

Assessment of Genetic Algorithm using Mendel's Principles

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ABSTRACT

Genetic Algorithms (GAs) takes huge search spaces and looks for optimal combinations of things' solution. It is the robust solution rated against fitness criteria so it avoids local optima and searches for global fitness. But still there are some cases and situations where solutions are provided on the basis of local optimal values which defeat the whole process of using Gas for optimization. Mutation operator in GA maintains uniqueness in the newly generated data values/chromosomes by modifying a part of data based on some predefined probability. The purpose of this paper is to optimize the conventional GA to prevent the population from stagnating at any local optima. In this paper, Mendelian inheritance theory describes the rules by which the genes are inherited by the offspring. New offspring always carries the properties of both of its parents but only one parent's properties are dominant. But it doesn't mean that the other parent's properties are suppressed, they are just hidden. After inter-off springs' mating one of the new offspring may have the properties of second parent as well. A modified version of this theory has been used in the paper to improve the problem of local optima. The proposed optimized algorithm is further evaluated with the help of De Jong's test functions. The results are compared with the conventional GA results and the performance is analysed.

Keywords: Genetic algorithm, Mendelian principle, Optimization, Population, Genetics.

1. Introduction

Genetic Algorithms are intense broadly useful streamlining look calculations based upon the standards of development saw in nature. The GA, initially created by Holland (1975), reproduces a developmental procedure of a living animal types, utilizing commonplace natural hereditary qualities operations, for example, "determination", "change" and "hybrid". Be that as it may, the structure of GA does not have an attractive doctrinal sense the same number of issues of hereditary qualities are not reflected in the calculation. The structure of chromosomes, creation of populaces, new eras, and hybrids, and so on are coded with arbitrary schedules, bringing about a non-natural presentation. In spite of the fact that the establishments of GA rely on upon populace hereditary qualities, traditional GA is fairly unsuitable as a delicate approach. The functional issues are frequently portrayed by a few non-commensurable and contending measures of execution or goals, with various confinements forced on the choice factors. The decision of an appropriate bargain arrangement from all non-second rate choices is not just issue subordinate; it for the most part depends likewise on the subjective inclinations of a choice specialist. Therefore, the last answer for the issue is the consequence of both an enhancement procedure and a choice procedure [1]. In the recent past, lot of research work has been done by several researchers for improving GA's local optimization issue using Mendel's principles but in this paper several changes have been proposed in the base theory (as described in the later sections) that is providing the better solution over the research happened till now.

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2. RELATED WORK

Yi Chen et.al [1] proposed a hereditary calculation utilizing Mendel's guideline (Mendel-GA), in which the arbitrary task of alleles from guardians to offsprings is inferred by the Mendel hereditary administrator to take care of the trade rates assurance issue. The demonstrating of trade rates developments is a testing undertaking in worldwide fund. Other than the customary hereditary administrators of choice, hybrid, and transformation, Mendel's standards are incorporated, as an administrator in the hereditary calculation's advancement procedure. In the quantitative examination of trade rates assurance, the Mendel-GA looks at the conversion standard variances at the short-run skyline. In particular, the point is to return to the assurance of high-recurrence trade rates and inspect the contrasts between the technique for hereditary calculations and that of the conventional estimation strategies.

Meera Kapoor et.al [2], presented streamline/boost de Jong's function1 in GA utilizing diverse determination plans (like roulette wheel, arbitrary choice, best fit/elitist fit rank choice, competition choice). The paper executes De Jong's function1 i.e. circle demonstrate utilizing diverse determination strategies utilized as a part of Genetic calculation and makes an examination of them in light of the wellness estimations of capacity at various number of emphases. The encoding plan for this issue of capacity expansion is esteem/genuine encoding. Thus an alternate kind of hybrid technique is connected for intersection the chromosomes in the populace for delivering better offspring's. Lastly the transformation strategy utilized is additionally not quite the same as the standard ones. Uniform transformation is utilized while actualizing this calculation.

Ryouei Takahashi et.al [3] proposed another half breed strategy by joining Ant Colony Optimization (ACO) and Edge Assembly Crossover (EAX). Authors devised an iterative Extended Changing Crossover Operator (i-ECXO) to take care of the Travelling Salesman Problem (TSP). The proposed conspire effectively looks for the ideal arrangement while keeping up the assorted qualities of the cyclic ways making the populace. EAX frames A-B cycles and produces kids by then again changing guardian Ant's edge for parent B's edge that showed up on the A-B cycles. One of the improving strategies that recreate the swarm insight in ants' sustaining conduct is ACO. In this paper, the differing qualities of eras were entirely measured by the entropy H in TDGA.

Basima Hani Hasan et.al [4] utilizes hereditary calculations to comprehend TSP, 0/1-Knapsack issue and Shubert Function. Change is one of the imperative administrators of hereditary calculations since the kind of transformation utilized regularly effects affects the outcomes. The review demonstrates that inclusion transformation is the best suite for TSP, Boundary and non-uniform changes are the best to use for Shubert Function issue, yet for 0/1 backpack issue all transformation sorts utilized gave about a similar outcome.

Mehmed Celebi et.al [5], proposes another approach in view of two express standards of Mendel trials and Mendel's populace hereditary qualities for the hereditary calculation (GA). These principles are the isolation and free grouping of alleles, individually. This new approach has been reenacted for the advancement of certain test capacities. The doctrinal feeling of GA is thoughtfully enhanced by this approach utilizing a Mendelian structure. The new approach has been tried on five De Jong's test capacities and on one Rastrigin work.

Grigorios N. Beligiannis et.al [6], proposed an approach to improve the execution of the exemplary Genetic Algorithm. Restarting a Genetic Algorithm is connected keeping in mind the end goal to acquire better learning of the arrangement space of the issue. Another administrator of "addition" is acquainted so as with adventure the data that has as of now been gathered before the restarting strategy. A settled number of genomes are chosen and included into the new populace. These genomes embody all the helpful data assembled about the arrangement space till that era. The part of the "vector that is a blend of Ritz qualities" is played by the arrangement of genomes that is passed to the new era.

Carlos A. Coello [7] give a far reaching survey of the most mainstream transformative based ways to deal with multi-target advancement, giving a few bits of knowledge into their operations inquire about roots, a brief portrayal of their principle calculations, their focal points and hindrances, and conceivable scope of relevance. Moreover, some illustrative certifiable utilizations of every approach are additionally included.

3. MENDELIAN PRINCIPLE

Mendel's standards have constantly connected to the hereditarily repeating animal in a common habitat. Mendel gave an unequivocal administer to how the genotypes of the posterity can be anticipated from the genotypes of their folks, and he likewise settled models for how genotypes were identified with characteristics. Mendel's work is known to a great extent through his Experiments in Plant Hybridization distributed in 1865. It provided details regarding eight years of experimentation with the garden pea. Mendel settled on a few ponder decisions for his investigations which were critical in empowering one to deduce the laws of legacy in his arrangement of examinations, basically inspecting extremely straightforward, now called Mendelian forms of inheritance. By experimenting with pea plant breeding, Mendel developed three laws of inheritance that described the transmission of genetic traits.

Mendel's laws are as follows:

3.1. The Law of Dominance

A dominant trait is a trait whose appearance will always be seen in offspring. In other words, dominance describes the relationship between two alleles. If an individual inherits two different alleles from each of its two parents and the phenotype of only one allele is visible in the offspring, then that allele is said to be dominant. Mendel's law of dominance states that if one parent has two copies of allele A — the dominant allele and the second parent has two copies of allele a— the recessive allele, then the offspring will inherit an Aa genotype and display the dominant phenotype.

To illustrate this consider a scenario where the pea shape trait of the pea plant is considered. One form of the gene (allele) codes for round and represented as "R", and the other allele codes for wrinkled represented as "r". Then the possible genotypes are:

Table 1
Possible Genotypes

<i>Genotype symbol</i>	<i>Genotype Vocab</i>	<i>Phenotype</i>
RR	Homozygous DOMINANT	Round
rr	Homozygous RECESSIVE	Wrinkled
Rr	heterozygous	Round

Considering one pea seed as pure round (RR) and other as pure wrinkled (rr):

Parent (P): RR X rr

The Punnet square for such combination will be as:

	R	R
r	Rr	Rr
r	Rr	Rr

The possible genotypes of the offspring (Generation 1) obtained from the parents is shown by punnet square. It shows that all the offsprings will have 100% Rr Genotype and the dominant trait 100% round.

3.2. The law of Segregation

A parent may have two distinct alleles for a certain gene, each on one copy of a given chromosome. Mendel's second law, the law of segregation, states that these two alleles will be separated from each other during meiosis. Specifically, in the second of the two cell divisions of meiosis the two copies of each chromosome will be separated from each other, causing the two distinct alleles located on those chromosomes to segregate from one another.

The above scenario can be further continued in segregation as:

$$Rr \rightarrow R + r$$

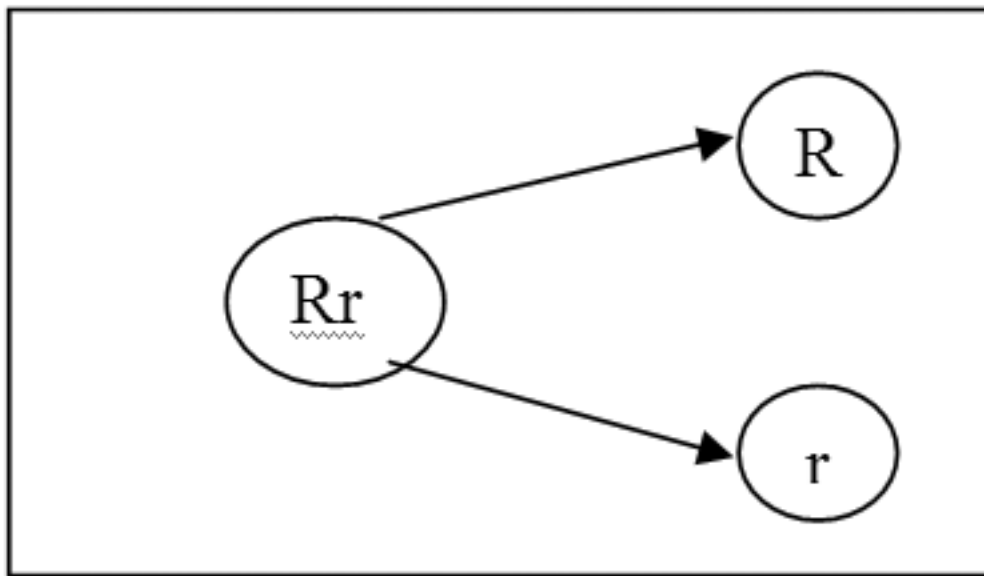


Figure 1: Heterozygous individual segregation

In this case, second generation off springs is generated from the first generation.

The Punnet square for the combination is:

	R	r
R	RR	Rr
r	Rr	rr

The P-square shows that now the genotypes are: 25% RR, 50% Rr and 25% rr. The 3 boxes out of 4 produce an offspring with dominant trait while one is producing with recessive trait.

3.3. The Law of Independent Assortment

Mendel's third law, the law of independent assortment, states that the way an allele pair gets segregated into two daughter cells during the second division of meiosis has no effect on how any other allele pair gets segregated. In other words, the traits inherited through one gene will be inherited independently of the traits

inherited through another gene because the genes reside on different chromosomes that are independently assorted into daughter cells during meiosis.

To illustrate this, more than one trait is considered in the parent pea plant.

Consider that the parent pea plant contain two different traits: 1. Seed Texture (round or wrinkled)

2. Pod color (green or yellow)

The genotypes are represented as:

R- Dominant allele for round seeds

r- Recessive allele for wrinkled seeds

G- Dominant allele for green pod

g- Recessive allele for yellow pod

The genotypes for parent pea plant will be:

$RrGg \times RrGg$

Table 2
The Punnet square for the new combination

	RG	Rg	rG	Rg
RG	RRGG	RRGg	RrGG	RrGg
Rg	RRGg	RRgg	RrGg	Rrgg
rG	RrGG	RrGg	rrGG	rrGg
rg	RrGg	Rrgg	rrGg	Rrgg

9/16 boxes (offspring) demonstrate predominant phenotype for both properties(round and green), 3/16 indicate dominance for first characteristic and latent for second (round and yellow), 3/16 also demonstrate recessive phenotype for first attribute and dominant structure for second (wrinkled and green), and 1/16 indicate recessive type of both characteristics (wrinkled and yellow). This likelihood can be condensed as 9:3:3:1 phenotypic proportion.

In any case, each posterity has a phenotype that is proportional to its genotype. Consequences of the hybrid for a person who have n-quality sets give 4^n conceivable outcomes. Here, n characterizes the quantity of quality sets.

Table 3
Genotype and phenotype numbers in Mendelian crossover

<i>Heterozygous gene pairs</i>	<i>Gamete Numbers</i>	<i>Genotype Numbers</i>	<i>Phenotype Numbers</i>
N	2^n	4^n	2^n
1	2	4	2
2	4	16	4
3	8	64	8
.....
N	4096	16,777,216	4096

4. PROBLEM IDENTIFICATION AND PROPOSED SCHEME

Genetic Algorithm is a sort of inquiry streamlining calculation in view of the hypothesis of advancement and the hereditary transformation hypothesis of Mendel. Regardless of being a hot research point in various

application ranges, the hypothesis of hereditary calculation is still blemished and there is much extent of advancement. Hereditary calculations involve the essential operation of encoding, determination, hybrid and change.

Different hereditary controls and change recreate the fundamental qualities of organic development from various points in Genetic calculations and are an imperative measure in the avoidance of neighborhood joining. Numerous analysts have enhanced the hereditary calculation from alternate points of view.

A genetic algorithm consists of three principal operations:

- A. The selection operation picks some of offspring to survive by as per some predefined rules. The quantity of the population is then under great control.
- B. The crossover operation creates the offspring from two picked mates in the whole population after selection, by exchanging a few bits of the two people. The offspring will then acquire a few attributes of their parents.
- C. The mutation operation generates the offspring by randomly changing one or several bits of individuals the offspring may then possess different characteristics from their ascendants. Mutation can then avoid local search in the searching space and increase the probability of finding the global optimum.

The primary impediment of GA's is that while taking care of ideal issues with unadulterated constant factors they are less effective than the inclination based calculations, as demonstrated by the way that significantly more emphases are required for joining. In this paper, Mendel's principles are incorporated with the customary hereditary administrators of determination, hybrid, and change, as an administrator in the hereditary calculation's development procedure. The Mendel operator is applied on the population generation step as it is an initial part of the genetic search. Like conventional GA, initial population is generated randomly and a data pool of certain variables is generated randomly. After population generation gametes are selected for a genotype and dominance and recessiveness are obtained. Further Mendel operator is applied to the obtained genotypes.

The working of the proposed approach can be understood by following flowchart:

Further the optimized Mendel-GA is evaluated with De Jong's Test functions. De Jong capacity is prevalent in genetic algorithm writing. It is otherwise called sphere model. It is persistent, raised and uni modal.

De Jong 1 capacity is one of the least complex test capacities. It has a consistent and convex format. The worldwide ideal esteem in the De Jong 1 capacity is zero and is free from the quantity of factors.

It can be defined as:

$$f(x) = \sum_{i=1}^n x_i^2 (i)$$

where $-5.12 \leq x_i \leq 5.12$, Global minimum is at: $f(x)=0$, $x(i)=0$, where, $i = 1:n$.

De Jong 2 function is a classic optimization function, and is known as a 'banana function' or the Rosenbrock's function. The global optimum point is in a deep parabolic shaped valley, its value is zero and is independent of the number of variables. As it is hard to find the minimum point in the valley, this function is used for the performance tests.

$$f(x) = \sum_{i=1}^n 100 \cdot (x_{i+1} - x_i^2)^2 + (1 - x_i)^2$$

where $-2.048 \leq x_i \leq 2.048$, Global minima is at $f(x) = 0$, $x_i = 1$, $i = 1:n$

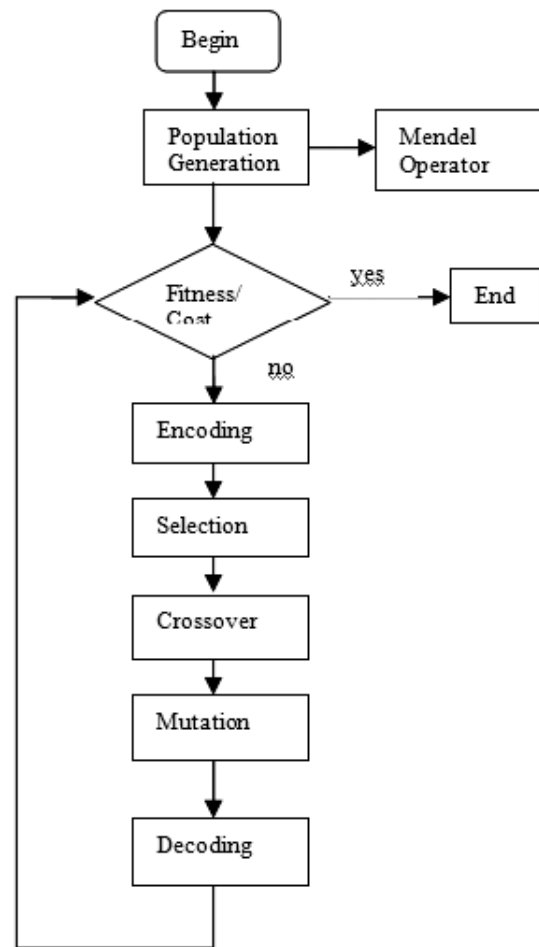


Figure 2: Proposed mechanism

5. SIMULATION ANALYSIS

The proposed scheme has been simulated in MATLAB and the results have been presented in this section. The optimized Mendel GA is simulated and is compared with the conventional GA to show the performance differences. The optimized GA is evaluated with De Jong test functions.

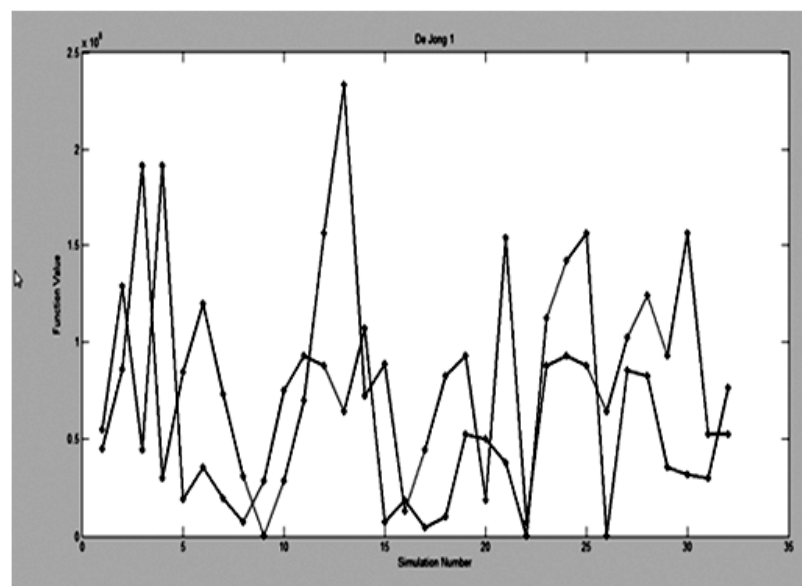


Figure 3: Fitness value of the optimized Mendel GA evaluated with De Jong test function

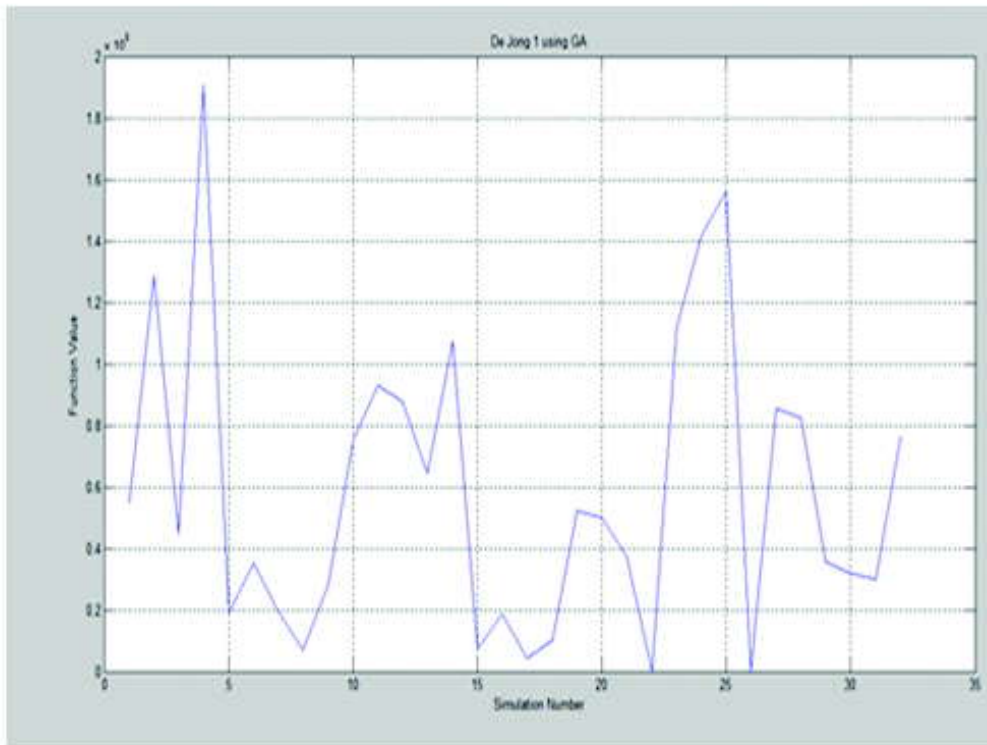


Figure 4: Traditional GA evaluated with De Jong Test Function

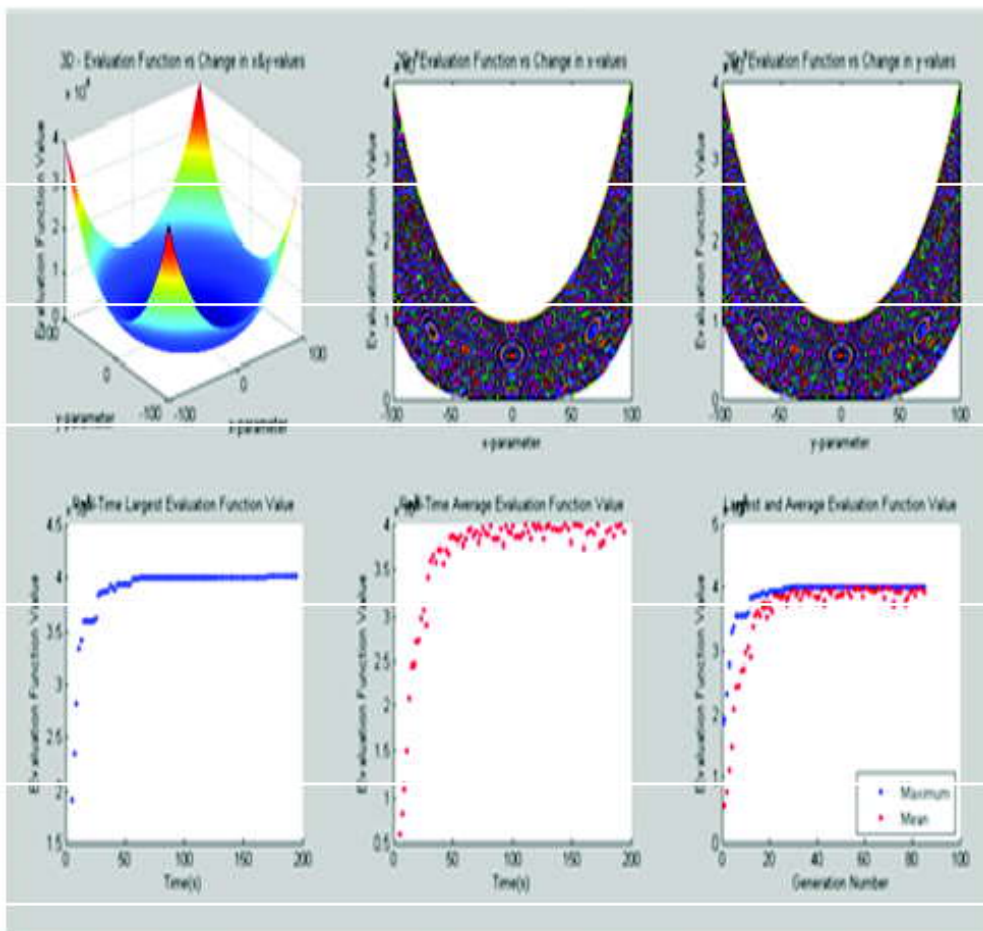


Figure 5: Optimized Mendel GA shows better performance than the conventional GA

Table 4
Conventional GA

<i>Population</i>	<i>Fitness function</i>	<i>Iteration</i>
8	$1.9087 * 10^8$	50
16	$1.9087 * 10^8$	37
32	$1.9087 * 10^8$	36
64	$3.4549 * 10^8$	38
128	$3.4549 * 10^8$	38

The table shows the fitness function and iterations in conventional GA according to the population size.

Table 5
GA with Mendel

<i>Population</i>	<i>Fitness function</i>	<i>Iteration</i>
8	$1.9686 * 10^8$	275
16	$1.9087 * 10^8$	390
32	$3.0541 * 10^8$	324
64	$3.5201 * 10^8$	696
128	$3.5542 * 10^8$	410

This table shows the iterations and fitness function values with respect to the population size in optimized Mendel GA.

6. CONCLUSION

Genetic algorithm (GA) is a extensive explore substance in the application domain and the academic circle at present. This paper elaborates the enhancement of impulsive meeting in GA used for optimizing multimodal mathematical problems. Mutation is the standard function in Genetic Algorithm (GA) for enhancing the degree of inhabitants range, but it is proved that it is not efficient often, mostly in traditional GA. To deal with this issue, Mendelian operator has been applied after the inhabitant's generation phase. The optimized Mendel GA has been evaluated with De Jong's test functions and the results are compared with the conventional GA. The results show that the optimized GA shows better performance than the conventional one. In proposed optimized GA mechanism execution rate is fast as compare to existing mechanism also proposed mechanism can be applied to real applications with high throughput and less execution time. In future we will try to apply proposed mechanism in some real life problems like TSP, QARP and VRP etc.

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