

# Inspired Feistel DNA Based Crypto System Using D-Box

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## Abstract

DNA Cryptography is one of the encouraging aspects in the field of DNA Computing. DNA Computing or Bio Molecular Computing is used to describe the properties of DNA in parallel computation. DNA is used not only as a computation tool but also as an information carrier. The inherent property of storing large information nearly Terabytes of data into a single gram of DNA, the random nature of DNA making the code unbreakable made the researchers give higher priority to DNA Computation. As we all know traditional crypto algorithms are broken, there is a need of harnessing the biology, chemistry and computer science into single discipline so that it gives better security to the information.

In this paper, we proposed a model which uses Feistel Cipher implementation and adding the Shannon properties of Confusion and Diffusion using DNA Codons. One more characteristic of the proposed scheme is it used randomly generated key in each round so that it is highly impossible to detect the plaintext in a cryptanalytic attack. It is also observed that when the performance of the proposed algorithm is measured through avalanche effect it is found that it is better than some of the existing algorithms.

**Keywords:** DNA Cryptography, Symmetric Cryptography, Feistel Cipher, DNA Codons.

## INTRODUCTION

The advancements in the technology look towards the improvement in the security issues. Many of the fields like banking sector, e-mailing system, etc., needs to increase the security policies to gain trust from the customers. To provide security to the information, they use cryptographic techniques. Cryptography is an art of converting the text information into unreadable format. The process of converting the text data into unscrambled message is called as Encryption process and the reverse process is called as decryption process. The unscrambled message is nothing but cipher text. Many cryptographic approaches exist to provide security to the information, namely Modern Cryptography, Quantum Cryptography and DNA Cryptography. In modern cryptography, it faced difficulty in prime factorization and in Quantum cryptography deals with photons and the signals are limited to 90 miles [1]. After that, Gehani et.al.[2] discovered the DNA structure used to provide security to the information. He designed DNA based cryptography with substitution and One Time Pads. DNA cryptography which is a branch of DNA computing is a new technique which secures the data using biological structure. DNA Computing uses DNA sequences for computation instead of traditional silicon based

chips. The year 1994 is the new era in the field of computer science when Adleman used the bio molecular technology to solve the mathematical complex problem such as Hamiltonian path [3] which is a NP-complete problem. He demonstrated the problem using seven cities and each city named as a DNA sequence. By using the process of DNA synthesizing, he generated all possible routes within lesser time when compared to traditional approach and this process is called massive parallelism. The approach of massive parallelism using DNA helped many researchers to solve several NP Hard problems. With the inspiration of Adleman, Lipton showed a Boolean formula to solve the satisfiability problem known as SAT[4] using DNA. Solving complex mathematical problems in all possible solutions at once is the central idea of a DNA computer, since it stores 700 terabytes data in 1g of DNA which is proved by George Church's team of Harvard University[5]. The idea behind using cryptography in the arena of biology is to come up with a remarkable technology which provides unbreakable algorithms. But the limitation of these algorithms are that it lacks the theoretical approach which become a hurdle to model the good DNA cryptographic schemes[6]. The present paper provides the DNA based security based upon the traditional Feistel cipher that uses the D-box, a key element in this algorithm and the encryption process is done based on the number of rounds required. The key is also generated randomly in each round. The detailed process is described in the next sections. The Section2 describes the related works done based on DNA cryptography. In Section 3 and 4 the scope of proposed scheme and the implementation results are shown. The Section 5 and 6 describes experimental results, the performance analysis of proposed algorithm and comparison of avalanche effect of existing algorithms with proposed scheme. Finally, Section 7 is the conclusion of the paper.

## BIOLOGICAL BACKGROUND

DNA is an acronym for Deoxyribonucleic Acid which is a hereditary element in human beings and all other living organisms. It is made up of nucleotides. Each nucleotide contains a Sugar group, a Phosphate group and a Nitrogen base. The four Nitrogen bases which are called as Adenine (A), Thymine (T), Cytosine(C) and Guanine (G)(Figure 1). The entire information is represented in the form of these DNA four bases as against the traditional methodology (0's and 1's)[7,8]. Generally, DNA exists as double helix in its nature by pairing the complimentary bases from a single strand. These DNA bases pair up i.e., A with T and C with G according to Watson-Crick Complementary rule. These pairs can be called as base pairs. The central dogma of biological

structure for processing the genes into proteins is described in two step process: one is transcription and other is translation. In the transcription phase the DNA bases are converted into RNA having four bases. The difference between DNA and RNA is, instead of Thymine, Uracil (U) is used in RNA. In the translation phase each three nucleotide bases can form an amino acid which is useful in protein synthesization. The amino acid also called as Codon. There are total 64 codons can be formed from the 4 bases. All 64 Codons can form only 22 amino acids[9,10].

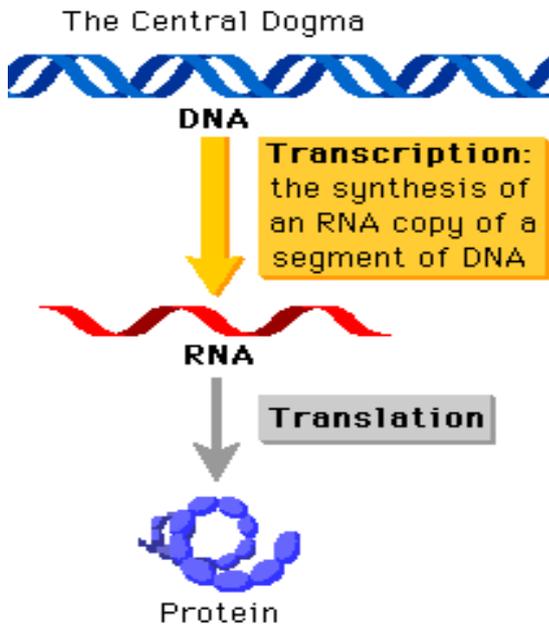


Figure 1: Central Dogma of DNA

Table 1: DNA Table

Base	Gray Code
A	00
C	01
G	10
T	11

**RELATED WORK**

The field of network security always looks for unbreakable cryptographic algorithms to protect the information while storing the data on a cloud or while passing the data on a network or some other security issues. In this regard, many researchers performed more work on cryptographic approaches and found that DNA carries the information more confidential and the size of DNA strand is large than the traditional methods. There are numerous works taking place for the last few years. In the beginning, Gehani et.al., introduced DNA based cryptography in which he used two methods. One is based on OTP that is One Time Pad is one of the encryption technique which cannot be cracked and the other one is steganography method [2]. Sherif et.al., implemented an encryption algorithm, each plaintext character replaces with four DNA nucleotides or the single

pixel of an image replaced by four DNA nucleotides[11]. Fatma et.al., proposed a new DNA Cryptographic algorithm which used amino acid codes for implementing OTP[12]. Carbon nanotube based information passage and DNA based cryptosystem was proposed by Jie Chen[13]. Pakaj Rajkheja added a DNA cipher layer to the traditional cryptography algorithm that is IDEA. In this algorithm the Key size is incremented to immune the information from cryptanalytic attacks[14]. LAI Xuejia, proposed DNA-Public Key Cryptography which provides digital signature along with encryption[15].

Now, in this paper an effort has been done to adding DNA cipher layer to traditional Feistel Cipher. Since several years there had been numerous modifications taking place in the classical Feistel cipher. It is a symmetric cryptography algorithm[16] which contains 64-bit message as a plaintext. In traditional Feistel cipher the plaintext is divided into two blocks, each block containing 32-bits. The left part L0 contains 32-bits and the right part R0 contains 32-bits. Apply round function with Key value and the right part of plaintext. A new key value is generated for each round. The derived result will be XOR with L0. Now, interchange the R0 and L0 values. Continue the same process for n number of rounds which is described as follows.

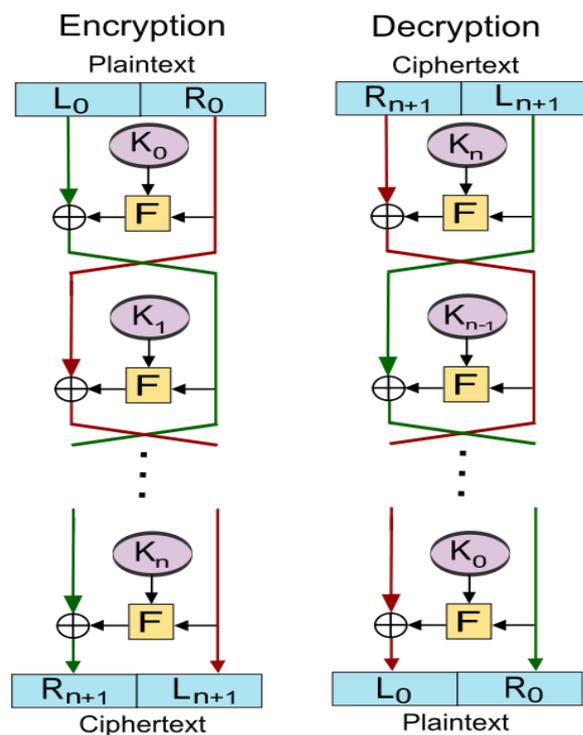


Figure 2: Traditional Feistel Cipher[16]

1. Divide the plaintext block into two equal parts(L0,R0).
2. For each round i=1,2,3,...n  
 $L_{i+1}=R_i$   
 $R_{i+1}=L_i \oplus F(R_i, K_i)$   
 Now the cipher text is (Rn+1, Ln+1).

In the decryption process perform the reverse process that is,

1.  $R_i=L_{i+1}$
2.  $L_i=R_{i+1} \oplus F(L_{i+1},K_i)$

Then  $L_0, R_0$  is the plaintext. Later it is modified by many researchers.

### FEISTEL INSPIRED DNA BASED CRYPTOSYSTEM

DNA cryptography when combined with Feistel cipher gives more security and can be utilized in DNA computers. The algorithm can be implemented by appending the DNA codon table which is called as D-box (Figure 4), to the traditional Feistel Cipher. The D-Box is used as Substitution box where the codons are arranged in  $4^3$  ways. A codon is a unit that consists of three adjacent bases on a DNA molecule and that determines the position of a specific amino acid in a protein molecule during protein synthesis. Out of 64 codons 61 codons are used to form 20 amino acids which can be represented as 20 alphabets (A, B, C...) and 3 codons are used as stop signals. For example, Phenylalanine amino acid (F) can be replaced with either TTT or TTC giving rise to ambiguity when these amino acids are used as an alphabet letter. So, in the present work, the names of amino acids are not mentioned and the 64-codons (Figure 3) are arranged in the D-BOX (8x 8 matrixes) in a random manner. The number of codons formed is 64 permutations of 3 DNA bases from the set of 4 DNA bases. The algorithm is implemented for 64-bit block. If the bits are less, zeroes are padded to make it 64.

		second base in codon				
		T	C	A	G	
T	TTT	Phe	TCT Ser	TAT Tyr	TGT Cys	T
	TTC	Phe	TCC Ser	TAC Tyr	TGC Cys	C
	TTA	Leu	TCA Ser	TAA stop	TGA stop	A
	TTG	Leu	TCG Ser	TAG stop	TGG Trp	G
C	CTT	Leu	CCT Pro	CAT His	CGT Arg	T
	CTC	Leu	CCC Pro	CAC His	CGC Arg	C
	CTA	Leu	CCA Pro	CAA Gln	CGA Arg	A
	CTG	Leu	CCG Pro	CAG Gln	CGG Arg	G
A	ATT	Ile	ACT Thr	AAT Asn	AGT Ser	T
	ATC	Ile	ACC Thr	AAC Asn	AGC Ser	C
	ATA	Ile	ACA Thr	AAA Lys	AGA Arg	A
	ATG	Met	ACG Thr	AAG Lys	AGG Arg	G
G	GTT	Val	GCT Ala	GAT Asp	GGT Gly	T
	GTC	Val	GCC Ala	GAC Asp	GGC Gly	C
	GTA	Val	GCA Ala	GAA Glu	GGA Gly	A
	GTG	Val	GCG Ala	GAG Glu	GGG Gly	G

Figure 3: Structured Codon Table

The proposed cryptosystem follows two functions one is  $E_K(M) = C$  which describes the encryption process and  $D_K(C) = M$  describes the decryption process. The plaintext (M) can be any character but the cipher text  $C \in \{A,C,G,T\}^*$ . First, the sender and receiver must agree for number of rounds to be performed and the random generated key for each round. The random generated Key  $K \in \{0, 1\}^*$ .

Suppose that Alice wants to transmit a message to the Bob, both agree for number of rounds to be performed and then Alice transmits the plaintext message into binary sequence(64 bit data) and split them into 32-bit each. Generate random key for each round and store them securely. Perform XOR operation for the generated key and the right part ( $R_i$ ) of the text. Then, pad 4 zeroes to the result to convert them into octal values. Based upon these values find the codon which is placed on the D-Box. Now the DNA sequence is converted into binary sequence (A-00, C-01, G-10, T-11)(Table1). Remove the last four padded bits and interchange Left and Right parts. After n rounds convert the binary sequence into DNA sequence which is cipher text sent to the Bob. The cipher text is transmitted to Bob through public channel and the stored keys through secure medium. Bob perform the reverse process and get the original plaintext message.

### 4.1 Algorithm for Encrvption

1. Read M, n
2. Convert the Plaintext M of 64-bit block into its equivalent Binary values.
3. Split into  $L_1$  and  $R_1$  each of 32-bits respectively.
4. For  $i=1$  to n
  - Begin
  - a. Generate Key ( $K_i$ ) Randomly.
  - b.  $R_i=R_i \oplus K_i$
  - c. Pad "0000" to  $R_i$
  - d. Split them into 3-bits string each and convert them into its equivalent octal value (We get total 12 strings out of 36 bits).
  - e. By taking two octal numbers we get 6 codons from the D-box(Refer point(c) in each round of encryption process).
  - f. Convert the above DNA strand into its equivalent binary according to DNA table.
  - g. Remove the last "1111" and the value is stored in  $R_i$ .
  - h. Interchange  $L_i, R_i$ .
  - end
5.  $C=L_i||R_i$ .
6. Convert C into DNA bases(00-A,01-C,10-G,11-T), the cipher text.

D-BOX	0	1	2	3	4	5	6	7
0	TTT	TTC	TCT	TTG	TTA	TCC	TCA	TCG
1	TAT	TAC	TGT	TAG	TAA	TGC	TGA	TGG
2	ATT	ATC	ACT	ATG	ATA	ACC	ACA	ACG
3	CAT	CAC	CGT	CAG	CAA	CGC	CGA	CGG
4	CTT	CTC	CCT	CTG	CTA	CCC	CCA	CCG
5	AAT	AAC	AGT	AAG	AAA	AGC	AGA	AGG
6	GTT	GTC	GCT	GTG	GTA	GCC	GCA	GCG
7	GAT	GAC	GGT	GAG	GAA	GGC	GGA	GGG

Figure 4: D-BOX

For e.g., the Codon which is placed in 4<sup>th</sup> row and 6<sup>th</sup> column is CCA.

**4.2 Algorithm for Decryption**

1. Read C, n
2. Convert C into binary from(A-00,C-01,G-10,T-11) and store the same in C.
3. Split the cipher text C into  $L_n$  and  $R_n$  each of 32-bits respectively.
4. For  $i=n$  to 1

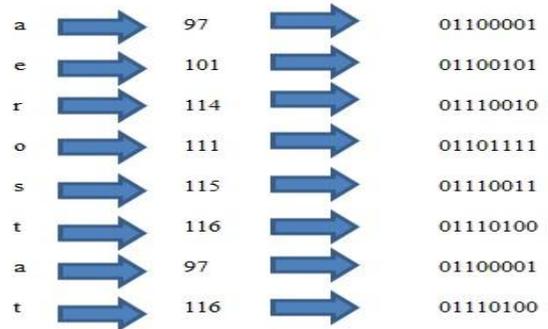
Begin

- i) Pad "1111" to  $L_i$ .
- ii) Convert the binary data into its equivalent DNA bases according to DNA table.
- iii) Split the DNA strand into codons.
- iv) Find the position of each codon from the D-Box.
- v) Convert the positions into its equivalent binary values.
- vi) Remove last "0000" from  $L_i$ .
- vii)  $L_i = L_i \oplus K_i$
- viii) Interchange  $L_i$  &  $R_i$ .

End

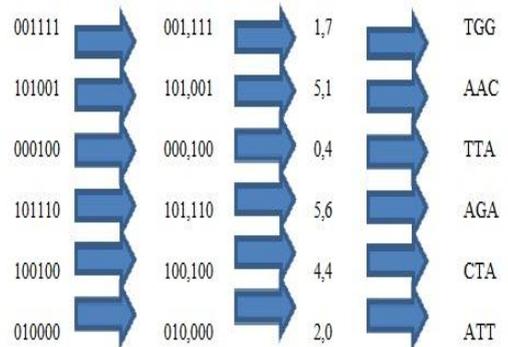
5.  $M = L_i || R_i$
6. Convert the above binary data into its equivalent ASCII Characters which is the plaintext M.
7. Write M.

3. Perform the following step for n number of rounds(let n=4)



**ROUND1**

- a) Generate the key randomly for each round.  
 $K1 = 01001101111001010100110111100101$
- b) Perform XOR operation between  $R1 \oplus K1$  and the result stored in  $R1$ . i.e.,  
 $R1 = 01110011011101000110000101110100$   
 $K1 = 01001101111001010100110100000101$   
 $R1 = 00111110100100010010110001110001$
- c) Divide the result into six bits each. Here total numbers of bits are 32, when we divide 32 with 6 it leaves the remainder 2. To make this remainder 2 as six we pad 0000. Convert three bits of them into equivalent octal values. Put the equivalent Codon form D-Box.
- d) Convert the above DNA strand into equivalent binary from DNA table.



TGGAACTTAAGACTAATT

111010000001111100001000011100001111

- e) Remove the last 1111 from the DNA strand and store the 32-bit result into  $R1$ .  
 $R1 = 11101000000111110000100001110000$
- f) Interchange  $L1$  and  $R1$  and store the result in plaintext M.  
 Now  
 $M = 111010000001111100001000011100000110$   
 $0001011001010111001001101111$ .

**ALGORITHM IMPLEMENTATION**

This section demonstrates the detailed implementation of proposed algorithm for the plaintext "aerostat". The section 5.1 represents the encryption process which converts the given plaintext into cipher text and the section 5.2 represents the decryption process which converts the cipher text into plaintext.

**Encryption Process**

Let the Plaintext M=aerostat

1. Convert the text into its equivalent binary values
2. Split them into 32-bits each and named as  $L1$  and  $R1$ .

**ROUND2**

a) Generate the key randomly for this round.

K2=10000111000011001000011100001100

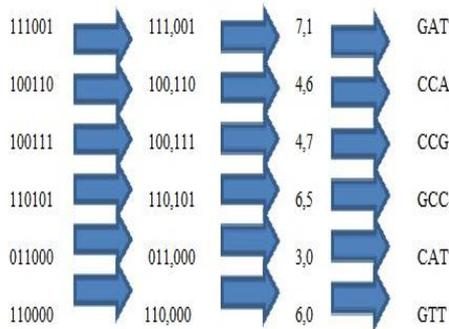
b) Perform XOR operation between R2⊕K2 and the result stored in R2. i.e.,

R2=01100001011001010111001001101111

K2=10000111000011001000011100001100

R2=11100110011010011111010101100011

c) Divide the result into six bits each. Here total numbers of bits are 32, when we divide 32 with 6 it leaves the remainder 2. To make this remainder 2 as six we pad 0000. Convert three bits of them into equivalent octal values. Put the equivalent Codon form D-Box



d) Convert the above DNA strand into equivalent binary from DNA table.

GATCCACCGGCCATGTT  
 100011010100010110100101010011101111

e) Remove the last 1111 from the DNA strand and store the 32-bit result into R2.

R2=10001101010001011010010101001110

f) Interchange L2 and R2 and store the result in plaintext M.

Now  
 M=100011010100010110100101010011101110  
 1000000111110000100001110000.

**ROUND3**

a) Generate the key randomly for this round.

K3=10101010001001011010101000100101

b) Perform XOR operation between R3⊕K3 and the result stored in R3. i.e.,

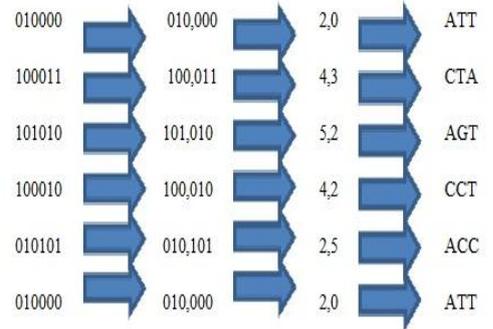
R3=11101000000111110000100001110000

K3=10101010001001011010101000100101

R3=01000010001110101010001001010101

c) Divide the result into six bits each. Here total numbers of bits are 32, when we divide 32 with 6 it leaves the remainder 2. To make this remainder 2 as six we pad 0000.

Convert three bits of them into equivalent octal values. Put the equivalent Codon form D-Box



d) Convert the above DNA strand into equivalent binary from DNA table.

ATTCTGAGTCCTACCATT  
 001111011110001011010111000101001111

e) Remove the last 1111 from the DNA strand and store the 32-bit result into R3.

R3=00111101111000101101011100010100

g) Interchange L3 and R3 and store the result in plaintext M.

Now  
 M=001111011110001011010111000101001000  
 1101010001011010010101001110.

**ROUND4**

a) Generate the key randomly for each round.

K4=01101101100100000110110110010000

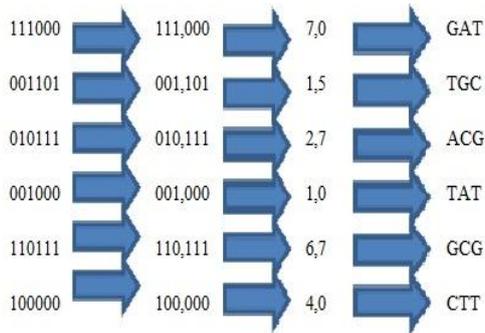
b) Perform XOR operation between R4⊕K4 and the result stored in R4. i.e.,

R4=10001101010001011010010101001110

K4=01101101100100000110110110010000

R4=11100000110101011100100011011110

c) Divide the result into six bits each. Here total numbers of bits are 32, when we divide 32 with 6 it leaves the remainder 2. To make this remainder 2 as six we pad 0000. Convert three bits of them into equivalent octal values. Put the equivalent Codon form D-Box.



d) Convert the above DNA strand into equivalent binary from DNA table.

GATTGCACGTATGCGCTT  
 100011111001000110110011100110011111.

e) Remove the last 1111 from the DNA strand and store the 32-bit result into R4.

R4=10001111100100011011001110011001

f) Interchange L4 and R4 and store the result in plaintext M.

Now

M=100011111001000110110011100110010011  
 1101111000101101011100010100.

4. After n rounds convert the Plaintext M into DNA form and store in to Cipher text C

Now Cipher text  
 C=GATTGCACGTATGCGATTCTGAGTCCTACCA.

### Decryption Process

The cipher text sent by the sender C=GATTGCACGTATGCGATTCTGAGTCCTACCA and convert this into binary form C=1000111110010001101100111001100100111101111000101101011100010100.

1. Perform the step for 4 rounds.

### ROUND4

a) Divide the binary string into two parts, each part contain 32-bit each and name it as L4 and R4.

L4=10001111100100011011001110011001

R4=00111101111000101101011100010100

b) Take the left part L4.

L4=10001111100100011011001110011001.

c) Convert the above binary strand into its equivalent DNA base as per the DNA table. Pad 1111 to this for converting them into equal length codons.

100011111001000110110011100110011111  
 1

GATTGCACGTATGCGCTT

GAT,TGC,ACG,TAT,GCG,CTT

d) Find the location of each codon in the D-Box and convert the values into equivalent binary values and remove last four zeroes to perform XOR operation.

7 0, 1 5, 2 7, 1 0, 6 7, 4 0

111000001101010111001000110111100000

e) Perform XOR operation with K4 and store it into L4. i.e.,  $L4=L4 \oplus K4$ .

L4=11100000110101011100100011011110

K4=01101101100100000110110110010000

L4=10001101010001011010010101001110

f) Interchange L4 and R4. Now the cipher text C=001111011110001011010111000101001000  
 1101010001011010010101001110

### Round3

a) Divide the binary string into two parts, each part contains 32-bit each and name it as L3 and R3.

L3=00111101111000101101011100010100

R3=10001101010001011010010101001110

b) Take the left part L3.

L3=00111101111000101101011100010100

c) Convert the above binary strand into its equivalent DNA base as per the DNA table. Pad 1111 to this for converting them into equal length codons.

001111011110001011010111000101001111

ATT,CTG,AGT,CCT,ACC,ATT

d) Find the location of each codon in the D-Box and convert the values into equivalent binary values and remove last four zeroes to perform XOR operation.

2 0, 4 3, 5 2, 4 2, 2 5, 2 0

0100001000111010100010010101010000

e) Perform XOR operation with K3 and store it into L3. i.e.,  $L3=L3 \oplus K3$ .

L3=01000010001110101010001001010101

K3=10101010001001011010101000100101

L3=11101000000111110000100001110000.

f) Interchange L3 and R3. Now the cipher text C=100011010100010110100101010011101110  
 1000000111110000100001110000.

### Round2

a) Divide the binary string into two parts, each part contains 32-bit each and name it as L2 and R2.

L2=10001101010001011010010101001110

R2=11101000000111110000100001110000

b) Take the left part L2.

L2=10001101010001011010010101001110

c) Convert the above binary strand into its equivalent DNA base as per the DNA table. Pad 1111 to this for converting them into equal length codons.

10001101010001011010010100011101111

GAT, CCA, CCG, GCC, CAT, GTT

- d) Find the location of each codon in the D-Box and convert the values into equivalent binary values and remove last four zeroes to perform XOR operation.

7 1, 4 6, 4 7, 6 5, 3 0, 6 0

111001100110100111110101011000110000

- e) Perform XOR operation with K2 and store it into L3. i.e.,  $L2=L2\oplus K2$ .

$L2=11100110011010011111010101100011$

$K2=10000111000011001000011100001100$

$L2=01100001011001010111001001101111$

- f) Interchange L2 and R2. Now the cipher text  $C=111010000001111100001000011100000110001011001010111001001101111$ .

**Round1**

- a) Divide the binary string into two parts, each part contains 32-bit each and name it as L1 and R1.

$L1=11101000000111110000100001110000$

$R1=01100001011001010111001001101111$

- b) Take the left part L1.

$L1=11101000000111110000100001110000$

- c) Convert the above binary strand into its equivalent DNA base as per the DNA table. Pad 1111 to this for converting them into equal length codons.

111010000001111100001000011100001111

TGG, AAC, TTA, AGA, CTA, ATT

- d) Find the location of each codon in the D-Box and convert the values into equivalent binary values and remove last four zeroes to perform XOR operation.

1 7, 5 1, 0 4, 5 6, 4 4, 2 0

001111101001000100101110100100010000

- e) Perform XOR operation with K1 and store it into L1. i.e.,  $L1=L1\oplus K1$ .

$L1=00111110100100010010111010010001$

$K1=01001101111001010100110100000101$

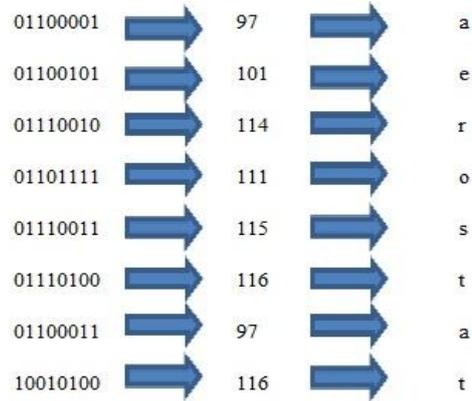
$L1=01110011011101000110001110010100$

Interchange L1 and R1. Now the ciphertext

$C=01100001011001010111001001101110111$

$0011011101000110001110010100$ .

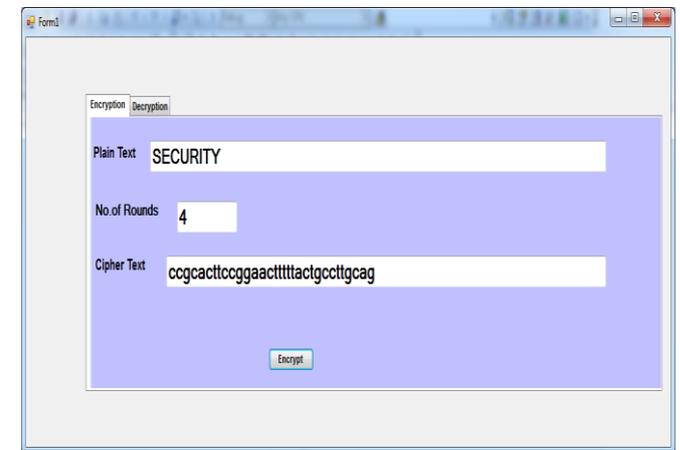
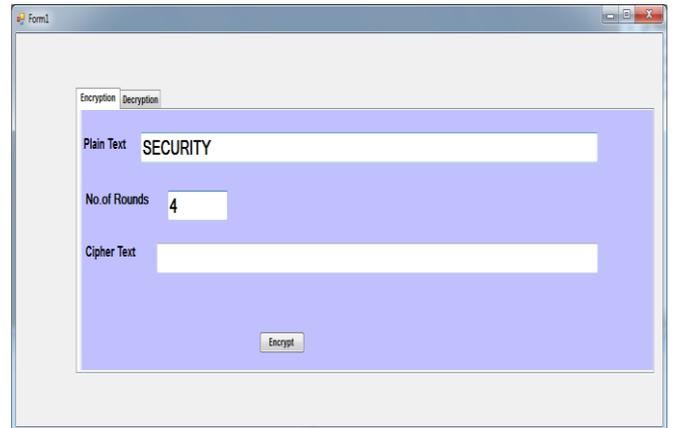
2. Converting the above cipher text into its equivalent ASCII Values, we get the original plaintext "aerostat".



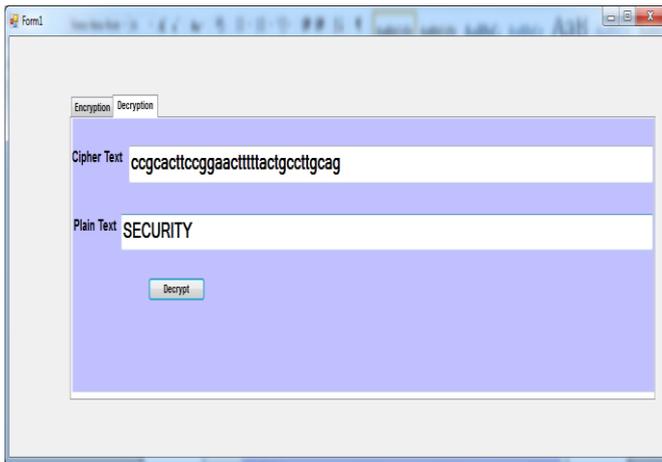
**PERFORMANCE ANALYSIS OF FEISTEL INSPIRED DNA BASED CRYPTOSYSTEM**

The above project is implemented using .NET environment and the simulations are performed by using .net programming on Windows 7 system. The system configuration used is core i3 processor/4 GB RAM and the results are displayed as below.

**Encryption Process**



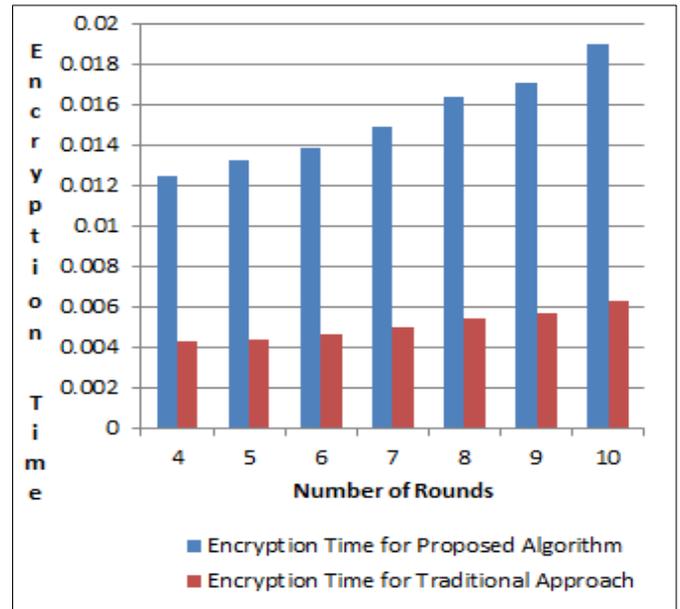
### Decryption Process



The following is the table (Table 2 & Table 3) which shows the time taken to encrypt and decrypt the text in Milliseconds. The performance analysis is compared with the existing algorithms and an effort has been done to propose a better algorithm. The analysis is observed for each round and it is witnessed that if the number of rounds increased then the time taken for encryption and decryption of text also increased.

**Table 2:** Performance analysis in relation to the number of rounds for the encryption of the “security” string.

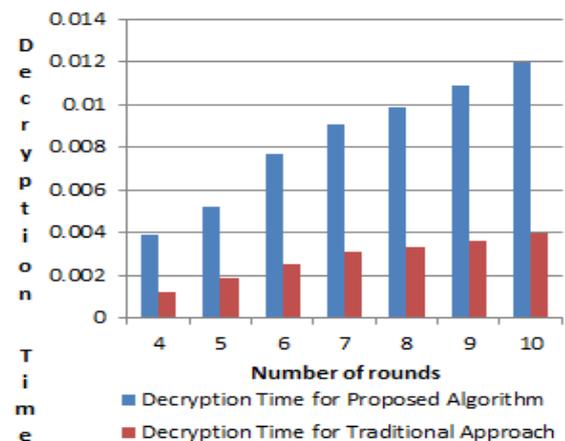
Number of Rounds	Encryption Time for Proposed Algorithm(in ms)	Encryption Time for Traditional Approach(in ms)
4	0.0124734	0.0042995
5	0.0132133	0.0044041
6	0.0138891	0.0046298
7	0.0148977	0.0049995
8	0.0163741	0.0054139
9	0.0170691	0.0057232
10	0.0190141	0.0063381



**Figure 5:** Performance analysis in relation to the number of rounds for the encryption of the “security” string.

No.of Rounds	Decryption Time for Proposed Algorithm	Decryption Time for Traditional Approach
4	0.0038991	0.0011874
5	0.0052451	0.0019017
6	0.0076595	0.0025531
7	0.0090701	0.0030923
8	0.009899	0.0033039
9	0.0108781	0.0036529
10	0.0119875	0.0039621

**Table 3:** Performance analysis in relation to the number of rounds for the decryption of the cipher text “GATTGCACGTATGCGCATTCTGAGTCCTACCA”.



**Figure 6:** Performance analysis in relation to the number of rounds for the decryption of the cipher text “GATTGCACGTATGCGCATTCTGAGTCCTACCA”.

**AVALANCHE EFFECT**

In cryptography, the avalanche effect refers to a desirable property of cryptographic algorithms. The avalanche effect is the outcome result, when an input is modified slightly (for example, flipping a single bit) the output changes consequently (Example half of the output bits flipped). In the case of high-quality block ciphers, such a small change in either the key or the plaintext is too difficult to afford.

The avalanche effect calculated by using the formula

$$\text{Avalanche Effect} = \frac{\text{Number of Flipped bits in cipher text}}{\text{Number of bits in the cipher text}} \times 100\%$$

The following is the calculation of avalanche effect to various cryptographic techniques and the proposed work. Let us take

Plaintext1:aerostat

Plaintext2:aerrstat

Key: eggplant

**1. Caesar Cipher**

Ciphertext1:01100110011010100111011101110100  
 01111000011110010110011001111001  
 Ciphertext2:01100110011010100111011101110111  
 01111000011110010110011001111001

**2. Playfair Technique**

Ciphertext1:01100111011000010111001101110001  
 01110001011000110111010001101000  
 Ciphertext2:0110011101100001011100101111001  
 01110001011000110111010001101000

**3. Vigenere Cipher**

Ciphertext1:01100101011010110111100001100100  
 01100100011101000110111001101101  
 Ciphertext2:01100101011010110111100001100111  
 01100100011101000110111001101101

**4. Blowfish**

The number of bits flipped in the cipher text in this algorithm is approximately 19 bits and the average percentage of avalanche effect is 29.68[17].

**5. Feistel Inspired DNA based crypto system**

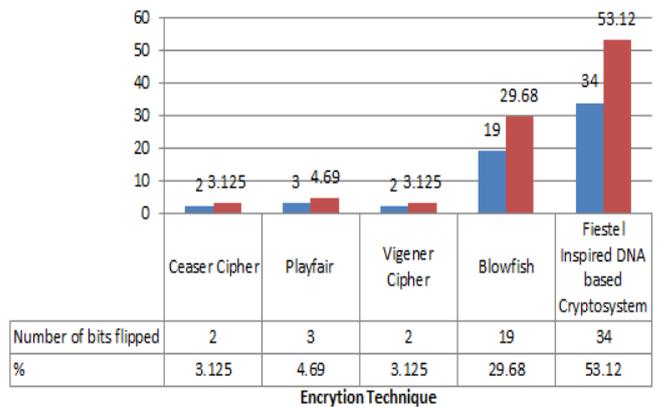
Ciphertext1:  
 100100001111101100111011110011110000110011  
 100001111100110111100  
 Ciphertext2:00001001101000101000100110010010  
 1001011000010010010011000010001

The following table (Table 4) shows the comparison results of various crypto algorithms and the proposed algorithm.

**Table 4:** Comparison results of Avalanche Effect for strings “aerostat” and “arrstat”

Name of the Algorithm	Number of bits flipped	%
Caesar Cipher	2	3.125
Playfair	3	4.69
Vigener Cipher	2	3.125
Blowfish	19	29.68
Feistel Inspired DNA based Cryptosystem	34	53.12

**Comparison of Avalanche Effect**



**Figure 7:** Comparison of the results of the avalanche effect for strings “aerostat” and “arrstat”

**CONCLUSIONS**

In this paper, we developed a symmetric block cipher algorithm as same as Feistel cipher adding a layer of DNA codons to it. Take a plaintext of 64-bit length which is further divided into two parts and perform XOR operation with randomly generated key with the right part and retrieve the codon values from the D-Box and then convert the DNA strand into its equivalent binary values, which is the cipher text. At the receiver end, the cipher text is divided into two blocks. Consider the left part and convert it into equivalent DNA bases and split them into Codons. Find the position of codons in the D-Box and convert the positions into equivalent binary values, perform the left bit XOR with the key value. It has been observed that, in the block of 64-bit plaintext more than half of the bits have changed in the cipher text in each round. Hence, it makes highly impossible for the intruders to identify the plain text. Further, comparison of performance analysis is done for the existing and proposed algorithms. Avalanche effect is also studied and recorded. Despite the fact that DNA is termed as a medium for ultra- compact information storage and the future appears promising to fascinate the researchers in the area of research, there are still some of the issues like-requirements of quantum attacks

and bio molecular labs that need to be taken into consideration.

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