

On the Computation of the Automorphisms Group of Some Optimal Codes Using Genetic Algorithm

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Abstract: Many research papers in coding theory have recently focused on designing high-rate codes or improving codes that exist through a better understanding and then improving the coding and decoding algorithm. As a result, this paper aims to investigate the computation of the Automorphisms groups of some optimal codes (e.g., some linear circulant codes where their distance meets the lower bound and nonlinear Nordstrom-Robinson (24, 28, 6) code). These Automorphisms groups provide information about the structure of the code, which aids in both the design and enhancement and improvement of decoding algorithms. A new genetic algorithm-based method is proposed, with a detailed description of its components, the fitness function, selection, crossover, and mutation, and is used to find an important collection of Automorphisms; the results obtained have shown that the proposed method is effective in finding stabilizers set for some types of codes.

Keywords: automorphism group, optimal codes, genetic algorithm, crossover, mutation.

用遺傳算法計算一些最優碼的自同構群

摘要：編碼理論方面的許多研究論文最近都集中在設計高速碼或通過更好地理解改進現有的碼，然後改進編解碼算法。因此，本文旨在研究一些最優碼的自同構群的計算（例如，一些距離滿足下限的線性循環碼和非線性諾德斯特龍-羅賓遜(24, 28, 6)碼）。這些自同構組提供有關代碼結構的信息，這有助於解碼算法的設計、增強和改進。提出了一種新的基於遺傳算法的方法，詳細描述了其組成部分、適應度函數、選擇、交叉和變異，並用於尋找重要的自同構集合；獲得的結果表明，所提出的方法可以有效地找到為某些類型的代碼設置的穩定器。

关键词：自同構群、最優碼、遺傳算法、交叉、變異。

1. Introduction and Preliminaries

There are several ways in coding theory that address its application, one of which is establishing the automorphism groups of codes, which allows us to determine the structure of the codes, classify them, and aid the decoding algorithm. This is still difficult because determining the whole automorphism group of codes is difficult, except for finite simple groups, which have been realized using sporadic groups [1] (e.g., the automorphism group of Golay Codes is Mathieu groups).

Recalling that the hamming distance between two codewords (vectors) c, c' in F_2^n is defined as the number of coordinates in which c and c' differ. A binary linear $[n, k, d]$ -code Cover F_2 is a k -dimensional subspace of the vector space F_2^n ,

$$d = d(C) = \min_{c \neq c' \in C} d(c, c') = \min_{c \in C \setminus \{0\}} wt(c) \quad (1)$$

and its generator matrix G is a $k \times n$ matrix whose rows are the basis of C .

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$$H_1 = \begin{bmatrix} 011100110000010111011010100000000000 \\ 111001100000101110110100010000000000 \\ 110011000001011101101001001000000000 \\ 100110000011111011010010000100000000 \\ 001100000111110110100101000010000000 \\ 011000001110101101001011000001000000 \\ 110000011100011010010111000001000000 \\ 100000111001110100101110000000100000 \\ 000001110011101001011101000000010000 \\ 000011100110010010111011000000001000 \\ 000111001100100101110110000000000100 \\ 00111001100100101110110000000000010 \\ 001110011000001011101101000000000001 \end{bmatrix} \quad (12)$$

1.3. Nordstrom-Robinson Code (NR)

Although there are infinite families of excellent nonlinear codes, no binary code compares to the nonlinear NR code (16, 8, 6). Snover has shown that NR code is the unique code of length 16, minimal distance 6, and 256 words. Different approaches to the construction of NR code has been established one of them is based on Golay code [24, 12, 8] (G) over F_2 , where the NR code consists of all vectors $y \in Z_2^{16}$ such that $xy \in G$ where

$x \in \{00000000, 11000000, 10100000, \dots, 10000001\}$.

Another approach is using $Z/4Z$ linear codes, which is proven by Forney et al., the binary image of the octal code (O_8) is the NR code, where the generator matrix is as follows [5]:

$$\begin{bmatrix} 10002111 \\ 01001213 \\ 00101321 \\ 00011132 \end{bmatrix} \quad (13)$$

where the binary image φ is defined as follows:

$$\varphi: Z_4 \rightarrow Z_2^2$$

$$\varphi(0) = 00, \varphi(1) = 01, \varphi(2) = 11, \text{ and } \varphi(3) = 10 \quad (14)$$

The generator matrix (15) is an extended Hamming code [8, 4, 4] over F_2 when octal code generator matrix entries are modulo 2. The Lee weight distribution of the linear Z_4 code is identical to the Hamming weight distribution of the F_2 image.

Now, we can obtain the generator matrix G_{NR} through the binary image φ of generator matrix vectors of O_8 code (14)

$$G_{NR} = \begin{bmatrix} 0100000011010101 \\ 0001000001110110 \\ 0000010001101101 \\ 0000000101011011 \end{bmatrix}$$

$G_{NR} \xrightarrow{\text{Standart form}} G_{NR}$

$$= \begin{bmatrix} 1000000011010101 \\ 0100000001110110 \\ 0010000001101101 \\ 0001000001011011 \end{bmatrix} \quad (15)$$

1.4. Genetic Algorithm

The GA is a member of the evolutionary algorithm family. C. Darwin observed that species evolution is dependent on two components: selection and reproduction, and the population of a GA evolves through genetic operators influenced by evolutionary biology. The reproduction of the fittest and most vigorous individuals is provided by the selection, while

reproduction is a phase in which evolution is carried out. Based on this evolution in nature, JH Holland invented the genetic algorithm in its first version, which was based on the programming of individuals in a binary system which improved significantly later. The task in such permutation problems (e.g., travel salesman problem (TSP), job-shop scheduling problem (JSP), bandwidth-reduction problem (BRP), and linear ordering problem (LOP)), which is a class of combinatorial optimization problems, is to arrange some objects in chromosomes in a certain order, with no duplicates. It should be done to optimize the objective function, where the representation of the chromosomes depends on [4]. GA solves the permutation problem by rapidly searching the search space. It employs the selection, crossover, and mutation operators, resulting in better chromosomes at the lowest possible cost [5]. The algorithm's effectiveness has been represented by several research papers, such as the research of Rajappa and Elsayed et al. Also, it has been demonstrated that using evolutionary algorithms to solve combinatorial optimization problems is effective [6-8]. There are powerful, nature-inspired algorithms such as the Gaining-sharing knowledge-based algorithm (GSK) [9-11], which have demonstrated superior results in solving optimization problems. The GA has several advantages such as:

- Only evaluate the objective function, regardless of its nature (e.g., continuity, derivation, and others), giving it more flexibility and a broader range of applications;

- Generation takes a parallel form by working on multiple points at once (population of size N), rather than a single iteration in classical algorithms;

- The use of probabilistic transition rules (selection, crossover, and mutation probabilities) rather than deterministic trajectories

Many researchers have indicated that comprehension of the GA parameters' interaction process, notably crossover probability, mutation probability, and population size, is the most important factor in evaluating the process. These variables are linked somehow and have an impact on GA efficiency. The optimal condition to use GA [18] is when there is diversity in the original population with a high crossover probability and a low mutation probability.

It is worth noting that the usual crossover operator cannot be used to solve permutation problems because chromosome ordering is crucial, and no genes should be duplicated or missed [12]. In addition, compared to other scenarios, it is more computationally expensive since a legalization step is necessary after each substring exchange for offspring with duplicate numbers. In this case, the time required to complete a crossover operation increases fast as chromosome size increases, reducing the efficiency of permutation-based GAs [13]. In their research publication [14], Chun Liu

and Andreas Kroll devised a genetic algorithm that did not employ the crossover operator. It is worth repeating that GA has been utilized to compute the minimum distance of linear block codes [15] and determine automorphisms set for some block codes like BCH and RQ codes of modest length.

2. Genetic Algorithm–Based Method

Our GA-based technique employs an encoding scheme that treats a permutation (chromosome) as a series of numbers ranging from 0 to the code length minus one. The following is how our method parameters work:

2.1. The Search Space and Fitness Function

The search space in which the GA-based method will search for permutations has $n!$ permutations (n is the code length). Each permutation matrix P_σ will be paired with its corresponding permutation for all permutations $\sigma \in S_n$.

$$P_\sigma = \sigma(I_n) \quad (16)$$

$S_c \subset C$ is a codewords set, such that, $\forall c_i \in S_c$,

$$c_i H^T = 0 \quad (17)$$

S_c is a matrix where its rows are formed by codewords.

$$S_c = \begin{bmatrix} c_1 \\ c_2 \\ \vdots \\ c_l \end{bmatrix} = \begin{bmatrix} c_{11} & c_{12} & \dots & c_{1n} \\ c_{21} & c_{22} & \dots & c_{2n} \\ \vdots & \vdots & \vdots & \vdots \\ c_{l1} & c_{l2} & \dots & c_{ln} \end{bmatrix} \quad (18)$$

Applying the action of S_n on $S_c, \forall \sigma \in S_n$ s.t,

$$y(S_c) = S_c P_y = \begin{bmatrix} c_{y(11)} & c_{y(12)} & \dots & c_{y(1n)} \\ c_{y(21)} & c_{y(22)} & \dots & c_{y(2n)} \\ \vdots & \vdots & \vdots & \vdots \\ c_{y(l1)} & c_{y(l2)} & \dots & c_{y(ln)} \end{bmatrix} \quad (19)$$

$$y(S_c) H^T = \begin{bmatrix} s_1 \\ s_2 \\ \vdots \\ s_l \end{bmatrix} \quad (20)$$

where H is the sparse parity check matrix

The permutation of S_c columns will generate another matrix of codewords if

$$\sigma(S_c) H^T = 0_{l \times (n-k)} \quad (5)$$

The fitness values of permutations will be used to choose the best permutations (individuals) using the fitness function, which is defined as follows:

$$f_\sigma = |\{C \cap \sigma(S_c)\}| = |\{s_i \in \{s_1, s_2, \dots, s_l\} | s_i = 0_{l \times (n-k)}\}| \quad (21)$$

The values of each permutation are computed using the fitness function (21) (a maximization function), with the best fitness value being equivalent to the best permutation that stabilizes the maximum of codewords in S_c [16].

Let ρ be a function defined on $S = \{0, 1, \dots, n-1\}$ such that:

$$\rho = \text{ChooseRandomly}(x, y), \text{ where } x, y \in S \quad (22)$$

$$\text{Offspring}[i] = \begin{cases} \rho(P_1[i], P_2[i]) & \text{when } i = 1 \\ \{ \{ \end{cases} \quad (23)$$

where $P_i = \text{Gene}_{i1}, \text{Gene}_{i2}, \dots, \text{Gene}_{in-1}$

The crossover operator that was shown above (23) is based on a random choice (22) with a probability of 0.5 for genes that do not exist in the offspring genes, otherwise choose the other, chosen to ensure that all produced individuals are within the search space and elements of S_n without relying on mutation due to the mutation operator's low probability.

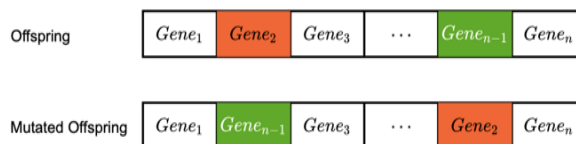


Fig. 1 Mutation operator

2.2. The Inputs and Outputs Method

The GA-based method works as follows:

Inputs:

- Codewords set S_c
- Initial population size N_i
- Number of generations N_{max}
- Crossover probability p_c
- Mutation probability p_m

Outputs:

- Set of Automorphisms permutations S_{Aut}

Data: Set of codewords S_c

Result: Set of Automorphisms S_{Aut}

$N_i \leftarrow$ Initial population of permutations σ_i randomly generated;

$N_g \leftarrow 1$;

while $N_g < N_{max}$ **do**

Evaluate the permutation fitness f_{σ_i} ;

Choose the N_e elite individuals from the sorted

current population into next

population f_{σ_i} ;

for $i=N_e$ to N_i , **do**

Select pair of elite individuals from N_e

individuals;

Apply the crossover and mutation operators;

end

$N_g \leftarrow N_g + 1$

end

Algorithm 1: GA-based method algorithm.

3. Results and Discussion

The default parameters of the GA-based method are given in Table 1, and the permutation is presented as a list of integers ranging from 0 to the length of the code minus one (each number representing a gene).

Table 1 Parameters of GA-based method

Parameter	Value
Initial population size	100
Selection	Elitism
Crossover probability	1
Mutation probability	0.07
Number of generations	200

Because every error-correcting code has an Automorphisms group, a set of Automorphisms permutations exists. We obtained a large number of stabilizers after running the algorithm. Tables 2 and 3, which include 15 Automorphisms permutations produced by our GA-based method for linear circulant [21, 14, 4] code and linear circulant [24, 16, 4] code, are cited as the example set of stabilizers for each code mentioned below.

Table 2 Automorphisms set of linear circulant [21, 14, 4] code of header 10010011101101

[4,9,16,3,7,12,8,14,2,10,20,5,15,6,19,18,13,1,11,0,17]
[14,7,2,15,10,12,20,13,0,8,19,11,3,17,9,5,4,1,18,16,6]
[2,1,0,3,13,11,6,19,8,9,20,5,12,4,14,15,16,17,18,7,10]
[13,3,0,11,8,4,1,17,20,5,18,2,19,6,7,10,14,15,16,9,12]
[19,20,15,13,18,4,9,10,1,7,5,12,14,2,8,17,11,0,6,3,16]
[7,4,0,12,9,15,10,19,16,13,14,18,3,1,20,11,8,17,5,2,6]
[17,18,13,11,7,10,8,9,14,12,5,16,19,1,6,2,20,15,4,0,3]
[7,3,0,12,2,6,10,1,20,5,18,14,4,19,16,17,8,11,13,9,15]
[13,3,0,10,8,4,15,12,5,20,9,2,19,6,7,11,14,1,16,18,17]
[3,0,11,7,4,1,14,9,18,5,16,19,6,17,12,8,10,13,15,2,20]
[18,7,12,16,17,2,1,8,15,5,13,4,10,9,3,19,14,20,11,6,0]
[13,9,16,18,19,12,1,17,20,0,2,11,15,6,7,3,4,8,5,10,14]
[9,11,1,16,6,0,12,18,3,10,19,7,13,14,17,5,8,4,20,15,2]
[15,13,8,4,9,11,20,5,7,6,14,2,10,3,1,16,18,17,19,0,12]
[0,1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20]

Table 2 Automorphisms set of linear circulant [24, 16, 4] code of header 1001011011011111

[5,4,3,6,1,0,7,2,10,17,8,15,20,21,12,23,18,9,16,11,22,13,14,19]
[0,5,6,3,4,1,2,7,18,9,10,23,22,21,14,15,16,17,8,19,20,13,12,11]
[6,1,0,3,2,5,4,7,15,20,21,12,11,10,17,18,13,14,19,8,9,16,23,22]
[0,1,2,3,4,5,6,7,8,17,10,23,12,21,14,19,16,9,18,15,20,13,22,11]
[6,3,0,1,2,7,4,5,20,19,8,17,16,11,22,21,18,15,14,13,12,23,10,9]
[0,1,2,3,4,5,6,7,8,9,16,23,12,13,20,19,10,17,18,15,14,21,22,11]
[4,5,2,7,0,1,6,3,22,21,20,15,18,17,10,23,14,13,12,11,16,9,8,19]
[2,7,4,1,6,3,0,5,16,11,12,13,14,19,18,17,22,23,10,9,8,15,20,21]
[3,4,1,2,7,0,5,6,23,22,13,20,15,18,17,16,21,12,11,10,9,8,19,14]
[1,0,3,2,5,4,7,6,16,17,18,15,14,21,12,11,8,9,10,23,22,13,20,19]
[1,2,7,0,5,6,3,4,13,14,15,8,17,10,23,12,19,20,21,22,11,16,9,18]
[3,2,5,0,7,6,1,4,18,19,20,21,22,23,10,9,14,15,8,17,16,11,12,13]
[7,2,1,0,3,6,5,4,18,15,20,13,12,23,16,9,14,19,8,17,10,11,22,21]
[6,7,4,5,2,3,0,1,16,23,12,13,14,19,18,9,22,11,10,17,8,15,20,21]
[1,0,3,6,5,4,7,2,16,17,8,19,20,21,22,11,18,9,10,23,12,13,14,15]

Table 4 contains, as an example, a set of 15 Automorphisms permutations produced by our GA-based method for NR (24,28,6) code. To be mentioned, any combination of two Automorphisms permutations is an Automorphism permutation for the abovementioned codes. This implies that if SAut includes all of the automorphism group's generators, then all of the others can be obtained.

Remarque, the octal code O_8 cannot supersedes the NR code due to their Automorphisms groups orders ($|\text{Aut}(\text{NR})|=16 \times 7!$ and $|\text{Aut}(O_8)|=|\text{GL}(4,2)|=1344$).

Table 3 Automorphisms set of NR (16,256,6) code

[8,3,9,2,5,4,6,7,0,1,13,15,11,10,12,14]
[8,9,1,2,4,7,6,5,0,3,15,14,10,12,11,13]
[8,1,2,9,5,7,4,6,0,3,10,12,11,13,15,14]
[8,2,9,3,7,4,6,5,0,1,11,15,13,14,12,10]
[0,2,3,1,6,4,7,5,8,9,12,13,14,15,10,11]
[0,9,1,2,4,6,7,5,8,3,15,14,10,12,11,13]
[8,2,3,9,7,5,6,4,0,1,12,14,13,15,11,10]
[0,3,2,1,4,6,5,7,8,9,12,11,10,15,14,13]
[8,1,2,3,5,4,6,7,0,9,10,11,12,13,14,15]
[0,9,3,1,6,4,5,7,8,2,13,12,14,10,15,11]
[0,3,2,1,7,4,6,5,8,9,12,11,10,15,14,13]
[0,1,9,2,6,4,7,5,8,3,15,13,11,12,10,14]
[0,9,2,1,4,5,6,7,8,3,11,12,10,14,15,13]
[8,1,9,3,6,5,7,4,0,2,15,11,13,12,14,10]
[0,2,1,3,4,6,7,5,8,9,10,15,14,13,12,11]

4. Conclusion

In this research, a genetic algorithm-based method for determining an important Automorphisms set of certain provided linear circulant codes, as well as the Nordstrom-Robinson (24, 28, 6) code, is suggested, which may be utilized to improve their decoding algorithms (the hard decision algorithm and the soft decision algorithm). The abovementioned codes yielded significant results, and we did not need a requirement on either the level of the search space or the level of the fitness function. This genetic algorithm-based method can be used to find a set of automorphisms groups of codes, and it can be completely relied on in decoding algorithms based on the set of automorphism permutations, especially in small and medium-length it can be used in a variety of codes. Our future work will improve the algorithm's efficiency and accelerate its function in finding automorphism groups for various types and lengths of codes by searching for a more efficient fitness function, adjusting the selection operator, the stochastic crossover operator, and the stochastic crossover operator.

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