Solving Linear Equations Systems Using Genetic Algorithm

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Abstract:

Genetic Algorithms (GA's) are a class of optimization algorithms. GA’s attempts to solve problems through modeling a simplified version of genetic process. There are many problems for which a GA approach is useful.

This paper aims to solve Linear Equations System (LES) for any number of variables using the GA. The application of this paper represented by cryptanalysis application, this done by attacking stream cipher systems, choosing one Linear Feedback Shift Register (LFSR), since its considered as a basic unit of stream cipher systems, in the performance of GA. The application divided into two stages, first, constructing LES's for the LFSR, and the second, is attacking the variables of LES's which they are also the initial key values the of LFSR.
1. Introduction

Cryptanalysis is the science and study of methods of breaking ciphers. It is a system identification problem, and the goal of Cryptography is to build systems that are hard to identify [1]. To attack a cryptographic system successfully the cryptanalysis is forced to be based on subtle approaches, such as knowledge of at least part of the text encrypted, knowledge of characteristic features of the language used,..., with some luck. The Cryptosystem are the systems which use the encryption and decryption processes.

The GA is an adaptive search method that has the ability for a smart search to find the best solution and to reduce the number of trials and time required for obtaining the optimal solution. The practicality of using the GA is to solve complex problems compared with traditional search techniques.

In this work, a LES of LFSR (which is considered as a basic unit of stream cipher systems) is constructed then solve it using the genetic algorithm.

2. Modern Cryptosystems

There are essentially two different types of cryptographic systems (cryptosystems), these cryptosystems are: public key and secret key cryptosystems [2]. First let us redefined some important notations:

- P is the Plaintext message and C is the Ciphertext message.
- Key space K: a set of strings (keys) over some alphabet, which includes the encryption key $e_k$ and the decryption key $d_k$.
- The Encryption process (algorithm) $E$: $E_{e_k}(P) = C$.
- The Decryption process (algorithm) $D$: $D_{d_k}(C) = P$.
- The algorithms E and D must have the property that: $D_{d_k}(C) = D_{d_k}(E_{e_k}(P)) = P$.

The public key cryptosystem also called asymmetric cryptosystems. In a public key (non-secret key) cryptosystem, the encryption key $e_k$ and decryption key $d_k$ are different, that is $e_k \neq d_k$. The secret Key Cryptosystem also called symmetric cryptosystems. In a conventional secret-key cryptosystem, the same key ($e_k = d_k = k \in K$), called secret key, used in both encryption and decryption; we are interest in this type of cryptosystems. The stream cipher systems one of the important types of the secret key cryptosystems [3].
3. Stream Cipher systems

In stream ciphers, the message units are bits, and the key is usual produced by a random bit generator (see fig.1). The plaintext is encrypted on a bit-by-bit basis.

![Stream cipher system diagram]

The key is fed into random bit generator to create a long sequence of binary signals. This “key-stream” $k$ is then mixed with plaintext $m$, usually by a bit wise XOR (Exclusive-OR modulo 2 addition) to produce the ciphertext stream, using the same random bit generator and seed.

Linear Feedback Shift Register (LFSR) systems are used widely in stream cipher systems field. A LFSR System consists of two main basic units. First, is a LFSR function and initial state values. The second one is, the Combining Function (CF), which is a boolean function. Most of all stream cipher systems are depend on these two basic units. Most practical stream-cipher designs center around LFSR. In the early days of electronics, they were very easy to build. A shift register is nothing more than an array of bit memories and the feedback sequence is just a series of XOR gates. A LFSR-based stream cipher can give a lot of security with only a few logic gates [4].

4. Genetic Algorithms (GA’s)

Genetic Algorithms (GA's) are search algorithms based on the mechanics of natural selection and natural genetics.
They combine survival of the fittest among string structures with a structured yet randomized information exchange to form a search algorithm with some of the innovative flair of human search [5]. GA is an iterative procedure, which maintains a constant size population of candidate solution. During each iteration step (Generation) the structures in the current population are evaluated, and, on the basis of those evaluations, a new population of candidate solutions formed. The basic GA cycle shown in fig. 2 [6].

An abstract view of the GA is:

Generation=0;
Initilaize G(P); \{G=Generation ; P=Population\}
Evaluate G(P);
While (GA has not converged or terminated)
    Generation = Generation + 1;
    Select G(P) from G(P-1);
    Crossover G(P);
    Mutate G(P);
    Evaluate G(P);
End (While)
Terminate the GA [6].
5. Constructing a Linear Equations System

Let’s assume that the tested LFSR is maximum LFSR (m-LFSR), then its period is \( P=2^r-1 \), where \( r \) is LFSR length. Let \( SR_r \) be a single LFSR with length \( r \), let \( A_0=(a_1,a_2,\ldots,a_r) \) be the initial value vector of \( SR_r \), s.t. \( a_j, 1 \leq j \leq r \), be the component \( j \) of the vector \( A_0 \), in another word, \( a_j \) is the initial bit of stage \( j \) of \( SR_r \), let \( C_0^T=(c_1,\ldots,c_r) \) be the feedback vector, \( c_j \in \{0,1\} \), if \( c_j=1 \) that means the stage \( j \) is connected else it’s not. Let \( S=\{s_i\}_{i=0}^{m-1} \) be the sequence (or \( S=(s_0,s_1,\ldots,s_{m-1}) \) read “S vector”) with length \( m \) generated from \( SR_r \). The generation of \( S \) depending on the following equation:

\[
s_i = a_i = \sum_{j=0}^{r} a_{j-r}c_j \quad i=0,1,\ldots \quad (1)
\]

Equation (1) represents the linear recurrence relation [7].

The objective is finding \( A_0 \), when \( r, C_0 \) and \( S \) are known. Let \( M \) be a \( r \times r \) matrix, which is describes the initial phase of \( SR_r \):

\[
M=(C_0 | I_{r \times (r-1)})\quad \text{where} \quad M^0=I.
\]

Let \( A_1 \) represents the new initial of \( SR_r \) after one shift, s.t.

\[
A_1=A_0 \times M=(a_1,a_2,\ldots,a_r)\begin{pmatrix} c_1 & 1 & L & 0 \\ c_2 & 0 & L & 0 \\ \vdots & \vdots & \vdots & \vdots \\ c_r & 0 & L & 0 \end{pmatrix}=(\sum_{j=0}^{r} a_{j-r}c_j,a_1,\ldots,a_{1-r}).
\]

In general,

\[
A_i=A_{i-1} \times M, \quad i=0,1,2,\ldots \quad (2)
\]

Equation (2) can be considered as a recurrence relation, so we have:

\[
A_i=A_{i-1} \times M=A_{i-2} \times M^2=\ldots=A_0 \times M^i \quad (3)
\]

The matrix \( M^i \) represents the \( i \) phase of \( SR_r \), equations (2) and (3) can be considered as a Markov Process s.t., \( A_0 \), is the initial probability distribution, \( A_i \) represents probability distribution and \( M \) be the transition matrix [8].
notice that:

\[ M^2 = [C_1C_0 | I_{r-r-2}] \]

and so on until get \( M^i = [C_{i-1} \ldots C_0 | I_{r-r-i}] \), where \( 1 \leq i < r \).

When \( C_0 = C_{C_0} \) then \( M^{P+1} = M \).

Now let’s calculate \( C_i \) s.t.

\[ C_i = M \times C_{i-1}, \ i=1,2,... \quad (4) \]

Equation (1) can be rewritten as:

\[ A_0 \times C_i = s_i, \ i=0,1,..,r-1 \quad (5) \]

if \( i=0 \) then \( A_0 \times C_0 = s_0 \) is the 1st equation of the LES,

if \( i=1 \) then \( A_0 \times C_1 = s_1 \) is the 2nd equation of the LES, and

if \( i=r-1 \) then \( A_0 \times C_{r-1} = s_{r-1} \) is the rth equation of the LES.

In general:

\[ A_0 \times \Psi = S \quad (6) \]

\( \Psi \) represents the matrix of all \( C_i \) vectors s.t.

\[ \Psi = (C_0, C_1, \ldots, C_{r-1}) \quad (7) \]

The LES can be formulated as:

\[ A = [\Psi^T | S^T] \quad (8) \]

\( A \) represents the extended (augmented) matrix of the LES.

**Example (1)**

Let the SR4 has \( C_0^T = (0,0,1,1) \) and \( S=(1,0,0,1) \), by using equation (4), we get:

\[
C_1 = M \times C_0 = \begin{pmatrix}
0 & 1 & 0 & 0 \\
0 & 0 & 1 & 0 \\
1 & 0 & 0 & 1 \\
1 & 0 & 0 & 0
\end{pmatrix}
\begin{pmatrix}
0 \\
0 \\
0 \\
1
\end{pmatrix} = \begin{pmatrix}
0 \\
1 \\
1 \\
0
\end{pmatrix}, \text{ in the same way, } C_2 = \begin{pmatrix}
1 \\
1 \\
0 \\
0
\end{pmatrix}, C_3 = \begin{pmatrix}
1 \\
0 \\
1 \\
1
\end{pmatrix}
\]

From equation (6) we have:
(0 0 1 1)
(0 1 1 0)
(1 1 0 1)
(1 0 0 1)

\[ \mathbf{A}_0 = (1,0,0,1), \] this system can be written as equations:

\[ a_3 + a_4 = 1 \]
\[ a_2 + a_3 = 0 \]
\[ a_1 + a_2 = 0 \]
\[ a_1 + a_3 + a_4 = 1 \]

Then the LES after using formula (8) is:

\[
\mathbf{A} = \begin{bmatrix}
0 & 0 & 1 & 1 & 1 \\
0 & 1 & 1 & 0 & 0 \\
1 & 1 & 0 & 0 & 0 \\
1 & 0 & 1 & 1 & 1
\end{bmatrix}
\] ....(9)

6. Use GA to Break Stream Cipher System

The GA will be used to solve LES's of LFSR with length \( r \), \( m=r \) equations are needed to solve the LES.

6.1 Coding Scheme

A LES is decoded by binary representation. As an example, the equation \( a_2 + a_5 = 1 \) of LFSR with length 5 decoded by the equation string (01001-1), where the absolute value (right side) of the equation is the real key of the LFSR. These equations are constant for fixed LFSR's length and connection function. As this representation indicates, the size of the equations space is \( 2^m - 1 \) (ignoring the zero string). By increasing the number of bits that are used for representing one continuous variable the accuracy of the representation can be increased [9].
6.2 Initial Population

For the initialization process we can initialize the population by a random sample of combinations of 0 and 1 with m-string length represents the probable initial values LFSR’s. The creation of the population must submit to what we called non-zero initial condition. The Initial Population Algorithm shown in fig.3.

\[ s_j := \text{RANDOM}(2); \]

6.3 Evaluation Function (Fitness Function)

This function is used to determine the “best” representation. The process of the evaluation function selection is as follows:

1. From population, a chromosome \( k, k=1,\ldots,m \) initial string of length \( m \) bits, so we get the string \( X_k = (X_{k1}, X_{k2}, \ldots, X_{km}) \).
2. The string bit \( X_{kj} \) product with corresponding equation string bit \( Y_{ij} \), where \( 1 \leq j \leq m \) s.t. the equation string is \( Y_i = (Y_{i1}, Y_{i2}, \ldots, Y_{im}) \) and calculate the observed value:

\[
O_{ki} = X_{k1} \odot Y_{i1} \odot X_{k2} \odot Y_{i2} \odot \ldots \odot X_{km} \odot Y_{im} = \sum_{j=1}^{m} X_{kj} \odot Y_{ij}
\]

3. Compare the observed value \( O_{ki} \) with key value \( K_i \) which represents the known output value of the cryptosystem, by using mean absolute error (MAE) s.t.
4. The Fitness value is

\[ \text{Fitness}_k = 1 - \text{MAE}_k = 1 - \frac{1}{m} \sum_{i=1}^{m} |O_{ki} - K_i| \]  

\[ \text{...}(12) \]

where

- \( m \): The length of the chromosome string or equation string.
- \( X_{kj} \): is the initial value \( j \) in String \( X_k \).
- \( Y_{ij} \): is the equation variable \( j \) in the string \( Y_i \).
- \( O_{ki} \): is the observed value \( i \) of string \( X_k \) calculated from equation (10).
- \( K_i \): is the key bit (actual value) \( i \).

When the measured (observed) value \( O_{ki} \) matches the key bit \( K_i \), for all \( 1 \leq i \leq m \), then the summation terms \( \text{MAE}_k \) in equation (11) evaluate to 0 so the fitness value is 1. The fact that a fitness value of 0 is never achieved does not affect the algorithm since high fitness values are more important than low fitness values. As a result, the search process is always moving towards fitness values closer to or equal 1. The steps of the Fitness Algorithm are shown in fig. 4:

**Fitness Algorithm**

**INPUT**: 1. READ X vector; \{(Initial string with length \( m \) from population\)

2. READ Y vector; \{(Equation string from data base file\)

3. READ K vector ;\{Actual key=absolute value of LES\)

**PROCESS**: 4. FOR \( i = 1 \) TO \( m \) DO

4.1 \( O_i := \sum_{j=1}^{m} X_j \ast Y_j \); \{XOR sum, \( O_i \) is observed key\}

4.2 \( \text{Dif}_i := |O_i - K_i| \);
6.4 Genetic Operators

I. Selection Operator

This method uses the roulette wheel selection method. The string with high fitness has a higher probability of contributing one or more offspring to the next generation.

II. Crossover Operator

Given a population of initial string, each one with a fitness value, the algorithm progresses by a high fitness value selecting two initial strings for mating. The two parents generate two children using crossover operation. The breeding process is achieved using the single (one)-point crossover select randomly one point in the range $[1, 2, ..., n]$, where $n$ is the length of the string. The first offspring is produced by deleting the crossover fragment of the first parent from the first and inserting the crossover fragment of the second parent at the crossover point of the first point. The second offspring is produced in a symmetric manner [10]. The Crossover Algorithm shown in Fig. 5.

**Crossover Algorithm**

```plaintext
INPUT : 1. READ Parent1, Parent2;
PROCESS  : 2. REPEAT
  2.1 x := RANDOM(m);
  UNTIL x > 1;
  3. Child1 := copy(parent1, 1, x) + copy(parent2, x + 1, m);
```

Fig.5 Crossover algorithm

III. Mutation Operator

After the new generation has been determined, the initial strings are subjected to a low rate mutation process. For this study when a bit in the initial string is selected for mutation randomly, it is swapped with the bit in the right (or left). If the fitness of the string is in the top half of the population, the character is
swapped with the character to its right. Of course, the swapping process subjects to non-zero initial condition.

Fig. 6 shows the Mutation Algorithm.

6.5 Genetic Parameters

The following parameters are been used: population size = 10, probability of crossover $P_c=0.7$, probability of mutation $P_m=0.05$, and 100 or 200 generations.

7. Experimental Results

Two stopping criterions are be used to stop the GA cryptanalysis system, first criterion, two hundred generations are enough to reach this level of fitness. The second, when the fitness value reaches (1.0), so no need to reach the high number of generation. The algorithm was fast enough that this took less than a minute on P4 PC. Let’s use 11 stages-LFSR, which has $1+x^2+x^{11}$ as characteristic primitive polynomial. The initial key value is: 10000000001. Table (1) shows the 11-stage equations of LES for single LFSR with binary representation.
Table (1) The 11-Stage Equations of LES for Single LFSR.

<table>
<thead>
<tr>
<th>I</th>
<th>Equation</th>
<th>Binary representation</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>$a_2 + a_{11} = 1$</td>
<td>01000000001 1</td>
</tr>
<tr>
<td>2</td>
<td>$a_1 + a_{10} = 1$</td>
<td>10000000010 1</td>
</tr>
<tr>
<td>3</td>
<td>$a_2 + a_9 + a_{11} = 1$</td>
<td>0100000101 1</td>
</tr>
<tr>
<td>4</td>
<td>$a_1 + a_8 + a_{10} = 1$</td>
<td>1000001010 1</td>
</tr>
<tr>
<td>5</td>
<td>$a_2 + a_7 + a_9 + a_{11} = 1$</td>
<td>0100010101 1</td>
</tr>
<tr>
<td>6</td>
<td>$a_1 + a_6 + a_8 + a_{10} = 1$</td>
<td>1000101010 1</td>
</tr>
<tr>
<td>7</td>
<td>$a_2 + a_5 + a_7 + a_9 + a_{11} = 1$</td>
<td>0101010101 1</td>
</tr>
<tr>
<td>8</td>
<td>$a_1 + a_4 + a_6 + a_8 + a_{10} = 1$</td>
<td>1001010101 1</td>
</tr>
<tr>
<td>9</td>
<td>$a_2 + a_3 + a_5 + a_7 + a_9 + a_{11} = 1$</td>
<td>0110101010 1</td>
</tr>
<tr>
<td>10</td>
<td>$a_1 + a_2 + a_4 + a_6 + a_8 + a_{10} = 1$</td>
<td>1101010101 1</td>
</tr>
<tr>
<td>11</td>
<td>$a_1 + a_2 + a_3 + a_5 + a_7 + a_9 + a_{11} = 0$</td>
<td>1110101010 1</td>
</tr>
</tbody>
</table>

For this example, only 10 initial keys were in the gene pool. The system began by generating 10 random initial key as shown:

Key 1: 00111000011 $\rightarrow$ Fitness: 0.64
Key 2: 10001011111 $\rightarrow$ Fitness: 0.55
Key 3: 11111101111 $\rightarrow$ Fitness: 0.55
Key 4: 10100111101 $\rightarrow$ Fitness: 0.64
Key 5: 00000011110 $\rightarrow$ Fitness: 0.45
Key 6: 00111110101 $\rightarrow$ Fitness: 0.36
Key 7: 01110110110 $\rightarrow$ Fitness: 0.55
Key 8: 11110100110 $\rightarrow$ Fitness: 0.45
Key 9: 11000111110 $\rightarrow$ Fitness: 0.73
Key 10: 11011101110 $\rightarrow$ Fitness: 0.55

The average fitness is 0.55. The best of these keys (key9) has a fitness value: 0.73.

After 31 generation, the pool begins to converge at a high rate of speed:
Key 1: 10001010001† Fitness: 0.82
Key 2: 10010100001† Fitness: 0.91
Key 3: 10010000001† Fitness: 0.73
Key 4: 10010000001† Fitness: 0.82
Key 5: 00101110001† Fitness: 0.64
Key 6: 10100100010† Fitness: 0.45
Key 7: 10100100011† Fitness: 0.64
Key 8: 10100100001† Fitness: 0.55
Key 9: 10101000001† Fitness: 0.73
Key 10: 10010100101† Fitness: 0.45

Average fitness has risen to 0.70 with the best key (key2) coming in at 0.91.

By generation 50, the gene pool looks like:

Key 1: 10000000001† Fitness: 1.00
Key 2: 10101000001† Fitness: 0.82
Key 3: 10010100001† Fitness: 0.91
Key 4: 10100000001† Fitness: 0.82
Key 5: 10101100001† Fitness: 0.64
Key 6: 10011010001† Fitness: 0.73
Key 7: 10000100001† Fitness: 0.36
Key 8: 10100010001† Fitness: 0.82
Key 9: 10000100001† Fitness: 0.73
Key 10: 10111001001† Fitness: 0.73

The average fitness is at 0.75. key1 turns out to have the highest fitness and on examination is the exact key. In table (2) we provide the generation number for which noted improvement in the evaluation function, together with the value of the function.
Table (2) Results of 50 Generations for Single LFSR.

<table>
<thead>
<tr>
<th>Gen.</th>
<th>Fitness</th>
<th>Average</th>
<th>key</th>
<th>Best Initial Key</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>0.73</td>
<td>0.55</td>
<td>9</td>
<td>11001111110</td>
</tr>
<tr>
<td>17</td>
<td>0.82</td>
<td>0.60</td>
<td>2</td>
<td>10010001001</td>
</tr>
<tr>
<td>31</td>
<td>0.91</td>
<td>0.70</td>
<td>2</td>
<td>10010100001</td>
</tr>
<tr>
<td>50</td>
<td>1.00</td>
<td>0.75</td>
<td>1</td>
<td>10000000001</td>
</tr>
</tbody>
</table>

The best initial key after (11) generations was (and equal to the real initial key):

1 0 0 0 0 0 0 0 0 1

8. Design of GA Cryptanalysis System

The GA cryptanalysis system can be view as two main parts, first, is the LES constructing which described LES constructing algorithm which is shown in Fig.7.

**LES Constructing Algorithm**

**INPUT** :
1. READ $A_0$; \(\text{\{}\text{ initial value of LFSR }\}\)
2. READ $C_0$ vector;
3. $M := (C_0 ||_{r-1})$;
4. READ $A_0$ vector; \(\text{\{}\text{ Initial values of LFSR }\}\)

**PROCESS** :
5. FOR $i := 0$ TO $m-1$ DO
   5.1 $C_i := M \cdot C_{i-1}$;
   5.2 $s_i := A_0 \cdot C_i$;

Fig. 7 LES Constructing Algorithm
The second part is the GA cryptanalysis part, which illustrated in GA cryptanalysis algorithm, shown in fig. 8:

**Fig. 8 GA Cryptanalysis Algorithm**

<table>
<thead>
<tr>
<th>GA Cryptanalysis Algorithm</th>
</tr>
</thead>
<tbody>
<tr>
<td>INPUT :</td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td>INITILIZE :</td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td>PROCESS :</td>
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<td></td>
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</tbody>
</table>

9. Conclusions

This research concludes the following aspects:

1. Although the proposed system is employed for small shift register length (r≤11), it was provide the base of building GA cryptanalysis system valid for long shift register attacking.

2. As be recommended, the mutation probabilities must be as low as possible and probability of crossover must be high in order to approach the correct or heuristic key of the LES obtained from the mentioned LFSR.

3. As a logical mathematical situation, if the proposed system gives a fitness value less than 1.0, this mean, no results obtained so we must run the system a gain, since the LES must has unique solution for fixed absolute values, no another solution gives fitness equal 1.0.

4. Percentages reported are based on number of tests and different numbers of the tests must be always used, and that what will done in this research.
References

حل نظام المعادلات الخطية باستخدام الخوارزمية الجينية

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المتخصصة:

تمثل مجموعة الخوارزميات الأسئلية. الخوارزمية الجينية تحاول حل المسائل من خلال بناي نموذج جبل بسيط من العملية الجينية. لقد نجحت الخوارزمية الجينية في حل الكثير من المسائل. وبالطبع فإن هذه الخوارزمية ستكون غير تقليدية في حالة تكون تحليل الشفرة هو أحد هذه المسائل.

هذا البحث يهدف إلى حل نظام المعادلات الخطية لأي عدد من المتغيرات باستخدام الخوارزمية الجينية. إن مجال تطبيق Stream (Cryptanalysis) (Ciphertext) (Linear Feedback Shift (Cipher Systems) (Register) باعتبار الوحدة الأساسية التي تدخل في بناء نظام التشغيل الإسبابي، معتمدين على أجز الخوارزمية الجينية. التطبيق في هذا البحث يتم في مرحلتين، الأولى، تمثل ببناء نظام معادلات خطية من مخرجات المسجل الزاحف، والمرحلة الثانية هي حل نظام المعادلات الخطية ومعرفة قيم المجاهيل والتي تمثل قيمة المفتاح الابتدائي للمسجل الزاحف.