algorithm to the joint source-channel case consists of the following steps:

1. A binary sequence of  $L_0 \cdot \log_2(q)$  bits is converted to  $L_0 GF(q)$  symbols.
2. The encoder measures  $T$  and  $P(j)$  for all the  $q$  symbols over the source, and transmits reliably this side information to the decoder.
3. The source is encoded according to (4.1), then reconverted to binary representation and transmitted.
4. The decoder maps the received signal back to  $GF(q)$ , and performs the regular decoding (4.2), but after every iteration of message passing, the prior for each source symbol is recalculated according to (7.2).

The complexity of calculating the  $q$  priors for a single symbol according to the posteriors of its neighbors is reduced from  $q^3$  to  $q^2$  by Eq. (7.2), hence the decoder's complexity remains linear, with total complexity of  $O(L_0 q u + L_0 q^2)$  per iteration.

However, there is still a need for the transmission of the side information ( $T$ ,  $P(j)$ ). Hence the size of the header is of the order of  $O(q^2)$ . For  $L \rightarrow \infty$  or more precisely for  $L \gg q^2$  the overhead of the transmitted side information is negligible; however, for a finite  $L \leq q^2$  it may cancel the benefits of the Markovian joint s/c decoder.

The header size can be reduced using the following observation: For sequences with enhanced autocorrelations, the structure of  $T$  was observed to be dominated by a small number of large elements. One can transmit only these dominated elements, and the remaining elements of each row of  $T$  will be filled evenly to maintain:  $\sum_{j=1}^q T_{ij} = 1$ .

Fig. 6 represents simulation results for the Markovian scheme. Triangles mark decoding using full  $T$ , while circles mark decoding using only the 8 (out of 64) most dominant elements of  $T$ . The performance seems to be only slightly affected by this approximation, which dramatically reduces the required transmitted side information.

An interesting open question is the effect of use of the sparseness of  $T$  on the complexity and performance of the decoder.

### §8. Efficient Joint s/c decoder with the lack of side information

The discussion in the previous sections indicates that the performance of the presented joint s/c coding is not too far from Shannon's lower bound and, most probably, using an optimized code (a better construction for the matrices  $A$  and  $B$

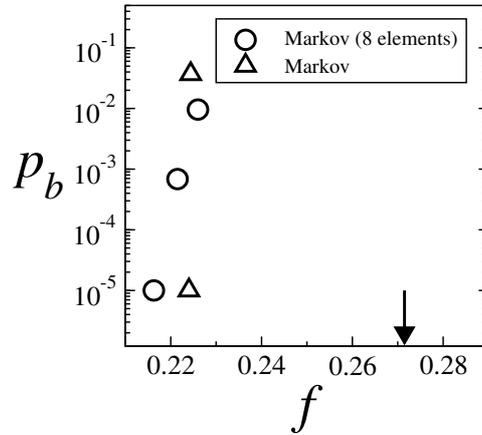


Fig. 6. The bit error rate,  $p_b$  versus the channel bit error rate  $f$  for  $L = 10,000$ ,  $R = 1/3$ ,  $q = 8$  with  $C_1 = C_2 = C_3 = 0.7$ . Decoding following the Markovian process, Eq. (7.2) (open triangle), decoding following the Markovian process where only 8 dominated elements of the transition matrix,  $T$ , are taken as a side information (open circle). Shannon's lower bound,  $f_c = 0.271$  ( $H = 0.47$ ), is denoted by an arrow.

of the MN code), the channel capacity can be nearly saturated. However, for a finite block length, the main drawback of our algorithm is the overhead of the header which must be encoded and transmitted reliably. One has to remember that the size of the header scales with  $q^2$  where the precision of each element is of the order  $O(\log L)$ . This overhead is especially intolerable in the limit where  $q^2 \log L / L \sim O(1)$ . Note that this is indeed the situation even for very large messages,  $L = 10^6$ , and the largest taken autocorrelation length is only  $\log_2 q = 8$ .

In this section we explain how the Markovian joint s/c can be implemented without any side information. The key points are the special properties of KS construction 12) (Fig. 2): the first  $L_0$  rows of  $A$  are characterized by one non-zero element per row and column, where the first  $L_0$  rows of  $B$  are characterized by 2 non-zero elements. Furthermore, due to the systematic form of  $B$ , each row cannot be written as a linear combination of the other rows. Hence, the first  $L_0$  bits of the syndrome, Eq. (4.2), are equal (up to a simple permutation) to the source, with an effective flip rate,  $f_{eff}$ . For  $GF(2)$  for instance,  $Z_j = x_i + n_j + n_{j+1}$  ( $i$  marks the position of the nonzero element in the  $j^{th}$  row of  $A$ ), and  $f_{eff} = 2f(1 - f)$ . The first  $L_0$  symbols of  $Z$  are therefore a result of a Hidden Markov Model (HMM). The underlying transition matrix,  $T$ , generating the source sequence, can be estimated by means of the EM algorithm 20). This is a standard tool for solving such *Parametric Estimation* problems, which has linear complexity. Having  $T$  (approximately) revealed, the Dynamical Block Priors can be used as described in 7.2.

For the general construction of the MN algorithm, one adds/subtracts rows of the concatenated matrix  $[A, B]$  and the corresponding symbols in  $z$  (see Eq. (4.2)), such that the following situation is finally reached: The first  $L_0$  rows of  $A$  are the identity matrix, regardless of the construction of the first  $L_0$  rows of  $B$ . From the knowledge of the noise level  $f$  and the structure of  $i^{th}$  row of  $B$  one can now calculate the effective noise level,  $f_{i,eff}$ , of the  $i^{th}$  received source symbol. Since all  $\{f_{i,eff}\}$  are functions of a unique noise level  $f$ , one can again estimate the parameters of the Markovian process using some variants of the EM algorithm. Note, that in the general case the first  $L_0$  rows of  $B$  contain loops, hence the  $\{f_{i,eff}\}$ , are correlated. However, these correlations are assumed to be small as the typical loop size is of the order of  $O(\log(L))$  21).

## §9. Concluding remarks

The only remaining major drawback of the presented Markovian joint s/c coding is that the complexity of the decoder per message passing, scales as  $O(Lq^2/\log_2(q))$ , this may considerably slow down the decoder even for moderate alphabet size. Note however, that for large  $q$ , such that  $q^2 \geq L$ , and low entropy sequences, the transition matrix,  $T$ , is expected to be very sparse (consider  $q = 1024$  vs. block size of  $L = 100,000$ ). Taking advantage of the sparseness of  $T$ , the complexity of the decoder can be further reduced.

The one-dimensional Markovian joint s/c decoder can be easily extended to coding of a two-dimensional array of symbols or even to an array of symbols in higher dimensions. The complexity of the DBP decoder scales as  $L_0^d q^{2d+1}$ , where

$L_0^d$  is the number of blocks in the array, and  $d$  denotes the dimension<sup>22</sup>). Using Markovian and Bayesian assumptions, the complexity can be reduced to  $O(L_0^d q^2)$ .

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- 16) For rate 9/8, for instance, the chosen construction for the matrices  $A$  and  $B$  is as follows.  $A$  has sets of rows with 1, 2, 3, 5, 9 random non-zero elements per row. the number of rows from each type is  $(m/16, m/4, m/16, m/16, m/16)$ , where  $m$  is the number of rows in  $A$ . The structure of  $B$  is the same as illustrated in Fig. 2, but 1.75 is replaced with 7/9. We ran simulations for this construction with  $C_1 = C_2 = 0.7$  and the corresponding entropy is  $H_2 = 0.513$  and  $L = 9,000$ . The extrapolation of  $t_{med}$  indicates that the threshold of this code for large  $L$  is  $f_c \sim 0.057$ . In the separation scheme using *optimal compression and error correction schemes* and with  $f_c = 0.057$  ( $R_{i.i.d} = 0.618$ ), one can find that the overall inverse rate of the communication channel is  $1/R = 0.513/0.618 \sim 0.83$ , which is only about 6% below our joint s/c inverse rate  $1/R = 8/9 \sim 0.89$ . One must remember that our MN construction can be further optimized, and the critical channel noise is expected to be enhanced,  $f_c > 0.057$ .
- 17) In practice we define  $t_{med}$  to be the average convergence time of all samples with  $t \leq$  the median time.
- 18) The PPMZ software used can be downloaded from [www.cbloom.com/src/ppmz.html](http://www.cbloom.com/src/ppmz.html)
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