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RANDOM BINARY SEQUENCES

by

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ABSTRACT. Two coding problems are considered. One is to obtain binary codes with desirable auto-correlation functions. The other is to obtain families of such codes with desirable cross-correlation functions. The results of a random computer search for the codes are described. An approximation to the expected number of trials in a search for families of codes is also presented.

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FOREWORD

This report describes the results of a random computer search for binary codes. The search was motivated by consultation with T. A. Westaway of the Electronic Systems Branch, Systems Development Department, on the problem of obtaining binary codes of length 100 with desirable correlation properties. In anticipation of future problems in this area, data were obtained on codes of other lengths.

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This report is released at the working level. Because of the continuing nature of the research coding problem, refinements and modifications may later be made in this study.

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INTRODUCTION

In electronic communications problems one is sometimes interested in obtaining finite length sequences of 1's and -1's with small auto- and cross-correlation functions. One way of acquiring such sequences, or families of such sequences, is to generate them randomly on a high-speed digital computer recording the desirable ones and discarding the rest. This report describes the results of such a random search with emphasis on sequences of length 100.

THE SET OF SEQUENCES

Let X_n be the set of all 2^n sequences of 1's and -1's of length n . Let x denote the sequence (x_1, x_2, \dots, x_n) . For $x, y \in X_n$ let

$$c_k(x, y) = \begin{cases} \sum \{x_i y_{i-k+n} : 1 \leq i \leq k\} & \text{if } 1 \leq k \leq n \\ \sum \{x_{i+k-n} y_i : 1 \leq i \leq 2n-k\} & \text{if } n+1 \leq k \leq 2n-1 \end{cases} \quad (1)$$

For $1 \leq k \leq n$, $c_k(x, y)$ is the value of the correlation of the left-most k elements of x with the right-most k elements of y . For $n+1 \leq k \leq 2n-1$, $c_k(x, y)$ is the value of the correlation of the right-most $2n-k$ elements of x with the left-most $2n-k$ elements of y .

The $(2n-1)$ -tuple $c(x, y) = (c_1(x, y), c_2(x, y), \dots, c_{2n-1}(x, y))$ of integers is called the cross-correlation function of x . The following integer-valued functions on $X_n \times X_n$ will be of interest.

$$M(x, y) = \begin{cases} \max \{|c_k(x, y)| : 1 \leq k \leq 2n-1\} & \text{if } x \neq y, \\ \max \{|c_k(x, y)| : 1 \leq k \leq 2n-1, k \neq n\} & \text{if } x = y \end{cases} \quad (2)$$

For simplification we let $m(x) = M(x,x)$. Estimates of μ_m , the mean of the function m , and σ_m , the standard deviation of m for values of n ranging from 10 to 1,000 were obtained by random sampling in the spaces X_n . Table 1 lists for each n the mean and standard deviation of the sample from X_n as well as a measure of the difference of the mean from $n^{2/3}$. It is interesting to note how closely $n^{2/3}$ approximates the sample mean for n between 40 and 300.

TABLE 1.

n	Sample size	μ_m	$n^{2/3}$	$\frac{\mu_m - n^{2/3}}{n^{2/3}}$	σ_m
10	500	4.24	4.64	-0.086	1.31
20	500	6.97	7.37	-0.054	1.79
30	500	9.28	9.66	-0.039	2.12
40	500	11.49	11.70	-0.018	2.61
50	500	13.54	13.57	-0.002	2.86
60	500	15.49	15.33	0.010	3.25
70	500	17.05	16.99	0.004	3.42
80	500	18.54	18.57	-0.002	3.67
90	500	20.17	20.08	0.004	4.00
100	500	21.53	21.54	0.000	4.05
110	500	22.91	22.96	-0.002	4.32
120	500	24.49	24.33	0.007	4.30
130	500	25.69	25.66	0.001	4.53
140	500	27.22	26.96	0.010	4.58
150	500	28.32	28.23	0.003	4.74
200	500	33.74	34.20	-0.013	5.07
300	500	44.19	44.81	-0.014	6.28
500	300	60.02	63.00	-0.047	7.68
1,000	200	91.69	100.00	-0.083	11.35

SELECTING SINGLE SEQUENCES

The first practical problem motivating this study was to find sequences $x \in X_{100}$ such that $m(x)$ is small. Generation of 500 random sequences indicated that the probability that $m(x) \leq 15$ is about 0.026. Among the sequences chosen none satisfied $m(x) \leq 10$. On the UNIVAC 1108 computer 500 members of X_{100} can be chosen and the corresponding values of m computed in a little less than one minute. If one is interested in obtaining only sequences x such that $m(x)$

is less than or equal to some fixed bound b , then one discards the sequence x immediately after computing the first $c_k(x,x)$ such that $|c_k(x,x)|$ exceeds b . This eliminates unnecessary calculation of the $c_k(x,x)$ for unacceptable sequences thus reducing the running time on the computer. Of course, if $b_1 \leq b_2$, then the number of sequences satisfying $m(x) \leq b_1$ will be less than the number satisfying $m(x) \leq b_2$. Table 2 shows the results of computer runs selecting sequences with upper bounds of 13, 15, and 20 for $m(x)$ in the set X_{100} . In the case of the 80 sequences found with an upper bound of 13, none had upper bound ≤ 10 .

TABLE 2.

Bound	No. of sequences tested	No. of good sequences	Running time
13	30,000	80	10 min, 31 sec
15	5,000	150	3 min, 1 sec
20	900	374	2 min, 48 sec

SELECTING FAMILIES OF SEQUENCES

The second practical problem under consideration was to obtain a family $x^{(1)}, x^{(2)}, \dots, x^{(f)}$ of sequences from X_{100} such that $M(x^{(i)}, x^{(j)})$ is less than some bound b for $1 \leq i \leq j \leq f$. Here one is confronted with the problem of determining values of f and b for which the computer will generate a solution in a reasonable length of time.

Suppose the probability that $m(x) \leq b$ is p for a random sequence x . The value $m(x)$ is the maximum of the first 99 values of $|c(x,x)|$ since

$$c_k(x,x) = c_{200-k}(x,x) \quad (3)$$

i.e., the function $c(x,x)$ is symmetric about $k = 100$. However, for $M(x,y)$ to be less than or equal to b , one must have $|c_k(x,y)| \leq b$ for $1 \leq k \leq 199$. If one assumes that the functions c_1, c_2, \dots, c_{199} are statistically independent, then the probability that $M(x,y) \leq b$ for two random sequences x, y is approximately p^2 .

Now suppose one wishes to select randomly a family $x^{(1)}, x^{(2)}, \dots, x^{(f)}$ of sequences from X_{100} as described above. First one chooses sequences until a sequence $x^{(1)}$ satisfies $m(x^{(1)}) \leq b$. Next one chooses sequences until a sequence $x^{(2)}$ satisfies $M(x^{(1)}, x^{(2)}) \leq b$ and $m(x^{(2)}) \leq b$. Continuing recursively one chooses at stage r a sequence $x^{(r)}$ satisfying

$$\begin{aligned} M(x^{(1)}, x^{(r)}) &\leq b \\ M(x^{(2)}, x^{(r)}) &\leq b \\ &\vdots \\ M(x^{(r-1)}, x^{(r)}) &\leq b \\ m(x^{(r)}) &\leq b. \end{aligned} \tag{4}$$

Having chosen sequences $x^{(1)}, x^{(2)}, \dots, x^{(r-1)}$ the probability that a randomly chosen sequence x satisfies the restrictions of Eq. 4 on $x^{(r)}$ is approximately

$$\underbrace{p^2 p^2 \dots p^2}_r p = p^{2r-1}.$$

It should be noted here that $(x^{(1)}, \dots, x^{(r-1)})$ is not actually the result of a random selection from the cross-product space

$$X_{100} \times X_{100} \times \dots \times X_{100}$$

since the family $x^{(1)}, \dots, x^{(r-1)}$ already satisfies inequalities similar to Eq. 4. Thus, p^{2r-1} is possibly a poorer approximation than it might seem.

Suppose $x^{(r)}$ is the N_r th sequence tested after $x^{(r-1)}$. Then N_r is a random variable, and p^{1-2r} approximates the mean of N_r . Let N denote the number of sequences tested in order to select the entire family $x^{(1)}, x^{(2)}, \dots, x^{(f)}$. N is also a random variable, and

$$N = N_1 + N_2 + \dots + N_f. \tag{5}$$

Since the mean of N is the sum of the means of the N_i , we have the following approximation to μ_N , where μ_i is the mean of N_i .

$$\begin{aligned}\mu_1 + \dots + \mu_f &= p^{-1} + p^{-3} + \dots + p^{1-2f} \\ &= p^{-1}(1+p^{-2}+\dots+p^{-2(f-1)}) \\ &= p^{1-2f} \left(\frac{1-p^{2f}}{1-p^2} \right).\end{aligned}\quad (6)$$

Since p^{2f} will in general be very small, we have

$$\mu_N \sim p^{1-2f}(1-p^2)^{-1}.\quad (7)$$

Table 3 shows estimates of μ_N based on approximation (Eq. 7) for three values of b . In the table p is the probability that $m(x) \leq b$ as estimated by a sample of 500 members of X_{100} .

TABLE 3.

b	p_b	N
15	0.026	9.25×10^{45}
20	0.464	5.98×10^8
25	0.840	5.34×10^2

On the basis of Table 3 it was decided to place the bound b for families of sequences from X_{100} at 25. Table 4 shows the results of ten computer runs each selecting a family of 15 sequences with auto- and cross-correlation bounded by 25. At the bottom of Table 4 the means of the sample N 's and running times are given. In view of the various assumptions made in approximating N in Table 3 the difference between the approximation 534 for $b = 25$ in Table 3, and the sample μ_N of 1020 in Table 4 cannot be considered too surprising.

TABLE 4.

N	Running time
1,582	9 min 55 sec
971	6 min 34 sec
563	3 min 55 sec
934	6 min 6 sec
1,678	10 min 22 sec
824	5 min 18 sec
1,561	9 min 26 sec
798	5 min 7 sec
919	5 min 58 sec
368	2 min 41 sec
1,020	6 min 32 sec

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