

CENTRAL WASHINGTON UNIVERSITY

COMPUTATIONAL INTELLIGENCE

WINTER 2019

Project 3 Report

Author:

Hermann YEPDJIO

Professor:

Dr. Razvan ANDONIE

February 20, 2019



Contents

1	Problem Description	2
2	Implementation	2
2.1	Input Data	2
2.2	Selection	2
2.3	Crossover	3
2.4	Mutation	3
3	Experimentation	3
3.1	Process	3
3.2	results	4
4	Conclusion	7

1 Problem Description

The genetic algorithm is a technique used for solving optimization problems. It follows the same process that drives biological evolution. It starts with an initial population of solutions which it modifies at each step to produce a new generation of the population usually better than the previous generation. The modification of the population is done through processes such as cross-overs and mutations.

The purpose of this project was to implement a genetic algorithm able to generate a magic square of size n . A magic square of size n is an $n \times n$ matrix which contains unique numbers from 1 to n^2 and with each column, row and diagonal having the same sum. After implementing the algorithm, we experimented using different sizes for the population and different mutation rates. The details of the implementation as well as the results of the experimentation are discussed below.

2 Implementation

2.1 Input Data

The program takes the following inputs from the user before the search can start:

- an integer greater than 3 for the size of the magic square (3 for 3×3 , 4 for 4×4 etc ...),
- an integer for the size of the initial population (the size of the population remains constant across generations),
- a float number between 0 and 1 for the mutation rate

2.2 Selection

For the crossover operation, a mating pool of half the size of the population is created and filled with chromosomes from the original population selected based on some probabilities calculated using their fitness.

2.3 Crossover

Since a magic square requires all numbers of the matrix to be unique, simple crossover techniques such as taking halves of the genes of 2 parents and merge them together to produce an offspring could not be considered. Therefore, we used an approach proposed on the paper Genetic Algorithm Solution of the TSP Avoiding Special Crossover and Mutation written by Göktürk Üçoluk. The approach basically follows the following steps to produce 2 offspring:

- calculate the inversion sequence for each parent,
- generate the children arrays,
- transforming the children arrays back to permutation representation.

The paper mentioned above provides more details on how those steps are performed.

2.4 Mutation

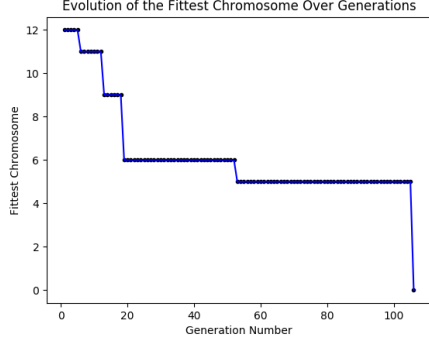
After the crossover is performed, the program randomly decides whether to apply a mutation or not based on the mutation rate provided by the user. After the mutation stage is over (no matter if genes were mutated or not) the program looks for the least two fitted chromosomes in the original population and replace them with the new children.

3 Experimentation

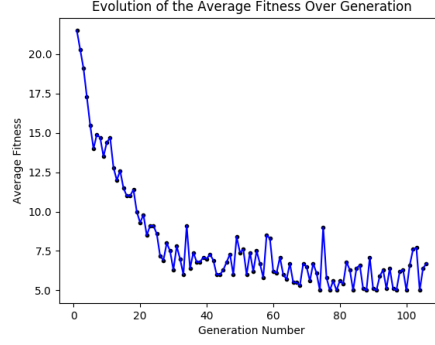
3.1 Process

During this stage, we experimented generating magic squares of different sizes using different mutation rates and population sizes. The results were recorded and are described in the next subsection.

3.2 results

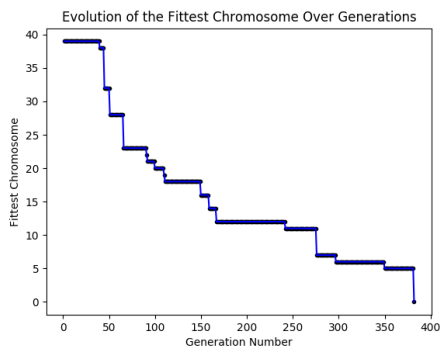


(a) 3*3 Magic Square Evolution of Fittest Chromosome

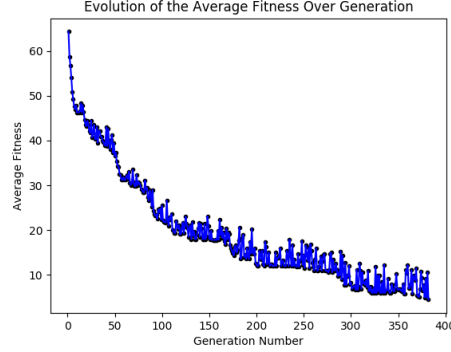


(b) 3*3 Magic Square Evolution of the Average Fitness

Figure 1: 3*3 Magic Square. pop_size = 10, mutation rate = 0.5



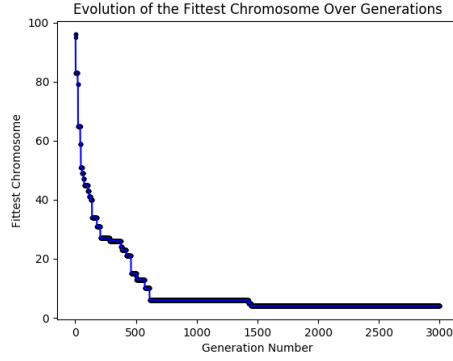
(a) 4*4 Magic Square Evolution of Fittest Chromosome



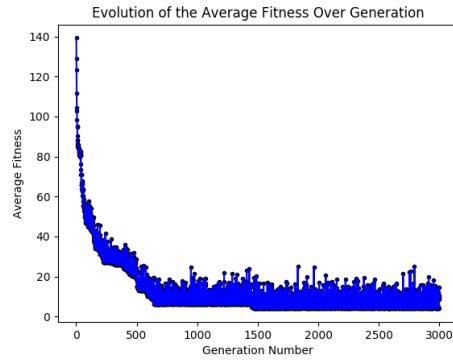
(b) 4*4 Magic Square Evolution of the Average Fitness

Figure 2: 4*4 Magic Square. pop_size = 10, mutation rate = 0.5

In Figures 1 and 2 above we can see that a 3*3 and a 4*4 magic squares were found (fitness = 0) at *generation* $\simeq 100$ for the first and *generation* $\simeq 400$ for the second. We can also see that both the fitness of the fittest chromosome and the average fitness of the population improve (get closer to 0) over the generations.



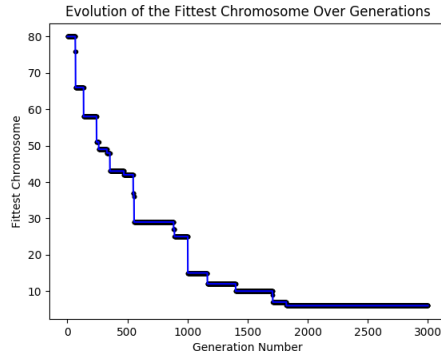
(a) 5*5 Magic Square Evolution of Fittest Chromosome



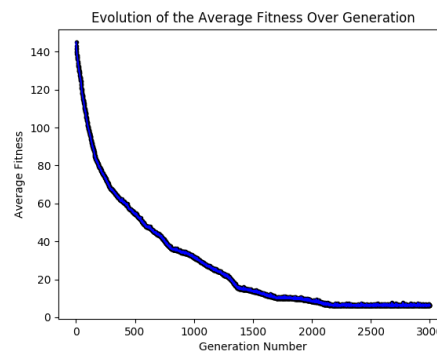
(b) 5*5 Magic Square Evolution of the Average Fitness

Figure 3: 5*5 Magic Square. pop_size = 10, mutation rate = 0.5

In Figure 3 we can see the same convergence of the average fitness and fitness of the fittest chromosome observed in Figures 1 and 2 even though a solution is not found before reaching generation 3000.



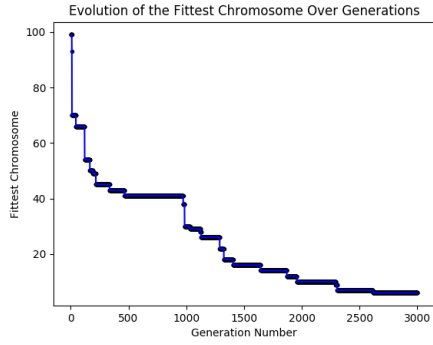
(a) 5*5 Magic Square Evolution of Fittest Chromosome



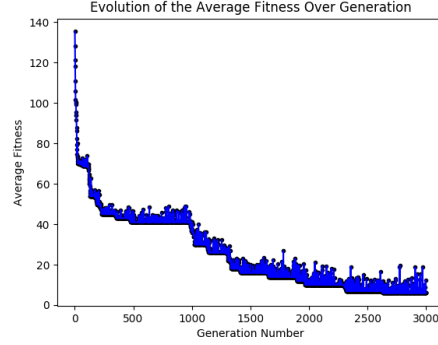
(b) 5*5 Magic Square Evolution of the Average Fitness

Figure 4: 5*5 Magic Square. pop_size = 100, mutation rate = 0.5

Figures 4 shows that the same convergence observed in Figure 3 happens when we increase the population size from 10 to 100. However, this convergence occurs more slowly than it does in Figure 3.



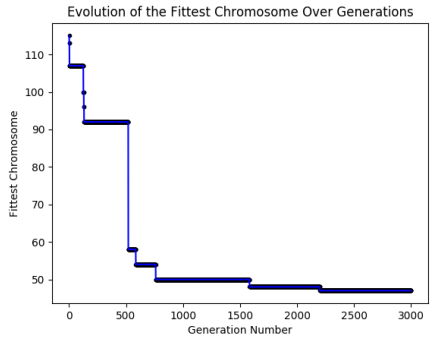
(a) 5*5 Magic Square Evolution of Fittest Chromosome



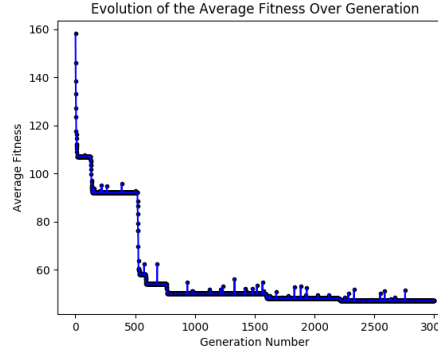
(b) 5*5 Magic Square Evolution of the Average Fitness

Figure 5: 5*5 Magic Square. pop_size = 10, mutation rate = 0.1

Figure 5 shows that reducing the mutation rate from 0.5 to 0.1 affect the speed at which the solutions converge and therefore also affect the optimal solution that can be found given a limited number of generations.



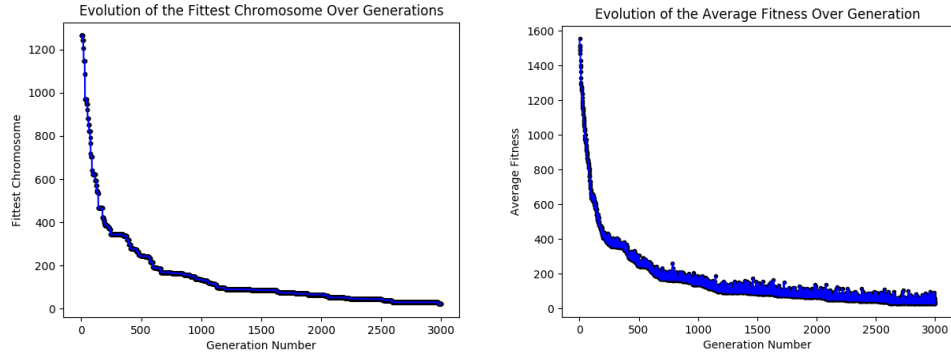
(a) 5*5 Magic Square Evolution of Fittest Chromosome



(b) 5*5 Magic Square Evolution of the Average Fitness

Figure 6: 5*5 Magic Square. pop_size = 10, mutation rate = 0.01

Figure 6 basically shows the same results observed in Figure 5.



(a) 10*10 Magic Square Evolution of Fittest Chromosome

(b) 10*10 Magic Square Evolution of the Average Fitness

Figure 7: 10*10 Magic Square. pop_size = 10, mutation rate = 0.5

Figure 7 shows how the solutions converge for a 10*10 magic square. We can see a huge improvement for the average fitness from above 1500 at generation 0 to less than 50 at generation 3000.

4 Conclusion

From the experimentation described above, we can conclude that choosing values for parameters such as the mutation rate or the population size depends on the type of problem we are trying to solve. Even though in general having a large population and a small mutation rate produces better results, in some cases like ours, that assumption does not hold true as we got better results using a small population size and a large mutation rate. However, figuring out what type of values we should use can only be done through experimenting with different values and observe which ones work better.