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Application of DNA Coding, the Lorenz Differential Equations and a Variation of the Logistic Map in a Multi-Stage Cryptosystem

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Abstract: The need for information security has become urgent due to the constantly changing nature of the Internet and wireless communications, as well as the daily generation of enormous volumes of multimedia. In this paper, a 3-stage image cryptosystem is developed and proposed. A tan variation of the logistic map is utilized to carry out deoxyribonucleic acid (DNA) encoding in the first stage. For the second encryption stage, the numerical solution of the Lorenz differential equations and a linear descent algorithm are jointly employed to build a robust S-box. The logistic map in its original form is utilized in the third stage. Diffusion is guaranteed through the first and third encryption stages, while confusion is guaranteed through the application of the S-box in the second encryption stage. Carrying out both confusion- and diffusion-inducing stages results in encrypted images that are completely asymmetric to their original (plain) counterparts. An extensive numerical analysis is carried out and discussed, showcasing the robustness and efficacy of the proposed algorithm in terms of resistance to visual, statistical, entropy, differential, known plaintext and brute-force attacks. Average values for the computed metrics are: Information entropy of 7.99, MSE of 9704, PSNR of 8.3 dB, MAE of 80.8, NPCR of 99.6 and UACI of 33. The proposed algorithm is shown to exhibit low computational complexity, encrypting images at an average rate of 1.015 Mbps. Moreover, it possesses a large key space of 2^{372} , and is demonstrated to successfully pass all the tests of the NIST SP 800 suite. In order to demonstrate the superior performance of the proposed algorithm, a comparison with competing image encryption schemes from the literature is also provided.

Keywords: chaotic maps; cryptography; image encryption; logistic map; Lorenz differential equations; S-box

1. Introduction

Security has become a matter of utmost significance because of the enormous advancements and complexity observed in today's wireless communication networks and big data applications [1–3]. Thus, utilizing cryptography [4,5], steganography [6,7], watermarking [8] and their combinations [9,10] to protect data has become essential for ensuring the secure functioning and use of millions of online services. For many years, well-established cryptosystems were put in use, for virtually every type of data. Those included the Data Encryption System (DES) [11], its variant, the Triple DES [12], as well as the Advanced

Encryption Standard (AES) [13,14]. However, it quickly became clear that not all cryptosystems are well-suited for application on multimedia such as 2D and 3D images and videos. This is because images and videos have vast amounts of data, redundancy, as well as strong cross-correlation among adjacent pixels. To this end, the literature on developing secure, robust image cryptosystems has been on the rise in recent years. Algorithms that carry out image encryption are based on mathematical operations that are derived from, or related to, chaos theory [15–18], electric circuits [19,20], DNA encoding [21–25], and cellular automata [26], to name a few. The following paragraphs highlight the significance of DNA cryptography and chaos theory in security applications, as well as their use in cutting-edge image encryption algorithms. Substitution boxes (S-boxes), a powerful component for introducing confusion into a cryptosystem, are discussed next.

DNA cryptography makes use of both biological and computational properties to offer more confidentiality over classical cryptographic algorithms while encrypting data [24]. Traditional cryptosystems often only provide one layer of protection, and it is possible that their secrecy is compromised as the underlying computational techniques are made public. On the contrary, DNA cryptography utilizes the self-assembling characteristics of DNA bases in combination with a cryptographic approach to provide many security measures that enhance the amount of data confidentiality [25]. For example, the authors of [22] convert ciphertext to a genomic form using amino acid tables. The tables' protein sequence composition add to the ciphertext's level of ambiguity. In [23], the authors propose a DNA encoding algorithm that is built on a unique string matrix data structure producing distinctive DNA sequences. They employ these sequences to encode plaintext as DNA sequences. While DNA cryptography has garnered the interest of scientists and engineers in recent years, it definitely has not gained as much attention as that dedicated to chaotic and dynamical systems use in image cryptosystems.

The intrinsic properties of chaotic functions as a random phenomenon in nonlinear systems are favourable for cryptography [27]. Specifically, their sheer sensitivity to initial conditions, control parameters, periodicity, pseudo-randomness, and ergodicity. These properties are incorporated into the design of image encryption algorithms. Broadly, such schemes are divided into two classes: (a) One-dimensional (1D) and (b) Multi-dimensional (MD). The utilization of 1D chaotic maps provides for simpler and more efficient software and hardware implementations. However, this also translates into less desirable characteristics, in terms of shorter chaotic periods, non-uniform distribution of their chaotic output, as well as a greater susceptibility to cryptanalysis. On the contrary, the utilization of MD chaotic maps in image encryption algorithms provides stronger security levels at the expense of increased complexity and, consequently, longer running times for software and hardware implementations. Extensive literature exists on the use of 1D and MD chaotic functions in image cryptosystems. The authors of [4], for instance, propose an image encryption algorithm that is based on a combination of encryption keys designed using the Arnold cat map, the 2D logistic sine map, the linear congruential generator, the Bernoulli map and the tent map. In a similar manner, the authors of [17] also employ multiple chaotic maps, however, in their implementation, they aim at reaching a minimum number of encryption rounds while maintaining a high degree of security and robustness. In [28], the authors employ a finite field aiming to generalise the logistic map and search for an auto morphic mapping between two logistic maps in order to compute parameters over the finite field Z_N . Shannon's ideas are fully put into use in the work of [18], where an LA-semi group is applied for confusion, and a chaotic continuous system is adopted for diffusion. The authors of [29] present an interesting work that employs a zigzag transform in a conjoint manner with a dynamic arrangement that alternates in a bidirectional crossover approach to image encryption. In their proposed cryptosystem, both the logistic map and a hyperchaotic Chen dynamical system are utilized. In actuality, this paragraph merely touches upon the use of chaos theory in color image cryptosystems. Recent writing on the subject is voluminous. The following paragraph focuses on a distinct but vital component of numerous image encryption algorithms: substitution boxes.

A crucial part of contemporary block cryptosystems is an S-box. It makes it easier to convert any given plaintext into ciphertext. The confusion property is provided by the straightforward addition of an S-box to a cryptosystem, which results in a non-linear translation between the input and output data [30]. An S-box provides better security the more uncertainty there is. For many block encryption techniques, the level of security offered by using one or more S-boxes closely correlates with how resistant they are against assaults. While such algorithms may have numerous stages, an S-box is typically the only non-linear stage that improves the security of sensitive data [31]. To be acceptable for real-time data encryption, the design of an S-box should be efficient and low in complexity. Recent literature provides multiple instances of design and utilization of S-boxes in image encryption algorithms. For example, the authors of [15] proposed an S-box utilising a third-order nonlinear digital filter. Its non-linearity was enhanced using a novel optimisation approach. In [31], the authors developed an optimization algorithm for a chaos-based entropy source to generate an S-box. Multiple stage image encryption schemes where an S-box is a core stage are rather popular among scientists and engineers, since such a combination satisfies Shannon's ideas of confusion and diffusion. Furthermore, employing more than one encryption stage provides security against known plaintext attacks [32]. The authors of [16] propose one such example of a 3-stage image encryption algorithm, where in one of the stages they utilize the S-box proposed in [33]. This S-box is based on a modular approach and is thus, highly nonlinear. In the other two stages, a Lucas sequence and a Sine logistic map are made use of to generate encryption keys. Another 3-stage algorithm is proposed in [34], where an S-box is also utilized as a core stage, sandwiched between the application of two encryption keys. The first key is a Mersenne Twister based PRNG, while the second key is a tan variation of the logistic map. In [35], the authors follow a similar approach, generating a PRNG-based S-box using Wolfram Mathematica[®], and utilizing the Rossler system and the Recaman's sequence, for each of the encryption keys.

Combining DNA cryptography with chaotic functions and S-boxes in image encryption algorithms gained the attention of many researchers in the field as an attempt to achieve better performance [36], either in terms of improved security or lower computational complexity and thus ever decreasing encryption and decryption times. The work presented in [37] introduced a cryptosystem for color images based on a combination of chaotic maps and DNA sequences. The reported theoretical and statistical analyses reflected the robustness of combining DNA with chaotic maps against statistical and brute force attacks [37]. In [38], a 2D Henon-Sine map and DNA coding was proposed. Exclusive-OR (XOR) and DNA random coding encryption operations were synthesized using an S-box for image diffusion, while image scrambling was carried out through swap operations on the pixels of the image. The work presented in [38] highlighted the merits of generalizing DNA encryption with S-box substitution in image encryption techniques. In [21], the authors proposed an image cryptosystem that utilizes parallel compressive sensing, chaos theory and DNA encoding. The authors of [39] provide an interesting work for grayscale image encryption that combines a 4D chaotic system with DNA encoding, the hash function SHA-2 and the random movement of a chess piece (Castle), through an iterative process. The authors of [40] not only combine the use of DNA encoding, SHA-512 hashing and multiple hyperchaotic maps, but also utilize a novel variation of a chaotic map, a logistic-tan map, as well as a pixel-shifting algorithm that is based on the Zaslavskii map.

The contributions of this paper are as follows:

- We propose a 3-stage image encryption scheme that makes use of Shannon's ideas of confusion and diffusion. In the first stage, DNA coding is employed, providing diffusion at the bit level. In the second stage, an S-box based on the numerical solution of the Lorenz differential equations and a linear descent algorithm is developed and used for confusion at the pixel level. In the third stage, the logistic map is utilized to produce an encryption key, providing diffusion at the bit level. The concatenation of these three stages guarantees output encrypted images to be completely asymmetric to their original (plain) counterparts.

- We propose an efficient and fast encryption scheme, with images of dimensions 128×128 encrypted in only 0.377145 s, achieving an average encryption rate of 1.015 Mbps.
- We propose a multi-stage image encryption scheme. Using more than one encryption stage provides security against known plaintext attacks.
- We propose an image encryption scheme that possesses a large key space of 2^{372} , and is effectively resistant to brute force attacks.
- We utilize both conventional (information entropy, pixel cross-correlation, MSE, PSNR, MAE, NPCR, UACI and NIST SP 800 suite) and unconventional performance evaluation metrics such as the Fourier transform and advanced bit dependency metrics to gauge the security and robustness of the proposed cryptosystem.

This paper is organized as follows. Section 2 outlines the mathematical background and describes the proposed image cryptosystem. Section 3 presents the computed numerical results and carries out a comparative analysis with counterpart cryptosystems from the literature. Finally, Section 4 draws the conclusions and suggests possible future research directions.

2. Proposed Image Cryptosystem

This section describes the proposed 3-stage image cryptosystem. It starts off by describing each stage separately, then how they complement each other forming one cryptosystem. Finally, the decryption process is described.

2.1. DNA Encoding Based on a Tan Variation of the Logistic Map

As mentioned before, key embedding is considered as a critically important component in image encryption. This is due to the role it plays in data diffusion. Moreover, in many of the more recent image encryption techniques, key embedding is performed two times, separated by a confusion step. As the proposed technique in this work follows the same model, the two stages of key embedding are performed differently. The third stage in this work utilizes bit-wise key embedding, as discussed in Section 2.3. For the first stage, which is the scope of this section, DNA encoding is utilized as a means for embedding a seed-based key into the raw data of an input image (to be encrypted), which is demonstrated in Section 2.1.2. Prior to that, the method of the key generation is discussed as well in Section 2.1.1.

2.1.1. Tan Variation of the Logistic Map

The logistic map is a recurrence relation, which is also a second-degree polynomial mapping that models how complex, chaotic behavior can be simulated by extremely simple non-linear dynamic equations. In recorded research history, the first mention of an equation which behaves accordingly dates back to 1976 [41]. The general mathematical definition of a logistic map is (Equation (3) in [41]):

$$X_{t+1} = \alpha \times X_t \times (1 - X_t) \quad (1)$$

Equation (1) shows a recurrent function where the value of X_{t+1} is directly calculated using the value of X_t , and the value of a scale factor α . Adopted in this work is a variation of the logistic map function, which utilizes the application of the tan function, beside introducing the use of a seed (as the value of X_0). The tan variation is equated as:

$$X_{t+1} = \alpha \times \tan(X_t) \times (1 - X_t) \quad (2)$$

Regarding the ranges for both keys (α and X_0), both are to be chosen such that for every X_t in the sequence, the range should remain $]0, 1[$. Starting by the lower limit, the 0 is avoided as $\tan(0) = 0$, which will result in, once a 0 appears in the sequence, all remaining values in the sequence will become 0. When it comes to the upper limit, given that the function in a tan variant, $\frac{\pi}{2} \approx 1.5708$ is to be avoided as $\tan(\frac{\pi}{2}) = \infty$. Moreover, plotting the tan curve, starting from 1 till 1.5708, the curve slope starts to be steep (in other words,

with a tangent slope greater than 1). The main effect of this is that the rate of increase in \tan values, given small steps in input, starts to be harder to limit (as it approaches infinity). As per that, initiating the value of X_0 , the range is considered optimal (for this scenario) as $]0, 1[$. According to the X_0 range, regarding the scale factor α , for a chaotic behaviour, the range is $[3.3, 3.6]$. This is confirmed visually through plotting the bifurcation diagram and Lyapunov exponent, in Figures 1 and 2, respectively.

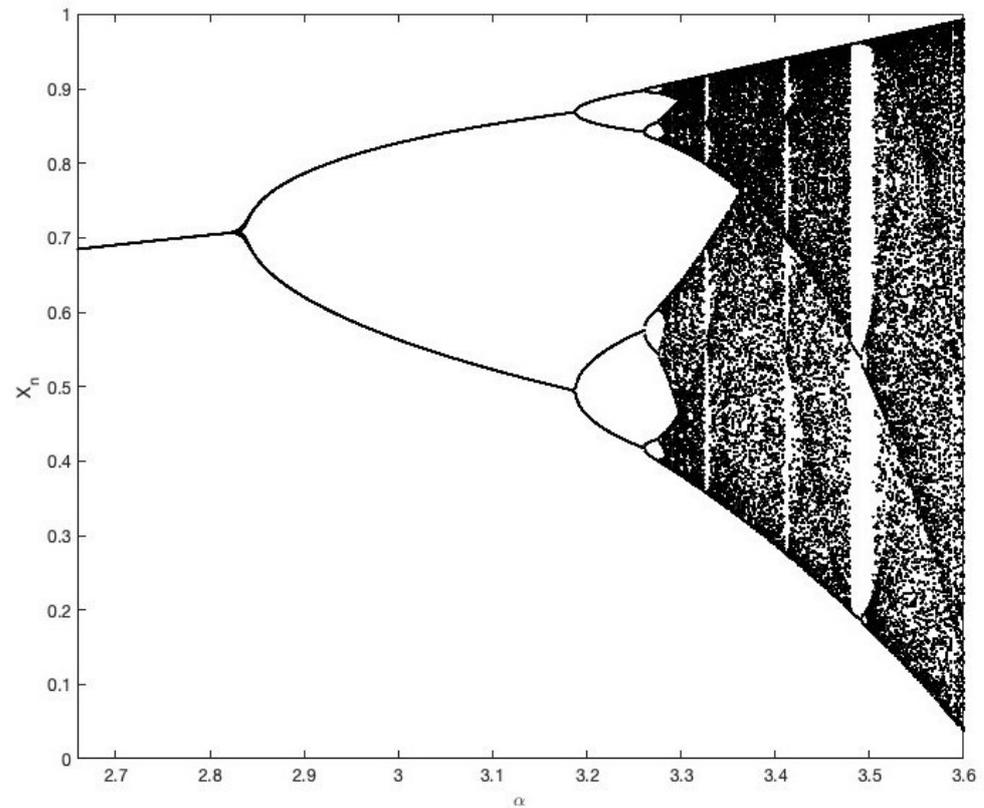


Figure 1. Bifurcation diagram of the proposed tan variation of the logistic map.

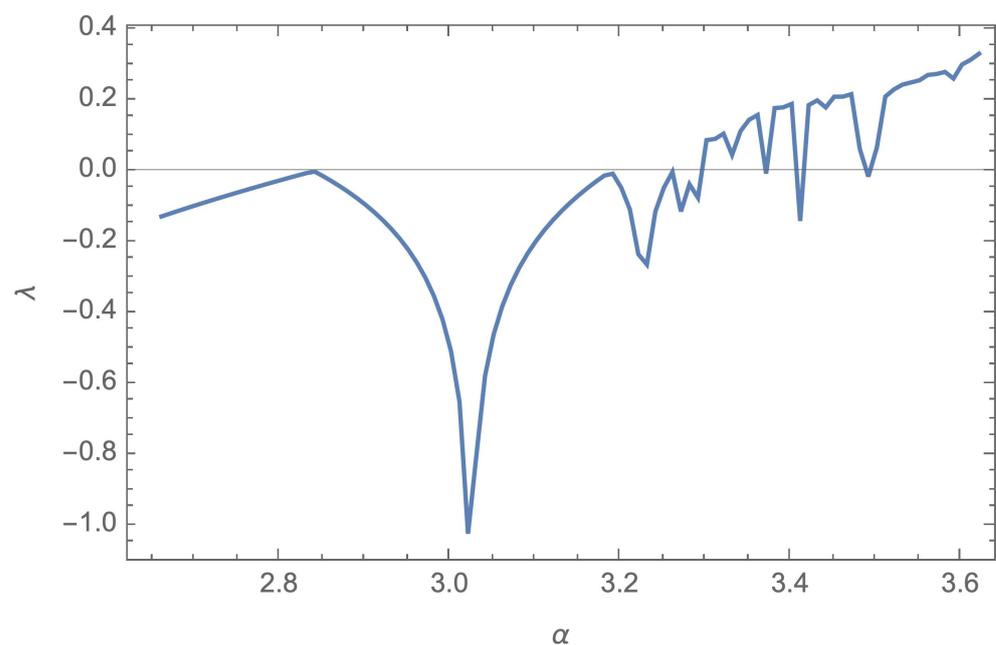


Figure 2. Lyapunov exponent diagram of the proposed tan variation of the logistic map.

2.1.2. DNA Encoding

Numerous studies have identified the DNA computer characteristics of vast parallelism, enormous storage, and ultra-low power consumption. DNA cryptography is a novel cryptography discipline that evolved from research on DNA computing, in which DNA is employed as an information carrier and contemporary biological technology is used as an implementation tool. As per that, in order to achieve better encryption, the DNA domain is the domain variation on which the image will be manipulated during this stage. The process of DNA encoding of an image is a modification of the bit stream representation in which each pair of successive bits is joined. In such a situation, the processes conducted would seem to be bit-level operations, but the attacker would not be able to readily trace them back using bit-level analyses.

DNA coding has three stages. First, it creates a DNA sequence from bit stream sequences by merging every 2 bits into 1 DNA base, as per the following relation:

$$\{(00 \rightarrow A), (01 \rightarrow T), (10 \rightarrow C), (11 \rightarrow G)\}. \quad (3)$$

Note that other relations also exist [21]. Table 1 shows eight possible permutations of complementary DNA encoding and decoding.

Second, applying a DNA level procedure. As reversibility is required, in this work, addition and subtraction are the inverse operations of choice, which are performed according to Table 2. In the last phase, a DNA sequence is converted into a bit stream.

The stated approach requires a DNA sequence of the same length as the image-generated DNA sequence in order to finish the encryption process. This work generates a PRNG seed-based bit stream, which is subsequently transformed into a DNA sequence. Given the two sequences (image and seed), the aforementioned processes may be carried out, with addition being used for encryption and subtraction for decryption. Figure 3 illustrates an example DNA sequence.

Table 1. Eight rules of complementary DNA encoding and decoding).

Rule	1	2	3	4	5	6	7	8
A	00	00	11	10	01	10	01	11
T	11	11	00	01	10	01	10	00
G	10	01	10	11	00	00	11	01
C	01	10	01	00	11	11	00	10

Table 2. Simple arithmetic operations on DNA bases (here, {Addition, Subtraction}).

	A	T	C	G
A	{A,A}	{T,G}	{C,C}	{G,T}
T	{T,T}	{C,A}	{G,G}	{A,C}
C	{C,C}	{G,T}	{A,A}	{T,G}
G	{G,G}	{A,C}	{T,T}	{C,A}

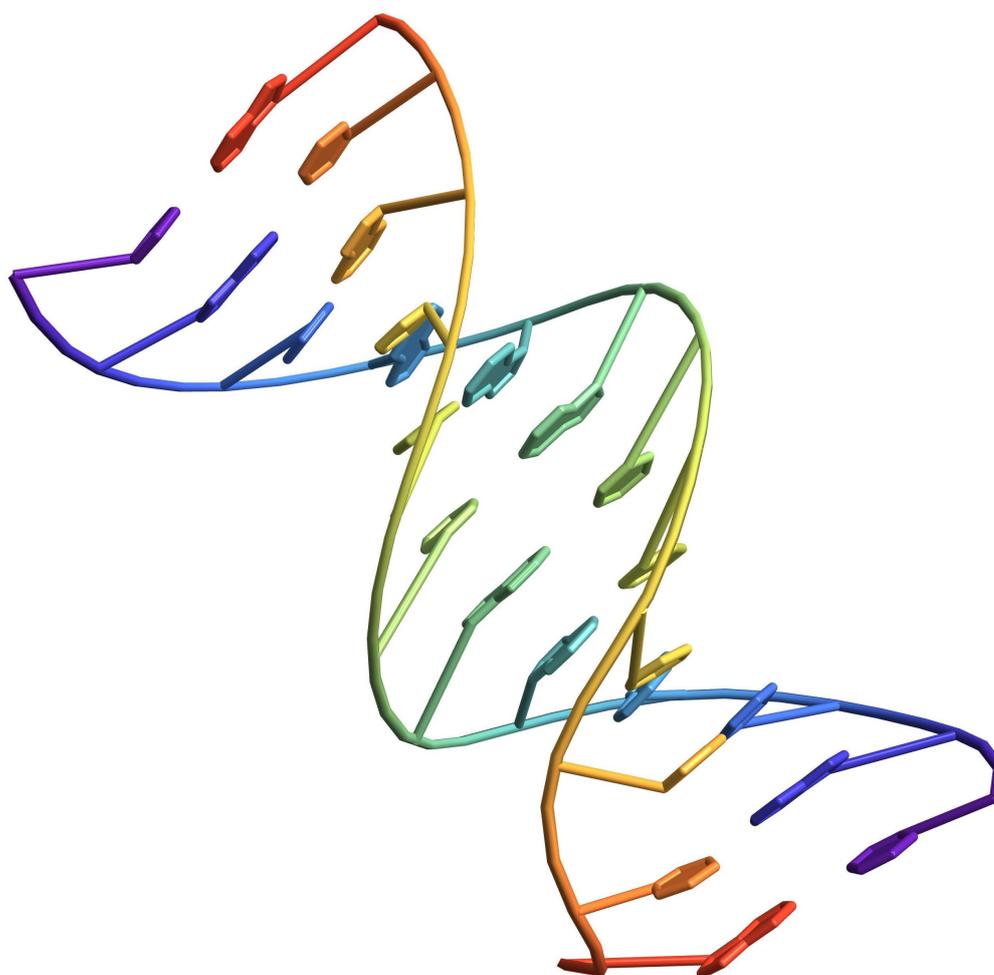


Figure 3. 3D plot of a DNA sequence.

2.2. Lorenz S-Box

An S-box is a vital non-linear component of any image encryption system. This is because it takes advantage of Shannon's property of confusion [42]. For the formation of the S-box, a PRNG building approach is applied. In other words, a random sequence is initially produced (using a seed since reconstruction is required), and then employed as a sequential selection factor until the S-box is completely constructed. For the production of PRNG sequences, the Lorenz system is used. The Lorenz system is a mathematical model for atmospheric convection. It was developed in 1963 by Edward Lorenz [43]. The mathematical model for this system is defined by the following three partial differential equations (where σ , ρ , and β are provided as seeds):

$$\frac{dx}{dt} = \sigma(y - x), \quad (4)$$

$$\frac{dy}{dt} = x(\rho - z) - y, \quad (5)$$

$$\frac{dz}{dt} = xy - \beta z. \quad (6)$$

The Lorenz system is numerically solved, resulting in a 3D geometry, as shown in Figure 4. Next, the x , y , and z coordinates of each point in the solution are flattened into a single 1D array, which is then turned into a bitstream using the median of the list as a threshold. Finally, each 8-bits are used to form a decimal number $\phi \in [0, 255]$. For example,

using the values $\sigma = 10$, $\beta = 8/3$, and $\rho = 28$, we obtain the first eight bits in the generated bitstream as $\{1, 1, 1, 0, 0, 1, 0, 0\}$, which we convert into a decimal value of 228, as shown in the first entry of the S-box in Table 3.

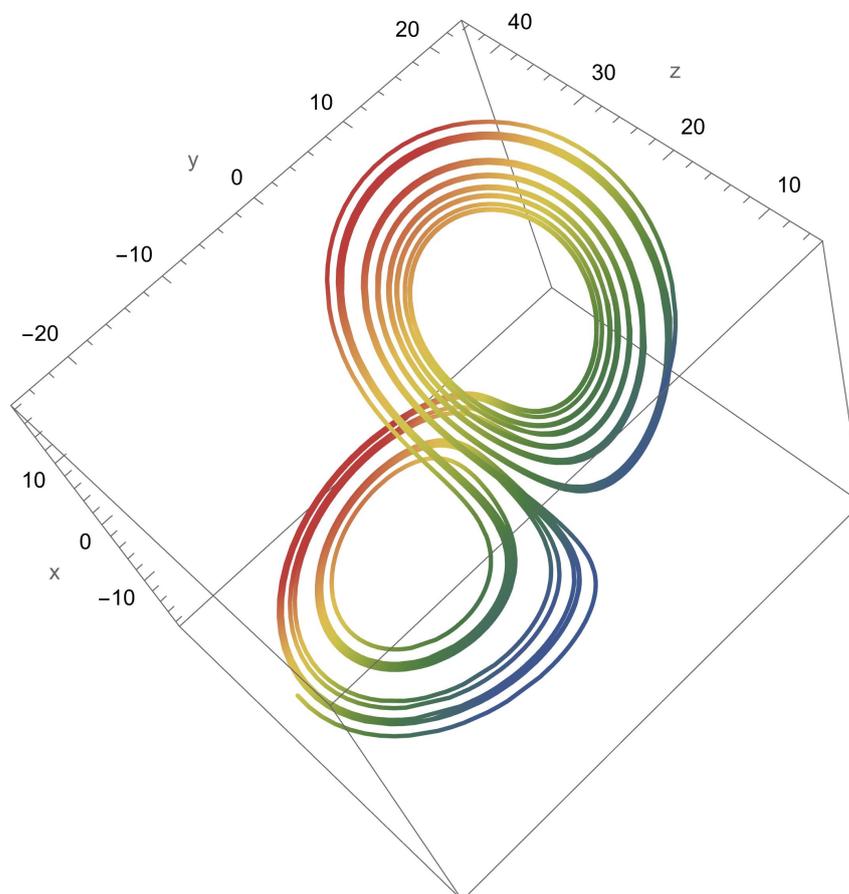


Figure 4. 3D plot of the numerical solution of the Lorenz system of partial differential equations.

Given a series S of 256 numbers (unsorted and containing repeated numbers), the S-box is constructed in constant time using the procedures of Algorithm 1.

Algorithm 1 Generate S-box given S and L

1. $L = [0 - 255]$
 2. $n = 0$
 3. $i = S_n \% \text{Length}(L)$
 4. append $L[i]$ to S-box
 5. delete $L[i]$ from L
 6. $n = n + 1$
 7. if $n \leq 256$: GoTo(2)
-

As an example, given seed values of $\sigma = 10$, $\rho = 28$, and $\beta = 8/3$, the generated S-box is as shown in Table 3.

Table 3. Lorenz system based S-box using the values $\sigma = 10$, $\beta = 8/3$, and $\rho = 28$.

228	251	73	62	161	206	233	5	76	37	165	78	41	154	212	3
201	81	39	158	218	40	159	84	69	163	86	42	167	94	46	170
89	7	31	32	49	189	95	14	186	97	51	181	250	202	43	152
54	191	249	50	196	134	52	188	103	85	209	105	44	79	112	58
254	109	68	219	111	227	204	117	92	220	121	235	210	174	55	102
182	122	20	179	100	229	119	96	242	123	57	231	60	87	234	127
59	238	171	126	2	156	66	246	101	63	248	135	64	255	211	147
19	215	116	9	140	70	15	131	125	21	142	185	98	237	74	27
160	77	34	222	82	80	199	88	203	178	13	35	177	99	75	180
38	36	29	113	114	184	108	128	200	53	124	141	166	143	198	129
150	22	173	67	33	213	172	223	149	197	214	195	194	240	130	253
148	151	232	176	137	244	252	132	4	6	56	18	12	192	47	30
153	93	104	205	8	65	168	216	90	175	190	72	187	17	136	239
25	118	164	139	183	83	221	224	241	245	236	247	10	115	208	207
71	26	225	133	45	107	1	138	23	24	144	146	243	162	16	157
193	120	48	217	28	155	0	106	61	11	145	226	110	169	230	91

2.3. The Logistic Map

The logistic map, as a 1D chaotic function, is a nonlinear deterministic system. It contains a variety of attributes, including strong sensitivity to initial conditions (used for seed effect modeling), and deterministic behavior (used for sequence reconstruction). Chaotic sequences generated by dynamical functions are pseudo-random sequences with very complicated and difficult to anticipate features. Therefore, chaotic systems may increase an encryption system's security. In the third stage of the proposed encryption approach, we make use of the logistic map, expressed mathematically as in (1), where two variables govern the function's behaviour. The first variable is the scale factor, $\alpha \in [3.7, 4]$, representing the rate of change of the returned data. The second variable is the initial value $X_0 \in [0, 1]$, determining the starting point of the graph.

2.4. Proposed Cryptosystem

The encryption procedure may be broken down into three (distinct) subroutines since it uses a 3-stage encryption approach. Each of the subroutines receives its own individual set of parameters (an image and a seed) and produces mostly only one result (encrypted image). In addition, each one is equipped with a reverse subroutine, which is essential for the decryption process. In Sections 2.4.1 and 2.4.2, the sequential procedures of both encryption and decryption processes are demonstrated, respectively.

2.4.1. Encryption Procedure

The encryption procedure, as mentioned, is divided into three stages. Each stage represents the interaction between the input image, and the seed-based key involved, in order to result in the final output, which is the encrypted image. Moreover, each stage is performed over multiple steps. For elaboration, this procedure can be enumerated in the following steps:

1. Stage 1: DNA Encoding Based on a Tan Logistic Map.

- (a) As a start, the input image, I of dimensions $M \times N$, is converted into a 1D bit-stream to produce I_{1D} , alongside calculating the length of this bit-stream:

$$BitStreamLength = M \times N \times 24 \quad (7)$$

- (b) Given a seed for the Tan variation of the logistic map, which consists of values for both X_0 and α , a sequence of numbers is generated (Seq_{tan}) using (2), which is equal in length to the bit-stream length.
- (c) Forming the Key_{tan} , the median (mid) of the sequence is generated, and each element is compared against this median value, converting the sequence into a bit-stream (comparing with the median value assures producing an equal number of 0's and 1's in the bit-stream, without sacrificing the PRNG aspect.) This step is performed as per the following equations:

$$C(n) = \begin{cases} 1, & n > mid \\ 0, & otherwise \end{cases} \quad (8)$$

$$Key_{tan} = \bigcup_{i=0}^{BitStreamLength} C(Seq_{tan}) \quad (9)$$

- (d) Both bit-streams I_{1D} and Key_{tan} are converted into DNA-streams using (3), generating $I_{1D,DNA}$ and Key_{tanDNA} , respectively.
- (e) Both DNA-streams are then added, using Table 2, generating $I_{1D,tanDNA}$, then converted back into a bit-stream, as per the following equation:

$$I_{1D,tanDNA} = AddDNA(I_{1D,DNA}, Key_{tanDNA}) \quad (10)$$

2. Stage 2: S-box Application.

- (a) Given a seed for the S-box, which consists of the Lorenz system inputs, σ , β , and ρ , the Lorenz system is calculated (as in Figure 4), then converted into a 1D sequence using (11) (as in Figure 5) of length 2048 (to form 256 numbers, 8 bits each).

$$Lorenz1D = \{P_1, P_2, \dots, P_M\} \rightarrow \{x_1, y_1, z_1, x_2, y_2, z_2, \dots, x_M, y_M, z_M\}. \quad (11)$$

- (b) As performed with the tan sequence, the median value will be utilized in combination with (8) in order to convert the Lorenz 1D sequence into a bit-stream. Then, each 8 bits are further converted to decimal, resulting in a list $S \in [0, 255]$, and $|S| = 256$.
- (c) List S is provided as input to Algorithm 1, producing the S-box. Finally, the S-box is applied as:

$$I_{1D,tanDNA,S-box} = S - box(I_{1D,tanDNA}) \quad (12)$$

3. Stage 3: Logistic Map Encoding.

- (a) Given a seed for the logistic map (Section 2.3), which consists of values for both X_0 and α , a sequence of numbers is generated (Seq_{log}) using (1), which is equal in length to the bit-stream length of the image ($BitStreamLength$).
- (b) As performed with the tan sequence (Stage 1, step c), the median value will be utilized in combination with (8) in order to convert the sequence into a bit-stream, generating Key_{log} .
- (c) The produced bit-stream, alongside the one produced the by the end of Stage 2 ($I_{1D,tanDNA,S-box}$), an XOR operation is performed as follows:

$$I_{1D,tanDNA,S-box,log} = I_{1D,tanDNA,S-box} \oplus Key_{log} \quad (13)$$

After performing the 3 stages, reshaping $I_{1D,tanDNA,S-box,log}$ back into a 2D image (of dimensions $M \times N$) results in the encrypted image I' . Figure 6 demonstrates a flowchart for the encryption procedure.

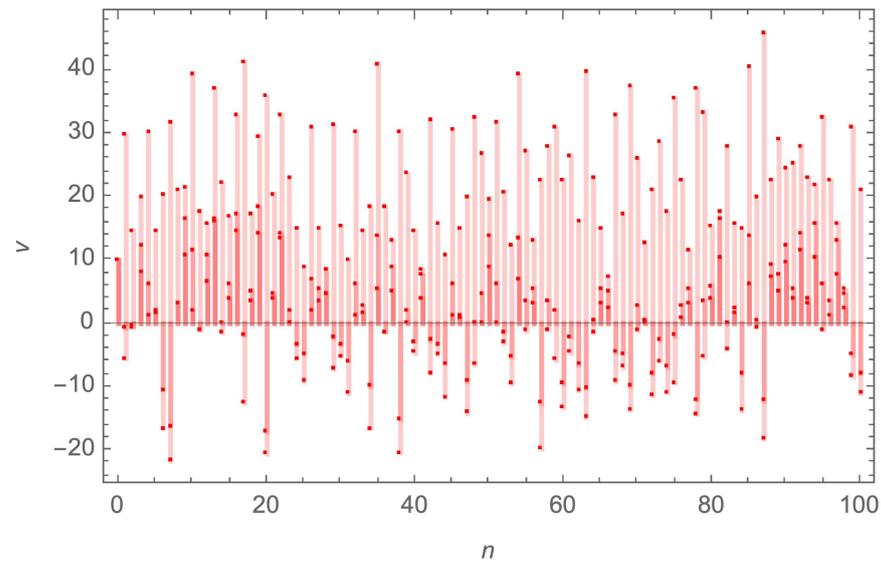


Figure 5. The first 100 points from the 1D array obtained from the 3D coordinates of the Lorenz system solution for the values $\sigma = 10, \beta = 8/3$ and $\rho = 28$.

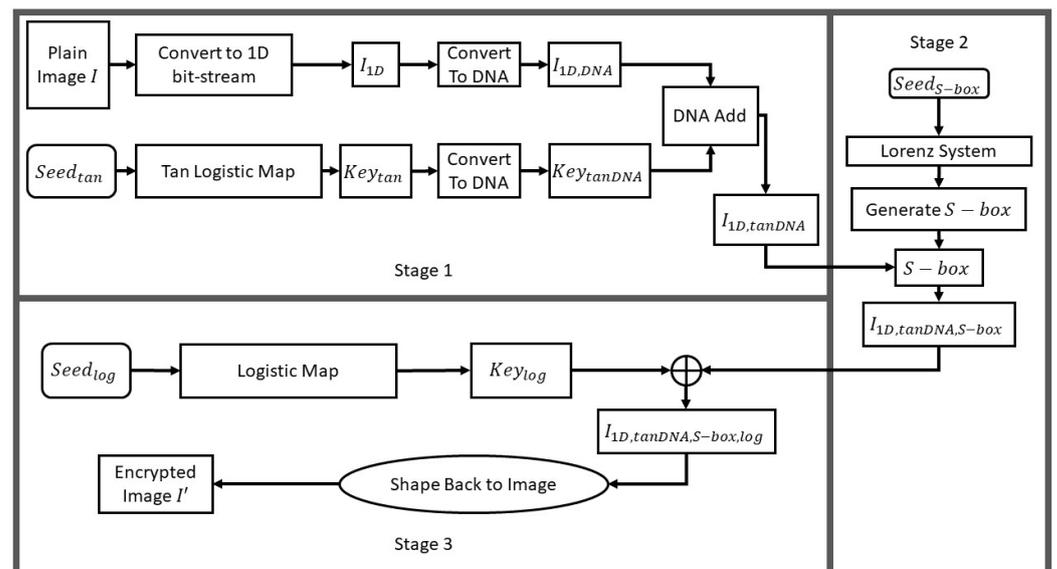


Figure 6. Flowchart of the proposed image encryption algorithm.

2.4.2. Decryption Procedure

As the reverse of the encryption procedure, decryption can be regarded as the procedure of unraveling encryption layers in order to retrieve the input image back. Hence, the previously mentioned steps are to be performed in a regressive order (as order of stages), in an inverse manner (as inverse of each performed process). Therefore, the decryption process starts with the encrypted image I' , alongside the seeds ($Seed_{tan}$, $Seed_{s-box}$ and $Seed_{log}$). Since the process of generating keys out of seeds is performed exactly the same way as in the encryption process (Key_{tan} using $Seed_{tan}$, $S-box$ using $Seed_{s-box}$, and Key_{log} using $Seed_{log}$), the decryption steps elaborated next are discussed in terms of keys instead of seeds. The decryption process is procedure as follows:

1. Stage 3: Logistic Map Encoding.
 - (a) Image I' of dimensions $M \times N$, is converted into a 1D bit-stream to reproduce $I_{1D,tanDNA,S-box,log}$.

- (b) Given $I_{1D,tanDNA,S-box,log}$, and Key_{log} , to retrieve $I_{1D,tanDNA,S-box}$, an XOR operation is performed, as follows:

$$I_{1D,tanDNA,S-box} = I_{1D,tanDNA,S-box,log} \oplus Key_{log}. \quad (14)$$

2. Stage 2: S-box Application.

- (a) Given $S - box$, the inverse $S - box$ is calculated, namely $S - box^{-1}$.
 (b) The inverse S-box is applied as:

$$I_{1D,tanDNA} = S - box^{-1}(I_{1D,tanDNA,S-box}) \quad (15)$$

3. Stage 1: DNA Encoding Based on a Tan Logistic Map.

- (a) The bit-stream $I_{1D,tanDNA}$ is first converted into a DNA sequence using (3).
 (b) In order to retrieve I_{1D} , Key_{tanDNA} is to be subtracted from $I_{1D,tanDNA}$ using Table 2, as shown in the following equation:

$$I_{1D} = SubtractedDNA(I_{1D,tanDNA}, Key_{tanDNA}) \quad (16)$$

Finally, I_{1D} is to be reshaped back into a 2D image (of dimensions $M \times N$) resulting in the input image I . Figure 7 demonstrates a flowchart for the decryption procedure.

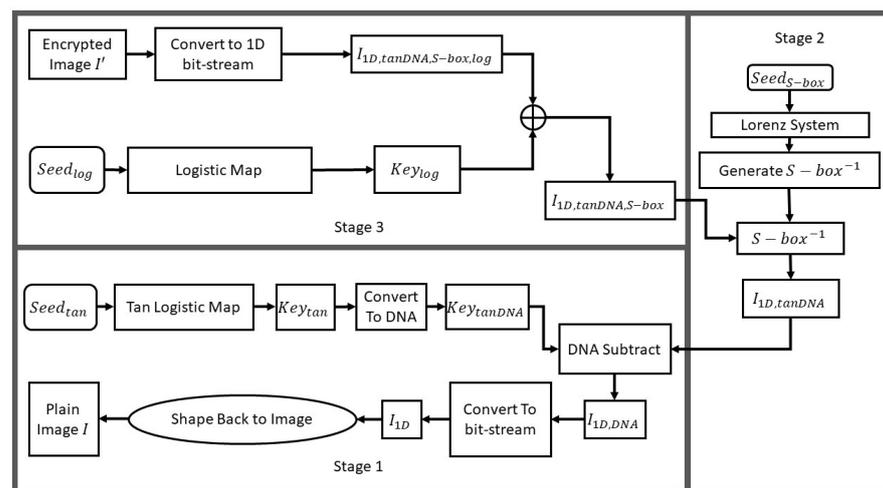


Figure 7. Flowchart of the proposed image decryption algorithm.

3. Numerical Results and Performance Evaluation

An encryption algorithm's performance is evaluated by how well it can withstand various visual, statistical, entropy, differential and brute-force attacks. In this section, the suggested image encryption algorithm's numerical findings are presented and discussed along with a comparison to counterpart algorithms from the literature. The various analyses were run on Wolfram Mathematica® v.13.1. The utilized computer had the following specifications: 2.9 GHz Intel® Core™ i9, 32 GB. For the sake of these tests, values used as keys for the experimental encryption process are assigned as follows: $\sigma = 10$, $\beta = 8/3$, $\rho = 28$ and $X_0 = 0.5$. Multiple images that are commonly used in image processing applications and experimentation were utilized, all of dimensions of 256×256 , unless otherwise stated. The performed tests are:

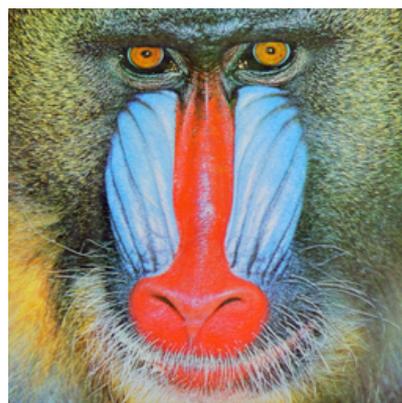
- Visual and Histogram Analysis (Section 3.1).
- Mean Squared Error (Section 3.2).
- Peak Signal to Noise Ratio (Section 3.3).
- Mean Absolute Error (Section 3.4).
- Information Entropy (Section 3.5).

- Fourier Transformation Analysis (Section 3.6).
- Correlation Coefficient Analysis (Section 3.7).
- Differential Attack Analysis (Section 3.8).
- The National Institute of Standards and Technology Analysis (Section 3.9).
- Key Space Analysis (Section 3.10).
- Histogram Dependency Tests (Section 3.11).
- Execution Time Analysis (Section 3.12).
- S-Box Performance Analysis (Section 3.13).

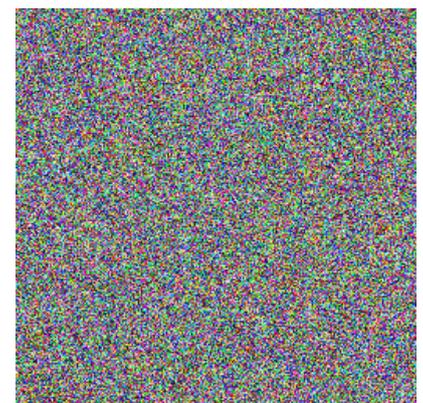
3.1. Visual and Histogram Analysis

In Figures 8–12 (including sub-figures), a number of input plain images, their encrypted forms along with their respective histograms, are shown. It is evident that the encrypted images' pixels are asymmetric. This results in the encrypted images being distorted to a high level, to the extent that all visual features in an input (plain) image being totally absent from the encrypted one.

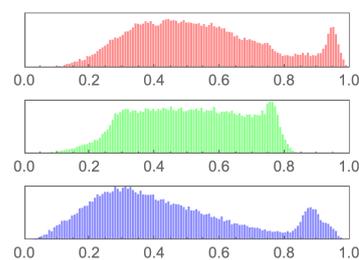
Moreover, histograms of the encrypted images are demonstrated as well. As the histogram of an image displays the frequency distribution of the pixels, the histogram of an encrypted image must be homogeneous to have a reliable encryption method. The core reason behind that is that a uniform histogram distribution reveals that each of the image's gray levels has a probability that is essentially equivalent. Therefore, the image will be more robust to statistical attacks as a result.



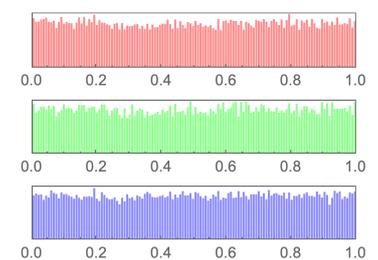
(a) Plain image.



(b) Encrypted image.



(c) Histogram of the plain image.

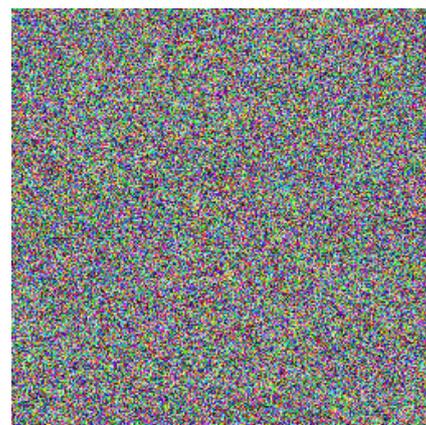


(d) Histogram of the encrypted image.

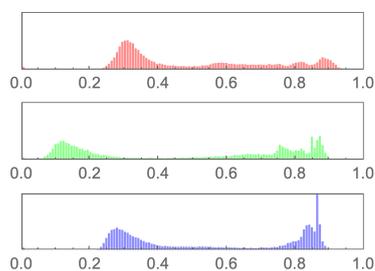
Figure 8. Mandrill image and histogram comparison before and after encryption.



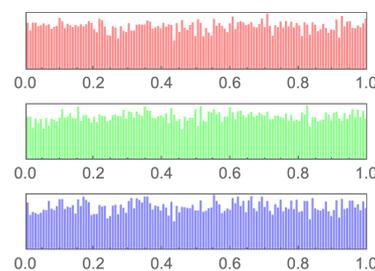
(a) Plain image.



(b) Encrypted image.

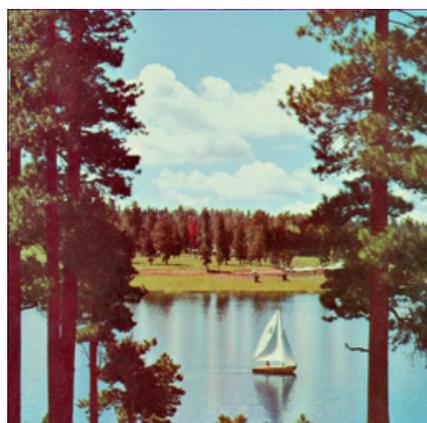


(c) Histogram of the plain image.

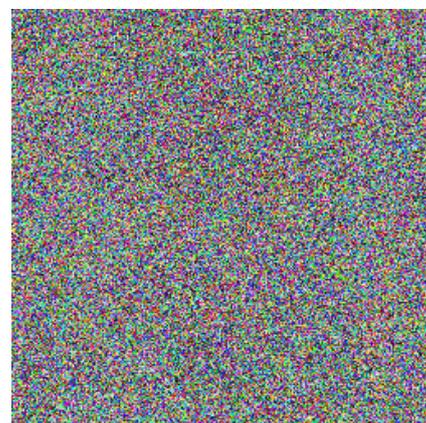


(d) Histogram of the encrypted image.

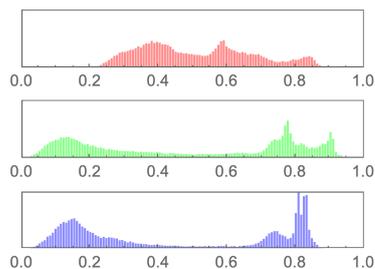
Figure 9. Tree image and histogram comparison before and after encryption.



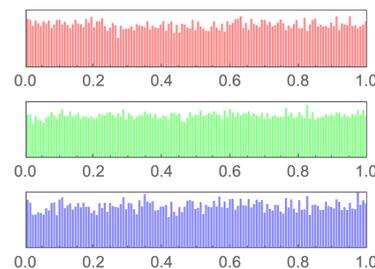
(a) Plain image.



(b) Encrypted image.



(c) Histogram of the plain image.

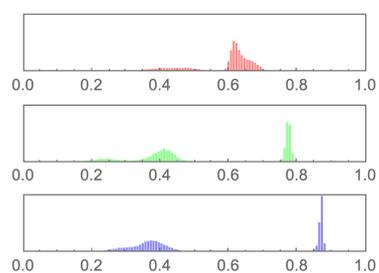


(d) Histogram of the encrypted image.

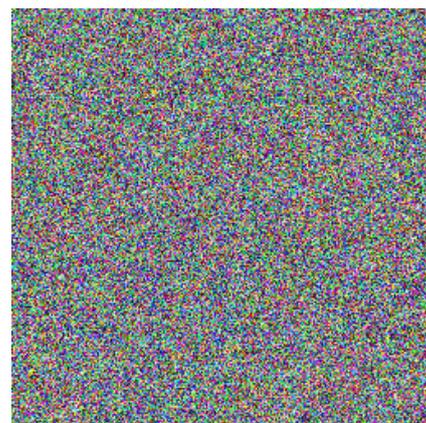
Figure 10. Sailboat image and histogram comparison before and after encryption.



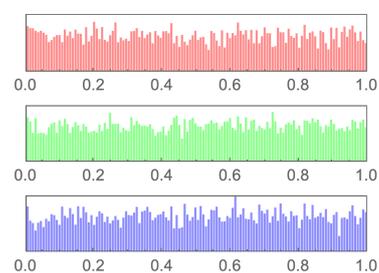
(a) Plain image.



(c) Histogram of the plain image.



(b) Encrypted image.

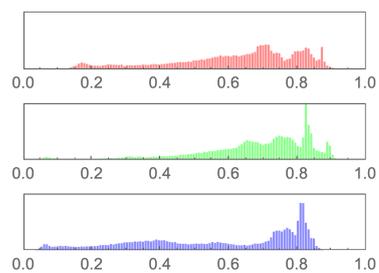


(d) Histogram of the encrypted image.

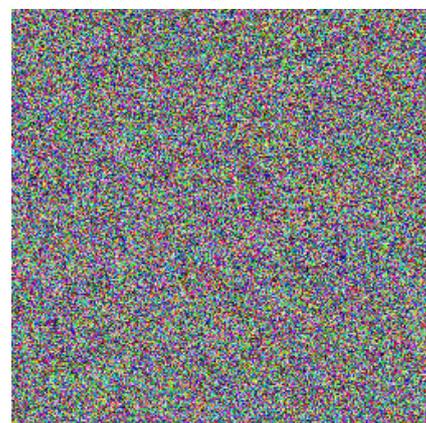
Figure 11. House image and histogram comparison before and after encryption.



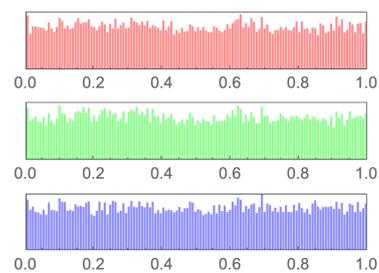
(a) Plain image.



(c) Histogram of the plain image.



(b) Encrypted image.



(d) Histogram of the encrypted image.

Figure 12. House2 image and histogram comparison before and after encryption.

3.2. Mean Squared Error

The Mean Squared Error (MSE) is one of the most common tools used in evaluating the similarity between two sets of numbers (in the most general form). As a variant of the Sum of Squared Differences (SSD), the same properties are inherited. For further elaboration, given two sets S and S' of same size n , the SSD is calculated as follows:

$$SSD = \sum_{i=1}^n (S_i - S'_i)^2. \quad (17)$$

In light of this representation, there are three main operations performed: subtraction, squaring and summing. Subtraction is mandatory as the operation desired revolves around detecting differences. Consequently, subtraction, as a mathematical operation, produces two results: direction and magnitude. As global difference (as over the whole set) is required, differences in opposing directions are to be added up (instead of cancelling each other out). Performing that, the importance of directional differences among corresponding individual elements in both lists is neglected. Mathematically, either absolute values (retaining the magnitude) or squared values (amplifying the magnitude) are used in order to remove the polarity, which is the main difference between the Sum of Absolute Differences (SAD) and the Sum of Squared Differences (SSD). Finally, summation allows all individual elements in the list to contribute (equally) to the final result, achieving a calculation of the global perspective.

In case of the MSE, the mathematical representation is modified to the following:

$$MSE = \frac{\sum_{i=0}^{M-1} \sum_{j=0}^{N-1} (I_{(i,j)} - I'_{(i,j)})^2}{M \times N}. \quad (18)$$

Given two images I and I' of dimensions $M \times N$, as images rectangular areas, two-dimensional summing is required. (Each dimension of the summation is starting at 0 ($i = 0$ and $j = 0$), and ending by $M - 1$ and $N - 1$ assuming the image (as a 2D array) is zero-indexed.) What can be perceived as a considerable alteration to the SSD is the division by the image dimensions. Such a step is performed in order to facilitate the comparison between MSE values in which image pairs' dimensions are different. For example, I_1 and I'_1 are both of size 256×256 with Mean Squared Error MSE_1 (both images must be of the same size). Comparing MSE_1 to MSE_2 which is calculated for I_2 and I'_2 of dimensions 512×512 is meaningful as both values are normalized with respect to images' scales. This entails that MSE can be preserved as the average error (in terms of SSD) per pixel, given two images.

In the scope of this work, MSE is evaluated for input images and their encrypted counterparts. In such a case, the ideal value for a well-performing encryption technique is expected to be high. In other words, as the target of encryption is to distort the visual attributes of images, the similarity should be minimal, resulting in a maximal error factor. Table 4 shows the computed MSE values for various input image examples, alongside showcasing how these values stand in comparison to other encryption techniques in the literature, demonstrating comparable results.

It is common practice to report MSE and Peak Signal to Noise Ratio (PSNR) values together when analyzing image encryption algorithms. This is because the computation of PSNR is based on the value of MSE. However, the authors of [39,40] only provide PSNR values in their respective works, with no mention of MSE values. This explains why Table 4 displays columns of N/A under the headings of [39,40].

Table 4. MSE values comparison with the literature.

Image	Proposed Scheme	[16]	[17]	[18]	[26]	[39]	[40]
Lena	9112.1	8926.96	10,869.73	4859.03	8888.88	N/A	N/A
Mandrill	8573.38	8290.84	10,930.33	6399.05	8295.21	N/A	N/A
Peppers	10,298.7	10045.1	N/A	7274.44	10,092.3	N/A	N/A
House	8427.04	8351.64	N/A	N/A	N/A	N/A	N/A
House2	9374.65	N/A	N/A	N/A	N/A	N/A	N/A
Girl	12,450.9	N/A	N/A	N/A	N/A	N/A	N/A

3.3. Peak Signal to Noise Ratio

Based on the MSE discussed in Section 3.2, PSNR, given a signal, aims at relating the error margin (represented by MSE), with respect to the peak value in the signal. In the scope of this work, the peak signal value is evaluated as the maximum pixel intensity in a given image. Accordingly, given image I , PSNR is equated as:

$$PSNR = 10 \log \left(\frac{I_{max}^2}{MSE} \right), \quad (19)$$

such that I_{max} is the maximum pixel intensity in I . Due to the fact that MSE is SSD based (which indicates that MSE is calculated to a squared order of magnitude), I_{max} is necessarily squared.

As shown in (19), PSNR is inversely proportional to MSE. Such mathematical representation steers the preference of the PSNR to be the inverse of the preference of MSE, hence a minimal value is more ideal. Table 5 presents the calculated values for the proposed cryptosystem as well as those of counterpart algorithms from the literature. In terms of MSE and PSNR, the proposed cryptosystem is shown to be superior to [16,18,26,39,40], but inferior to [17].

Table 5. PSNR values comparison with the literature.

Image	Proposed Scheme	[16]	[17]	[18]	[26]	[39]	[40]
Lena	8.53462	8.6237	7.7677	11.3	8.64233	8.5674	8.617
Mandrill	8.79929	8.9448	7.7447	10.10	8.94253	10.0322	8.9695
Peppers	8.00296	8.11128	N/A	9.55	8.94253	N/A	8.1156
House	8.87405	8.91309	N/A	N/A	N/A	N/A	8.935
House2	8.41125	N/A	N/A	N/A	N/A	N/A	8.5343
Girl	7.17879	N/A	N/A	N/A	N/A	N/A	7.282

3.4. Mean Absolute Error

Building on the argument presented in Section 3.2, an alternate technique to Sum of Squared Differences, SSD, would be Sum of Absolute Differences, SAD. In such alternative, the task of eliminating the polarity of the per-pixel error is performed by the absolute operation instead of the square one. Therefore, parallel to the SSD equation (Equation (17)), SAD is equated as:

$$SAD = \sum_{i=1}^n |S_i - S'_i|. \quad (20)$$

As discussed before in Section 3.2, the role of squaring, beside eliminating polarity, is to amplify the magnitude. On the other hand, using the absolute instead maintains the linearity of the behaviour of the error distribution among pixels, which is accordingly maintained in the global perspective of the whole image.

Upon such variation in the core function, MAE is represented mathematically as:

$$MAE = \frac{\sum_{i=0}^{M-1} \sum_{j=0}^{N-1} |I_{(i,j)} - I'_{(i,j)}|}{M \times N}, \quad (21)$$

for two images I and I' . As per this work (similar to the MSE scenario), MAE is evaluated for input images and their encrypted images. Moreover, for a well-performing encryption technique, MAE value is preferred to be maximal. Table 6 presents the numerical results of performing the MAE test on three images (Lena, Peppers, and Mandrill) in comparison to counterpart algorithms from the literature. As the numerical results demonstrate, the proposed algorithm fares comparably to them.

Table 6. MAE analysis of the Lena, Peppers and Mandrill images.

Image	Proposed Scheme	[26]	[17]	[44]	[39]
Lena	78.3564	77.3752	87	77.35	77.96
Peppers	82.3273	81.7740	N/A	74.71	N/A
Mandrill	81.913	75.1659	92	73.91	67.85

3.5. Information Entropy

In the domain of gray-scale images, information entropy is employed to measure the randomness of the distribution of gray pixel values of an image. According to Shannon's theory, information entropy is calculated as:

$$H(m) = \sum_{i=1}^M p(m_i) \log_2 \frac{1}{p(m_i)}, \quad (22)$$

where $p(m_i)$ refers to the probability of occurrence of symbol m , while M represents the total number of bits for each symbol. With respect to images, as a gray scale image has 256 different values $[0 - 255]$, which are 2^8 different possible permutations, the entropy value of an encrypted image at greatest evaluations approaches 8. Therefore, information entropy can be used to evaluate the degree of randomness of encrypted images. Entropy values of the proposed algorithm for the the images displayed and tested in this paper, along with counterpart algorithms from the literature are displayed in Table 7. The computed entropy values for the various images are very close to the ideal value of 8, which means that the proposed algorithm is resistant to entropy attacks. Furthermore, the differences across entropy values of the various cryptosystems is shown to be diminutive.

Table 7. Entropy values for encrypted images.

Image	Proposed	[16]	[17]	[45]	[18]	[26]	[39]	[40]
Lena	7.9856	7.999	7.999	7.997	7.996	7.997	7.9972	7.999
Mandrill	7.9905	7.999	7.999	7.999	N/A	7.996	7.9969	7.9991
Peppers	7.9951	7.999	7.9991	N/A	7.997	7.9969	N/A	7.9991
House	7.9577	7.999	N/A	N/A	N/A	N/A	N/A	7.999
House2	7.9847	N/A	N/A	N/A	N/A	N/A	N/A	7.999
Girl	7.9789	N/A	N/A	N/A	N/A	N/A	N/A	N/A

3.6. Fourier Transformation Analysis

In order to showcase the co-relation among pixels before and after encryption, the application of the Fourier transformation (for both images) is utilized, more accurately, for Discrete Fourier Transform (DFT). The main aim is, in the frequency domain, visual features such as edges and regions (which are not easily definable in the spatial domain) separate into different frequency ranges. This facilitates visual analysis and comparisons of images. Such separation takes place as a result of the interaction between the pixels in the spatial image with the increasing frequencies of the sine and cosine waves. Aiming

towards transforming a spatial domain image into the frequency domain, the following Fourier transformation equation is used:

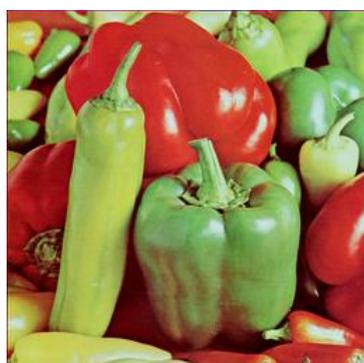
$$F(k,l) = \sum_{i=0}^{N-1} \sum_{j=0}^{N-1} f(i,j)e^{-i2\pi(\frac{ki}{N} + \frac{lj}{N})}, \quad (23)$$

such that $f(a,b)$ is the image representation in the spatial domain, with the exponential term being the basis function that corresponds to each point $F(k,l)$ in the Fourier space.

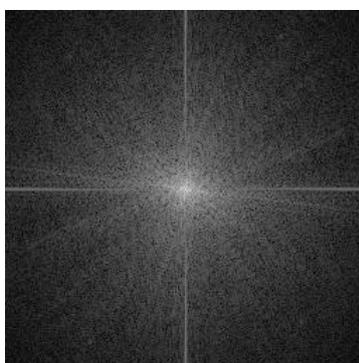
Interpreting a Fourier transformation of an image (presented as 2D data), two main regions of high relevance are to be looked at [46]. The first region is the middle area of the 2D grid of the Fourier transformed image. The main significance of this area is that it represents the amount of pixels with high similarity in values on the pixel level. Therefore, if a Fourier transformation image is generated out of an image with large regions (flat, same color areas), it is expected for the middle area to contain high values.

The second region is the center row, column and alongside diagonals of the Fourier transformed image. This is due to their representation of vertical, horizontal, and diagonal edges in the input image, respectively. (The Fourier image is considered to be a transposed matrix of the input image.) Thus, if the input image includes only vertical edges, the Fourier transformed image is expected to have a bright middle row, and vice versa. The rest of the Fourier transformed image represents the other features existing in the input image.

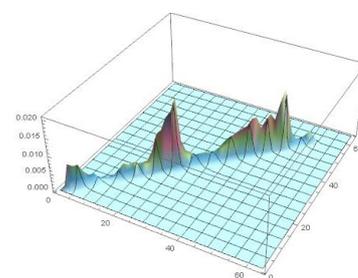
Conclusively, for a naturally looking image with wide regions and profound edges (Peppers in this example), the resulting Fourier transformed image contains a bright plus sign at its center, as observed in Figure 13b. On the other hand, on a distorted (encrypted) image, an equal description of values in the Fourier transformed image is expected due to the lack of profound regions or edges, as shown in Figure 13e.



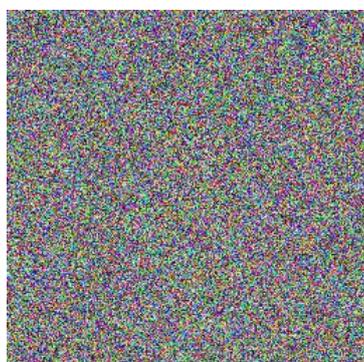
(a) Plain image.



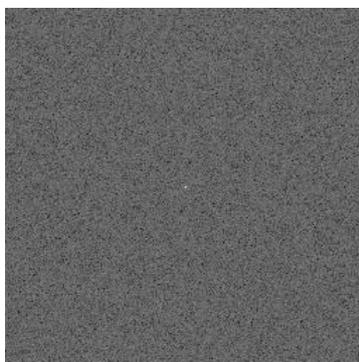
(b) Plain image Fourier.



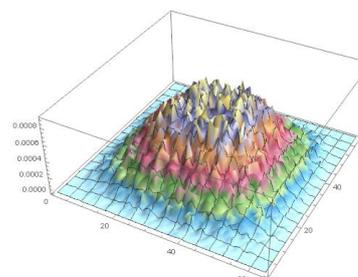
(c) Plain image co-occurrence.



(d) Encrypted image.



(e) Encrypted image Fourier.



(f) Encrypted image co-occurrence.

Figure 13. Peppers image alongside its Fourier transformation and 3D plot of its co-occurrence matrix before and after encryption.

3.7. Correlation Coefficient Analysis

In this evaluation method, the consistency of a single image is evaluated. The aim of such evaluation method is to assess (or provide an estimated coefficient for) the cohesion of near-proximity pixels. In other words, correlation coefficient analysis, in the domain of images, aims at calculating the percentage of the uniform regions with respect to the edge transitions. Hence, in a normal image case, a relatively high correlation coefficient value is expected, as it consists more of regions than edges (in terms of pixel count). On the other hand, as high distortion is aimed for in encrypted images, a smaller correlation coefficient is anticipated for it.

For mathematical evaluation, the correlation coefficient is calculated through the following equations:

$$\rho(x, y) = \frac{\text{cov}(x, y)}{\sqrt{\sigma(x)}\sqrt{\sigma(y)}}, \quad (24)$$

where:

$$\text{cov}(x, y) = \frac{1}{N} \sum_{i=1}^N (x_i - \mu(x))(y_i - \mu(y)), \quad (25)$$

$$\sigma(x) = \frac{1}{N} \sum_{i=1}^N (x_i - \mu(x))^2, \quad (26)$$

$$\mu(x) = \frac{1}{N} \sum_{i=1}^N (x_i). \quad (27)$$

Starting by the mean average of each distribution as in (27), two distribution behaviours are measured. The first behaviour is the dispersion (using (26)), which represents the uncertainty of the distribution. The second behaviour is the covariance (using (25)), which evaluates the linear direction similarity. Combining both mathematical phenomena results in reaching an evaluation for the correlation coefficient, represented in (24).

As previously mentioned, for a normal image (the input), a high correlation value is expected. On the other hand, a highly distorted image would result in having a low correlation coefficient. Table 8 demonstrates performing the correlation coefficient analysis on three images (Lena, Peppers and Mandrill), for both images, input and encrypted. Moreover, as the covariance is a directional relation, the three main directions are calculated, which are horizontal, vertical and diagonal. As shown by the numerical results, the input image showed a value approaching 1 in all cases, while the encrypted showed a value approaching 0. Table 9 presents the comparison between the proposed approach and counterpart algorithms from the literature, which showcases nearly similar results. Moreover, Tables 10 and 11 show the results of numerical comparison among the proposed algorithm and its counterparts from the literature, focusing on the color channels separately, with respect to the three directions for the images Lena and Mandrill, respectively.

Table 8. Correlation coefficients of plain and encrypted images.

Image	Plain Image			Encrypted Image		
	Horizontal	Diagonal	Vertical	Horizontal	Diagonal	Vertical
Lena	0.96734	0.94821	0.98276	0.003265	−0.00413	0.002451
Peppers	0.95595	0.95371	0.97939	0.004331	0.001856	0.001043
Mandrill	0.92203	0.87049	0.90303	−0.00484	0.003429	0.003522

Table 9. Correlation coefficients comparison between Input and encrypted Lena images.

Scheme	Horizontal	Diagonal	Vertical
Proposed	0.003265	−0.00413	0.002451
[26]	0.002287	−0.00132	−0.00160
[47]	0.0022	−0.0017	0.0001
[17]	0.0054	0.0054	0.0016
[48]	0.000199	0.003705	−0.000924
[49]	0.0681	0.0128	0.0049
[50]	0.001862	0.003768	0.000710
[51]	−0.0082	−0.0012	−0.0128
[52]	0.000546	0.000192	0.000514
[53]	−0.0029	−0.0045	−0.0001
[54]	0.0023	−0.0059	0.0029
[39]	−0.0061	−0.0018	0.0067

Table 10. Comparison between the three directions of correlation coefficients, for plain and encrypted Lena images, performed on each color channel separately, with respect to results from the literature.

Channel	Direction	Plain Image	Encrypted Image	[55]	[56]	[57]	[26]
Red	Horizontal	0.95722	0.006559	0.001365	0.0021	0.9568	−0.00364
	Diagonal	0.93389	−0.00145	0.000232	−0.0026	0.0075	0.00016
	Vertical	0.97889	0.002	0.004776	0.0018	−0.0376	0.000697
Green	Horizontal	0.94321	0.00295	0.003294	−0.0006	0.0020	0.000118
	Diagonal	0.91931	−0.001739	0.004807	0.0001	−0.0046	0.00177
	Vertical	0.97137	0.001745	−0.000579	0.0004	−0.0013	−0.0011
Blue	Horizontal	0.92845	−0.00278	0.002060	−0.005	0.0071	−0.00164
	Diagonal	0.90068	0.000744	−0.004043	−0.0104	−0.0009	−0.00523
	Vertical	0.95593	0.0051	0.000194	0.001	−0.0423	0.006041

Table 11. Comparison between the three directions of correlation coefficients, for plain and encrypted Mandrill images, performed on each color channel separately, with respect to results from the literature.

Channel	Direction	Plain Image	Encrypted Image	[55]	[56]	[26]
Red	Horizontal	0.94741	−0.00383	0.001391	0.0005	−0.00428
	Diagonal	0.90413	0.000245	0.000334	0.0014	−0.00009
	Vertical	0.92152	−0.00571	0.004650	0.0059	0.000706
Green	Horizontal	0.87266	−0.00357	−0.008134	0.0078	0.00340
	Diagonal	0.79341	0.003297	0.005334	−0.001	0.00282
	Vertical	0.83905	0.006606	0.000829	0.0042	−0.0016
Blue	Horizontal	0.92153	0.000063	−0.00889	0.0021	−0.00253
	Diagonal	0.87668	−0.00334	0.001710	−0.0114	−0.00635
	Vertical	0.91432	0.001022	0.000056	−0.0039	−0.00003

Alongside the numerical analysis provided by (24), directional covariance can be visualized by the plotting of the co-occurrence matrix. In case of images with natural visual aspects (more pixels representing homogeneous regions than transitional edges), values of high similarity tends to co-exist with a higher probability, resulting in a mostly linear distribution of magnitudes within the matrix. Oppositely, in a highly distorted (encrypted) image, a more equal distribution of values is expected to take place instead. For demonstration, this is carried out in Figure 13, where Figure 13c is for the plain image, while Figure 13f is for its encrypted version. It is clear that the 3D plot for the plain image is diagonal in nature, unlike that of the encrypted image, which resembles a mountain

in 3D space, as expected for an encrypted image, signifying random pixel locations. For further demonstration, focusing on the Papers image, Figures 14–16 show 3D plots of the co-occurrence matrices for the red, green and blue color channels, respectively. As demonstrated, pixel correlations are fully distorted on each color level individually.

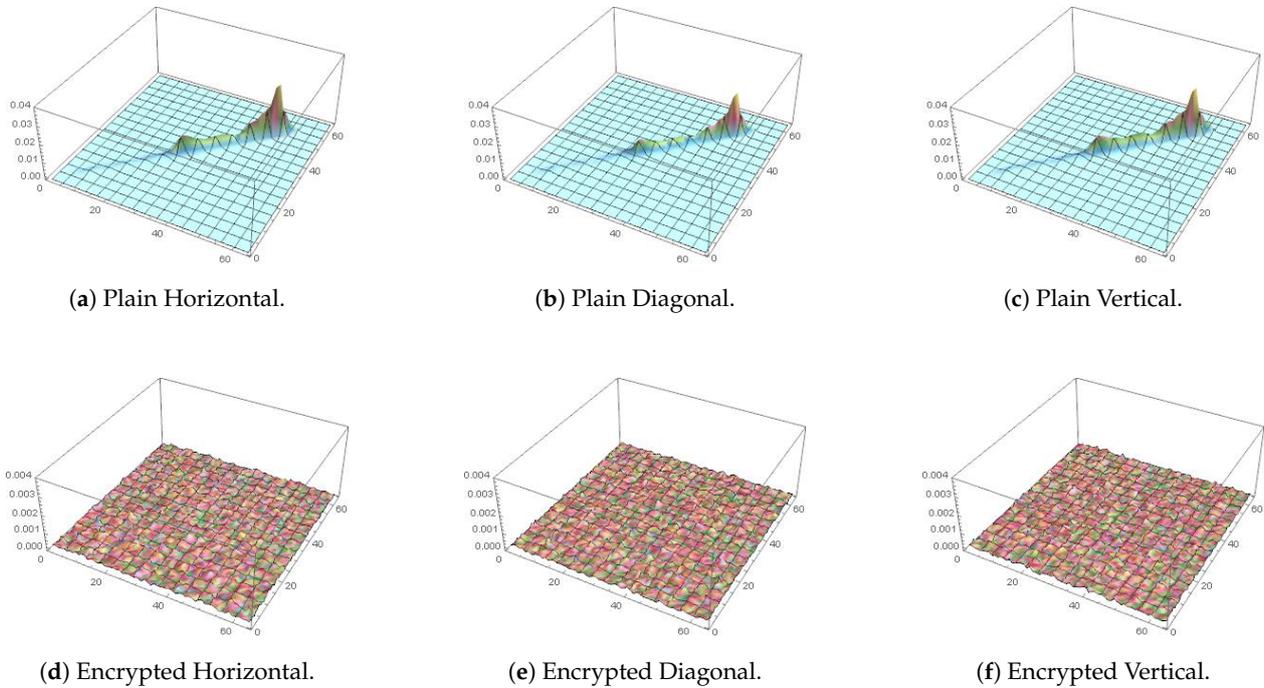


Figure 14. Peppers 3D plot of its co-occurrence matrix before and after encryption for red channel.

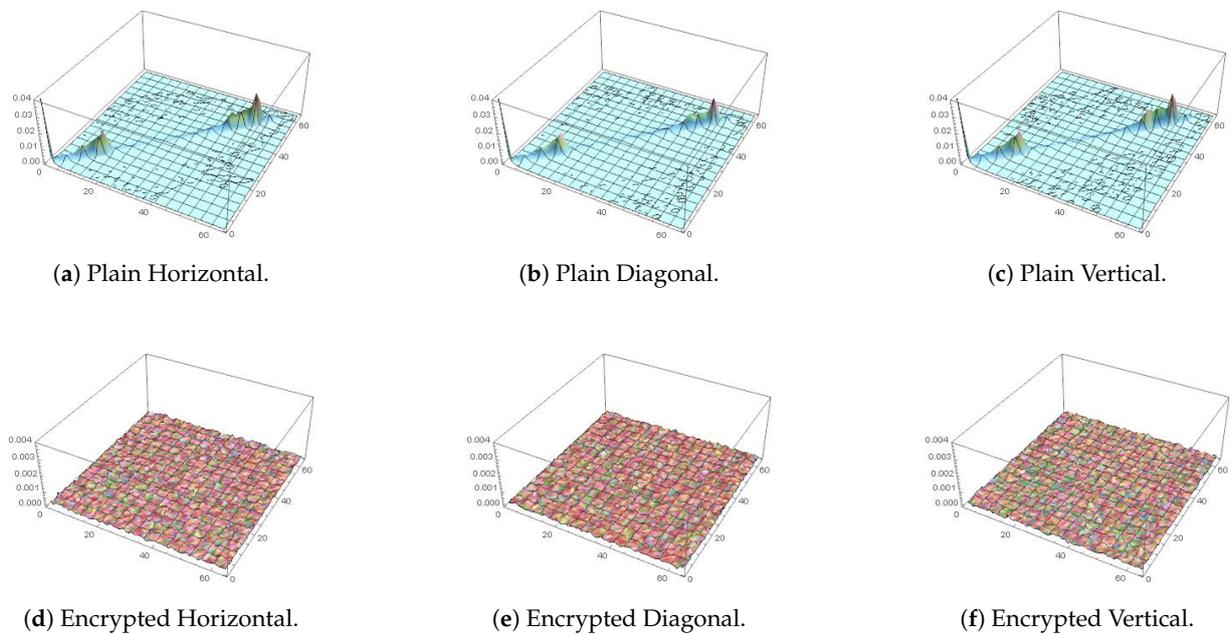


Figure 15. Peppers 3D plot of its co-occurrence matrix before and after encryption for green channel.

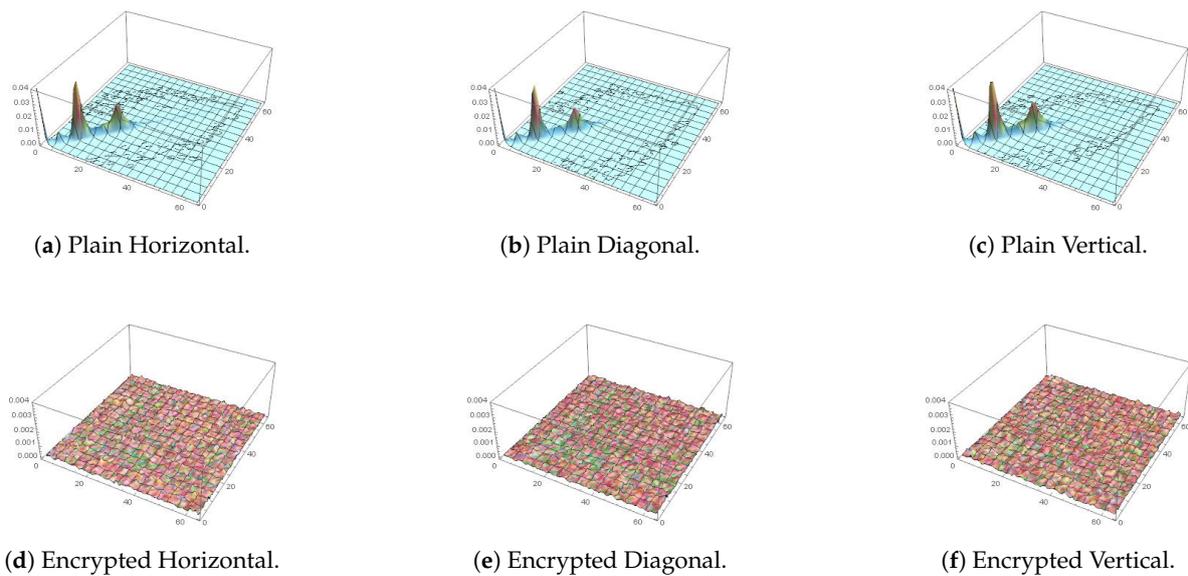


Figure 16. Peppers 3D plot of its co-occurrence matrix before and after encryption for blue channel.

3.8. Differential Attack Analysis

In this section, the quality of image cryptosystems is judged on the grounds of the direct difference between the input and the encrypted images. In other words, the input image is directly compared with the encrypted on pixel by pixel, or mean average bases. Such evaluation is performed in order to calculate a numerical percentage showcasing the difference in color intensities (per pixel, or as a mean average) taking place as a result of the encryption process. According to the fact that the lack of similarity among corresponding pixels within both images is encouraged, such pixel by pixel evaluation is necessarily performed. Moreover, a more global perspective of the cumulative pixels change rates among images (presented in the mean averages) is evaluated, which denotes the existence of general color intensity similarity among these images. In the literature, two tests are most commonly performed to fulfill these test aspects: NPCR for pixel by pixel comparison, and UACI for the mean average difference evaluation.

Number of pixels changing rate (NPCR) represents the percentage evaluation of the amount of changed pixels. The difference between pixels is performed with a strict equality perspective. Given two images I_1 and I_2 (of dimensions $M \times N$), the difference per pixel $D(x, y)$ (where x and y are the coordinates of the pixel) is calculated as:

$$D(x, y) = \begin{cases} 0 & I_1(x, y) = I_2(x, y) \\ 1 & \text{Otherwise} \end{cases} \Big|_{x \in [1, M] \& y \in [1, N]} \quad (28)$$

Accordingly, NPCR is equated as:

$$NPCR = \frac{\sum_{x=1}^M \sum_{y=1}^N D(x, y)}{M \times N} \times 100. \quad (29)$$

As per this representation, a higher percentage denotes larger difference between the two images. As a large difference is desired, in the literature, 99% is the target NPCR score for a good encryption technique.

In another perspective, the unified average change intensity (UACI) evaluates the difference between two images in terms of the mean averages. Mathematically, UACI is equated as:

$$UACI = \frac{1}{M \times N} \sum_{x=1}^M \sum_{y=1}^N \frac{|I_1(x, y) - I_2(x, y)|}{255} \times 100. \quad (30)$$

In the literature, a target percentage of more than 33% denotes a strong encryption technique. (With respect to the color range [0, 255], 33% is approximated to 85 steps of difference in intensity.)

Table 12 shows the result of performing NPCR and UACI, comparing the input and the encrypted image generated by the proposed algorithm, using various images as inputs. (As discussed before, the NPCR is greater than 99.6% and the UACI should also be greater than 33%.) For these tests, the proposed algorithm is also compared with its counterparts from the literature, as shown in Tables 13 and 14. As shown, the computed NPCR value of the proposed algorithm is >99% in all cases. On the other hand, the UACI value did not meet the optimal value. However, the exception to this case is the Girl image, which resulted in 32.1283, 36.4526, and 37.4855 for color channels red, green and blue, respectively, alongside 35.3554 as an overall.

Table 12. NPCR and UACI of different images.

Test Type	Image	Result
NPCR	Lena	99.647
	Peppers	99.6348
	Mandrill	99.6124
	House	99.6063
	House2	99.6155
	Girl	99.617
UACI	Lena	29.4651
	Peppers	32.1832
	Mandrill	29.3973
	House	29.3973
	House2	30.716
	Girl	35.3554

Table 13. NPCR and UACI values of different images' color channels comparison.

Test Type	Image	Color Channel	Proposed	[26]	[58]
NPCR	Lena	Red	99.6262	99.6109	99.6355
		Green	99.6201	99.6109	99.6256
		Blue	99.6948	99.6375	99.6159
	Peppers	Red	99.6658	99.6032	99.6307
		Green	99.6262	99.6032	99.6250
		Blue	99.6124	99.3750	99.6213
	Mandrill	Red	99.5605	99.5880	99.6102
		Green	99.6353	99.5880	99.6134
		Blue	99.6414	99.5880	99.6057
UACI	Lena	Red	33.0311	33.4158	33.4657
		Green	30.7273	30.3902	33.4552
		Blue	27.6116	33.2420	33.4550
	Peppers	Red	28.9610	33.3459	33.4832
		Green	33.7841	33.4702	33.4904
		Blue	33.8043	33.4357	33.4619
	Mandrill	Red	29.5056	33.4273	33.5002
		Green	28.0120	33.4635	33.4711
		Blue	30.6723	33.7951	33.4951

Table 14. Average NPCR and UACI of the Lena image comparison.

Scheme	NPCR	UACI
Proposed	99.65	30.4567
[26]	99.63	30.3432
[40]	99.63	33.48
[17]	99.52	26.7933
[59]	99.61	33.4342
[60]	99.52	26.7933
[39]	99.61	33.5160

3.9. The National Institute of Standards and Technology Analysis

The National Institute of Standards and Technology (NIST) SP 800 analysis is a set of statistical tests which ensure the necessary cryptographic properties of the random number sequences are met. The encrypted images' equivalent binary stream are run through the NIST analysis. To ensure resilience to cryptographic attacks, the results ought to surpass a p -value of 0.01. Table 15 shows the outcome of running the analysis. The results reflect the cryptographic robustness of the proposed scheme, with all the tests' outcomes larger than 0.01. Hence, we can safely conclude the validity of our proposed cryptosystem.

Table 15. NIST analysis on Lena encrypted image.

Test Name	Value	Remarks
Frequency	0.504134	Success
Block Frequency	0.715728	Success
Run	0.185355	Success
Long runs of ones	0.897733	Success
Rank	0.368065	Success
Spectral FFT	0.783087	Success
No overlapping	0.979028	Success
Overlapping	0.224884	Success
Universal	0.937934	Success
Linear complexity	0.750498	Success
Serial	0.012139	Success
Approximate Entropy	0.369179	Success
Cumulative sum (forward)	0.252025	Success
Cumulative sum (reverse)	0.338189	Success

3.10. Key Space Analysis

Key space analysis is calculated as the Cartesian product of the domains of the key values involved in the encryption procedure. Such a step is carried out in order to compute the amount of unique keys that can be utilized in the encryption procedure, which accordingly results in creating various possible encryption instances for the same input image. In the proposed image encryption algorithm, there is a total of seven variables involved in a single encryption procedure. The first two variables are used (in the first stage Section 2.1) for the generation of the bit-stream using the tan variation of the logistic map, which is used in the DNA encoding, namely α and X_0 . In the second stage (Section 2.2), a Lorenz system is used for the S-box generation, which demands three variables (σ , ρ and β). Finally, in the last stage discussed in Section 2.3, two keys are used, which are r and $t_r(0)$. As the largest machine precision is 10^{-16} , the key space is about $10^{7 \times 16} = 10^{112} \approx 2^{372}$, which exceeds the threshold earlier proposed in [61] as 2^{100} . This means that our proposed scheme can resist brute-force attacks. Furthermore, an examination of key space values of related image encryption schemes from the literature, as in Table 16 indicates that the proposed scheme utilizes a comparably much larger key space than most of its counterparts. The only exception here being the work of [39], in which one of the encryption stages relies on the random movement of a chess piece (Castle), which results in a very large key space.

Table 16. Key space values comparison.

Algorithm	Key Space
Proposed algorithm	2^{372}
[28]	2^{256}
[29]	2^{256}
[39]	2^{604}
[48]	2^{187}
[62]	2^{345}
[63]	2^{128}
[64]	2^{312}
[65]	2^{128}
[66]	2^{219}

3.11. Histogram Dependency Tests

According to the aim of image encryption (distorting image details in a reversible manner), all forms of correlation between the plain image and its encrypted version is to be absent. In such a test scenario, the images (plain and encrypted) are evaluated on the histogram level. Moreover, these tests are performed on the image as one unit, as well as on the color channels separately. Therefore, given two histograms of images, the comparisons performed aims at evaluating the level of the linear dependency between them. As any dependency (as a form of correlation) test evaluates the level of association between two variables [67], in a well performing encryption technique, the dependency level should be as low as possible. Accordingly, calculating the dependency coefficient as a value in the range $[-1, 1]$, it is desirable to be as close to 0 as possible (where 1 means strong dependency, and -1 means strong inverse dependency). In other words, given two distributions, dependency between them is evaluated as the alignment of one with respect to the other, which is either both are following the same linearity (evaluating to 1), or following perpendicular linearity (evaluating to -1), or there is no linearity (evaluating to 0). As mentioned, in the context of this work, the pair distributions of variables to evaluate are the histograms of both the input and the encrypted images. Out of many dependency evaluation techniques, in this work, five tests are performed, namely: Blomqvist β , Goodman-Kruskal γ , Kendall τ , Spearman ρ , and Pearson correlation r [68].

As a medial correlation coefficient, Blomqvist evaluates correlation between two distributions of variables X and Y , with their medians \bar{x} and \bar{y} , respectively, as per the following Equation:

$$\beta = \{(X - \bar{x})(Y - \bar{y}) > 0\} - \{(X - \bar{x})(Y - \bar{y}) < 0\}. \quad (31)$$

Considering the median as a reference point, pairs of elements across the two distributions of variables are either on the same side of the median (creating a linear correlation), or not (breaking the linear correlation).

Based on the relative order of succeeding elements in the two distributions of variables, the Goodman-Kruskal measure of monotonic association is used in a pairwise manner. Transforming the two histograms into one set of pairs in a 1 to 1 formation (containing pairs of the form $(H_1(n), H_2(n))$), comparing two pairs $((H_1(i), H_2(i))$ and $(H_1(j), H_2(j))$ for example), they are either aiding the linear correlation (concordant pairs) or breaking it (discordant pairs). Counting both concordant pairs and discordant pairs provides two counts, n_c and n_d , respectively. Given these two counts, Goodman-Kruskal correlation is equated as:

$$\gamma = \frac{n_c - n_d}{n_c + n_d}. \quad (32)$$

Based on the same concept of concordant pairs and discordant pairs, Kendall evaluates correlation with respect to the sample size n , equating τ as follows:

$$\tau = \frac{n_c - n_d}{\frac{n(n-1)}{2}} \tag{33}$$

For a rank-based correlation test, Spearman rank correlation test relates the rank of an element (its position if the list was sorted), with respect to the mean rank value. Spearman rank correlation is equated as:

$$\rho = \frac{\sum(R_{ix} - \bar{R}_x)(R_{iy} - \bar{R}_y)}{\sqrt{\sum(R_{ix} - \bar{R}_x)^2 \sum(R_{iy} - \bar{R}_y)^2}} \tag{34}$$

where x and y are the two evaluated variables, R_{il} is the rank of element i in list l , and \bar{R}_l is the mean of ranks of l .

Finally, Pearson correlation, as the most popular and straightforward correlation technique, simply relates elements in the distributions directly to their mean averages. Pearson correlation is equated as:

$$r = \frac{\sum(X_i - \bar{X})(Y_i - \bar{Y})}{\sqrt{\sum(X_i - \bar{X})^2 \sum(Y_i - \bar{Y})^2}} \tag{35}$$

where \bar{X} and \bar{Y} are the means of the distributions X and Y , respectively.

Table 17 shows the results of performing the six tests on various images. As all scores are approaching 0, there is a very minimal dependency between the input and encrypted images in terms of histograms over all color channels.

Table 17. Histogram dependency tests for various images.

Image	Color	β (31)	γ (32)	τ (33)	ρ (34)	r (35)
Lena	Red	-0.109375	-0.0701499	-0.0682424	-0.102114	-0.101001
	Green	-0.0589406	-0.0859604	-0.0851644	-0.126553	-0.118587
	Blue	-0.078125	-0.077486	-0.0733405	-0.105547	-0.116345
	Combined	0.0432231	0.0110378	0.0110029	0.0169526	0.0136941
Peppers	Red	0	0.0288035	-0.000855709	0.0436675	0.0261068
	Green	-0.0755594	-0.0436322	-0.0433383	-0.0637195	-0.0674477
	Blue	-0.146559	-0.0669877	-0.066303	-0.0992495	-0.112761
	Combined	0.125	0.0554493	0.055267	0.0827907	0.00394907
Mandrill	Red	-0.0476265	-0.0467586	-0.0464472	-0.0708594	-0.0700444
	Green	0.00394524	-0.00409397	-0.00403802	-0.00467083	0.0128912
	Blue	0.0711601	0.025111	0.0249474	0.0376744	0.0287206
	Combined	0	0.00135451	0.00135127	0.000814718	-0.00796008
House	Red	0	0.0124787	0.0121948	0.0217842	0.0585462
	Green	-0.0510818	0.00839867	0.00835041	0.0114007	0.0658291
	Blue	-0.122302	-0.0590104	-0.0567747	-0.081944	-0.077054
	Combined	0.0941184	-0.00570442	-0.0056861	-0.0037896	-0.0856705
House2	Red	0.019725	0.0458756	0.0453159	0.0647523	0.058933
	Green	0	-0.0115376	-0.0114824	-0.0167042	-0.056937
	Blue	0.0591722	0.0672817	0.0664841	0.0960327	0.150064
	Combined	-0.046875	0.0259157	0.025856	0.0381908	0.051188
Girl	Red	0.0942594	0.0389334	0.0330354	0.0431319	0.00880097
	Green	-0.0359493	0.0201752	0.0167873	0.0231847	-0.0228
	Blue	-0.0205128	-0.0861264	-0.0701142	-0.0957639	-0.0820709
	Combined	0.015625	0.0153764	0.0147366	0.0201923	-0.0580052

3.12. Execution Time Analysis

Encryption and decryption times are used to determine an algorithm’s complexity and suitability for real-time applications. Table 18 displays those values for the Lena

image at various dimensions, $N \times N$, where $N \in \{64, 128, 256, 512, 1024\}$. Depending on the image dimensions, the overall encryption and decryption time varies from 0.228822s, for $N = 64$, to just under a minute, for $N = 1024$. In addition, Table 19 presents a comparison of the encryption time among the proposed algorithm and its counterparts from the literature. Note that the differences in encryption time depend on numerous factors, including the algorithm's complexity, the machine specifications on which the algorithm is executed (i.e., processing power and accessible memory), and the software package or programming language used to execute the algorithm. In this work, Wolfram Mathematica[®] is employed, while in [39,66,69–71] Mathworks Matlab[®] was the software of choice. The average encryption speed for the proposed scheme is 1.015 Mbps.

Table 18. Encryption time, decryption time, and their added values of the proposed scheme for the Peppers image at various dimensions.

Image Dimensions	t_{Enc} [s]	t_{Dec} [s]	t_{Add} [s]
64 × 64	0.094082	0.13474	0.228822
128 × 128	0.377145	0.515267	0.892412
256 × 256	1.52554	2.05997	3.5855
512 × 512	6.05749	8.24135	14.2988
1024 × 1024	24.8238	34.2197	59.0435

Table 19. Encryption time comparison for various schemes of the Lena image having dimensions 256 × 256.

Scheme	t_{Enc} [s]	Machine Specifications (CPU and RAM)
Proposed scheme	1.42545	2.9 GHz Intel [®] Core [™] i9, 32 GB
[39]	2.7236	2.7 GHz Intel [®] Core [™] i7, 8 GB
[66]	3.45	N/A
[69]	1.1168	3.4 GHz Intel [®] Core [™] i7, 8 GB
[70]	1.112	3.4 GHz Intel [®] Core [™] i3, 4 GB
[71]	4.98	2.5 GHz AMD [®] , 4 GB

3.13. S-Box Performance Analysis

As a stable component, which is almost always at the core of any image encryption technique, also, as it holds the responsibility towards implementing Shannon's property of confusion in a cryptosystem, an S-box should be evaluated in isolation from the total perspective of the whole encryption process. There are five tests most commonly performed in order to evaluate the confusion capability of an S-box. The first test is nonlinearity [72], which represents the measure of how many bits in the truth table of a Boolean function need to be changed in order to approach the closest affine function (optimal value of 120, with most commonly achieved of 112). The second test is linear approximation probability (LAP) [73], which identifies the probability of bias for a given S-box (optimal value of 0.0625). Third test is a differential approximation probability (DAP) [74], which is a technique that examines the impact of specific variations in inputs and its effect on encrypted output (optimal value of 0.0156). Fourth, are the bit independence criterion (BIC) [75], which evaluates the relation between encryption procedures and the repeated patterns in the encrypted output (optimal value of 112). Finally, strict avalanche criterion (SAC) [75], which calculates the rate of change in the encrypted output with respect to the change in the input on a bit by bit level (optimal value of 0.5).

In commonly used methods for S-box generation, the procedure followed in the process of S-box generation takes the evaluation methods into consideration. In this work, as demonstrated in Section 2.2, a PRNG approach is followed, which introduces some advantages alongside some disadvantages. The main advantage is that the overall encryption process requires more keys, increasing the overall key space, resulting in more resistance to attacks. On the other hand, as per the keys provided to the S-box generation process, the evaluation scores for each S-box generation scenario is not fixed. As per

that, the relation between various keys in the key space and encrypting strengths of the generated S-boxes is to be correlated, as a future work.

Evaluating an S-box generated using keys: $\sigma = 10$, $\beta = 8/3$, and $\rho = 28$ (shown in Table 3) utilizing the aforementioned evaluation methods results in the findings presented in Table 20. As the evaluations demonstrate, not all optimal values were met. More precisely, while nonlinearity and SAC showed near optimal scores, DAP scored average, and LAP was far from optimal. For better reference, Table 21 shows scores comparison with popular S-boxes in the literature, which represent acceptable scores overall. These shortcomings are a natural result of the mechanism we adopted for the S-box generation, which is completely random, with full disregard to major S-box design criteria, which aim at avoiding fixed points and short ring cycles [76,77]. On the other hand, they can be regarded as trade-offs to increasing the key space of the overall encryption process, due to the addition of the new three tunability parameters (σ , ρ and β of the Lorenz system). In other words, instead of having a fixed (well-performing) S-box as one of the encryption stages, a randomly-generated S-box will allow for increasing the number of encrypted images per a single input plain image. In the case of the proposed work, the factor of increase in encrypted images per a single input plain image is 2^{19} , as thoroughly discussed in Section 3.10.

Table 20. Evaluation for S-box generated using keys: $\sigma = 10$, $\beta = 8/3$, and $\rho = 28$ (shown in Table 3).

Evaluation Method	Optimal Score	Scored of Proposed Scheme
Nonlinearity	112	106
SAC	0.5	0.5019
BIC	112	112
LAP	0.0625	0.1328
DAP	0.0156	0.0391

Table 21. Comparison between proposed S-box ($\sigma = 10$, $\beta = 8/3$, $\rho = 28$) and those provided in the literature.

S-Box	Nonlinearity	SAC	BIC	LAP	DAP
Proposed	106	0.5019	112	0.1328	0.0391
AES [13]	112	0.5058	112	0.0625	0.0156
Khan et al. [17]	111	0.5036	110	0.0781	0.0234
APA [78]	112	0.4987	112	0.0625	0.0156
Gray [79]	112	0.505	111.46	0.0664	0.0156
Zahid et al. [80]	107	0.497	103.5	0.1560	0.0390
Farwa et al. [81]	103.5	0.5065	103.3	0.1328	0.0468
Aboytes et al. [82]	112	0.4998	112	0.0625	0.0156
Hayat et al. [83]	100	0.5007	104.1	0.0390	0.1250
Nasir et al., (S4) [84]	112	0.5	112	0.0625	0.0156

4. Conclusions and Future Works

This research work proposed a 3-stage image encryption scheme. DNA encoding, along with a tan variant of the logistic map, were made use of in the first stage of encryption. An S-box based on the numerical solution of the Lorenz differential equations and a linear descent algorithm was developed for the second stage, and utilized to carry out bit confusion. Finally, the logistic map in its original form was utilized to produce an encryption key for the third stage of encryption. Performance of the proposed image cryptosystem was evaluated utilizing commonly used metrics from the literature, as well as newly adopted ones. The computed values reflect a secure and robust cryptosystem that is resistant to visual, statistical, entropy, differential, known plain text and brute-force attacks. This has been validated through comparison with the performance of counterpart schemes from the literature. The proposed cryptosystem was shown to exhibit comparable performance to them, if not superior at times. Future works could: (a) attempt to replace the

utilized 1D chaotic function with a higher-dimensional one. While this would invariably improve the security even more, it is expected that it might affect the computational complexity, and thus the execution time of the algorithm; and (b) optimize the generated S-box design, to achieve better values in all of the evaluation metrics. While the proposed S-box design did not score optimal values in all metrics, this was compensated in the proposed cryptosystem as a whole, due to the other two encryption stages. Furthermore, a future work could replace the utilization of the Lorenz system for the construction of the S-box. The reason behind such a recommendation is that the Lorenz system is a continuous one, with its computation heavily depending on the numerical method adopted in solving it. A better choice would be any discrete chaotic function.

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Abbreviations

The following abbreviations are used in this manuscript:

DNA	Deoxyribonucleic acid
MAE	Maximum absolute error
MSE	Mean Square Error
NIST	National Institute of Standards and Technology
NPCR	Number of Pixel Changing Ratio
PSNR	Peak Signal-to-Noise Ratio
UACI	Unified Averaged Change Intensity

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