

# ARTIFICIAL INTELLIGENCE - CS F407

## GENETIC ALGORITHM FOR 3-SAT Problem

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A submission for **Assignment 1** made in partial fulfilment of course Artificial Intelligence - CS F407 by

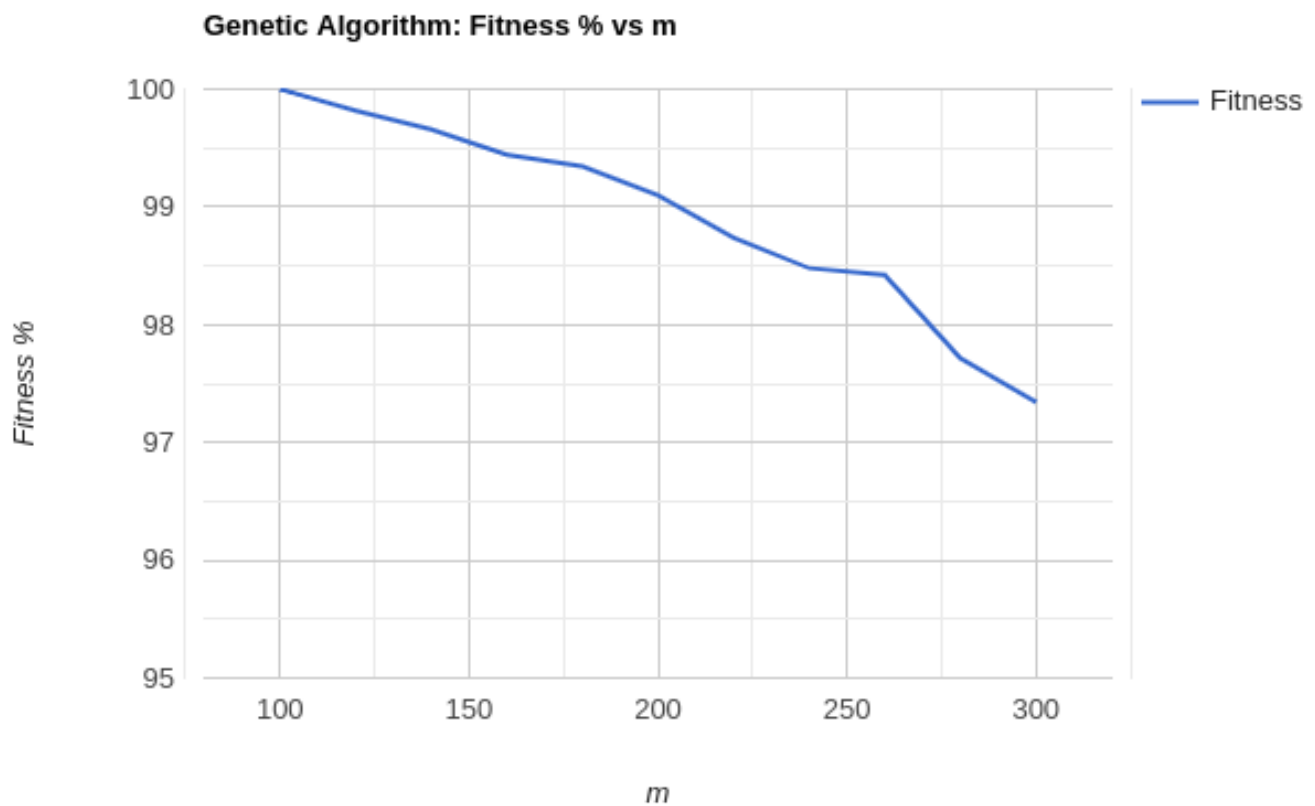
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## Q1: Graph of Fitness% value vs m, the number of clauses:

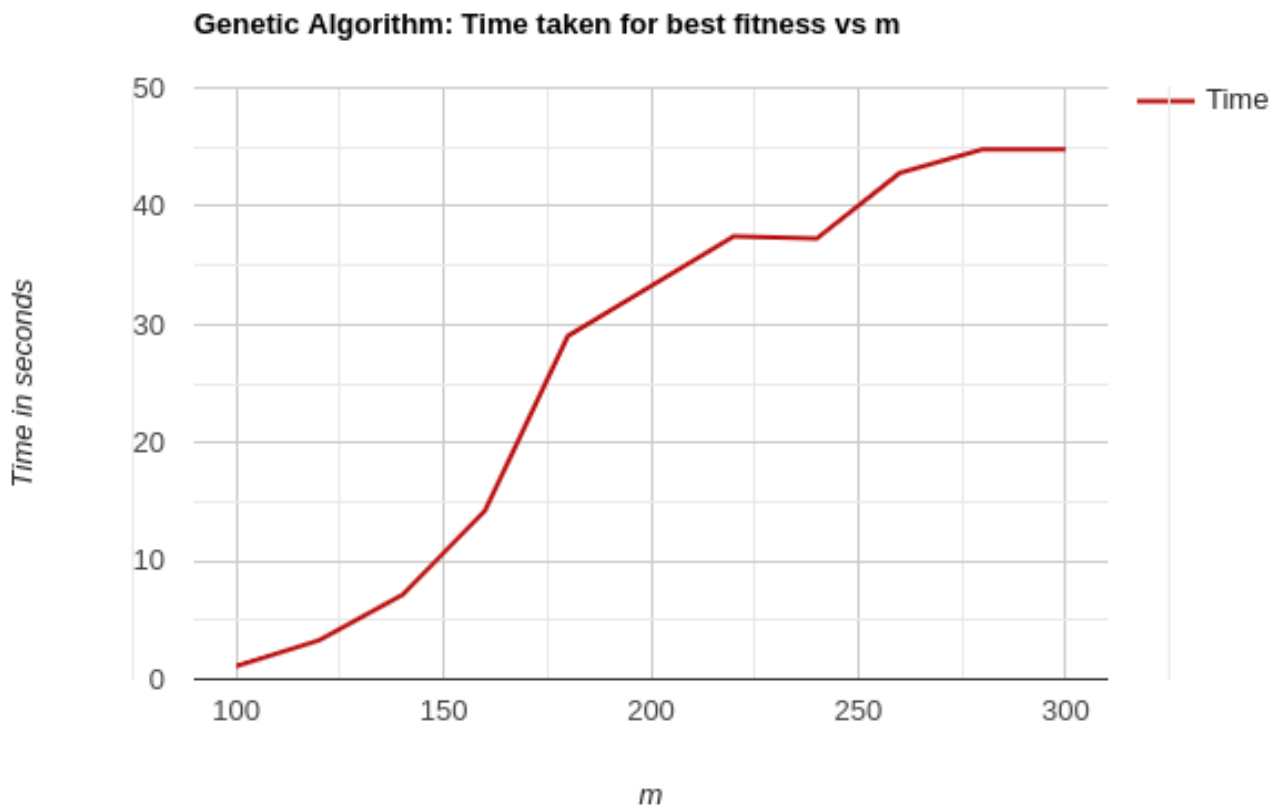


The fitness value is almost 100% for lower values of m, and it decreases to about 97.4% on average for m = 300

m	100	120	140	160	180	200	220	240	260	280	300
F %	100.0	99.81	99.66	99.44	99.24	99.10	98.74	98.48	98.42	97.71	97.34

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## Q2: Graph of Time taken for best Fitness% value vs m, the number of clauses:



The average time is greatly increased if GA gets stuck at a certain local optima. In this case, the average time is increased until the program is terminated or 45 seconds is completed.

m	100	120	140	160	180	200	220	240	260	280	300
Time	1.132	3.30	7.121	14.23	29.05	37.25	37.45	42.82	44.83	44.82	44.82

**Note:** I had auto-terminated the program completely at t = 44.8

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### Q3 How did I improve the GA Algorithm, and some failed approaches:

#### General Method, and population info:

The **members of the population** were taken to be Bit-Strings of length 50, since  $n$  remains constant = 50 throughout the experiment. If the Bit-String at index  $i == 0$ , it means that Variable  $\#(i + 1)$  is set to negative (negation), and vice versa (positive) for value at index  $i == 1$ .

The **fitness function** used is direct, and measures the number of Clauses satisfied, given an assignment of variables (a bit-string). The population size is kept constant = 20, for all values of  $m$ , as this was found to be most efficient.

With the version of GA described in the Textbook, I was getting a moderate final fitness function value: I Was able to satisfy CNF, 100 clause-sentences, however, the value wasn't hitting even close to 98-99% for 100-120 clauses, and the time taken was in general, slow.

#### I made the following changes which helped improve GA Algorithm:

##### **1. Selecting best 2 from the population at every iteration to form the next generation**

This method boosted up the running time of the algorithm by a great margin. Instead of assigning weights to each member on the basis of their fitness value, and hence choosing a member based on their weight (since the probability of a member with greater fitness to be picked is more), I picked the best 2 from the population to reproduce.

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This process first involved ordering the population on the basis of fitness function values, then, taking only a certain fraction (elitism rate) of population onto the next generation. From this new generation, best 2 are picked and their offspring are added to the current generation to fill up for those members “killed” during transfer

## **2. 2-point crossover:**

Instead of the general 1-point crossover, I introduced a better, 2-point crossover, for “reproducing” 2 members of the population. Here, we randomly select 2 points  $x, y$  ( $x$  and  $y$  represent the gene number, or the “variable” number in the chromosome) ( $x < y$ ) and exchange the genetic material of these 2 parents in between  $x$  and  $y$  Only. The 2 parents would produce 2 children, 1 of which was similar to Parent1 in regions not belonging to  $(x, y)$  and the other was similar to Parent2 in regions not belonging to  $(x, y)$

## **3. Extensive Multi-bit mutation:**

Upon experimentation, I found that extensive mutation helps to improve diversity of the population, so that if we proceed along a path that “looks correct”, but will not converge to a maximum value, i.e, the path would probably lead to a local optimum, we still add diversity by extensive mutation.

Initially, I mutated only 1 bit in the entire child produced after crossover. This didn't give great results in terms of final fitness%. So, I decided to mutate multiple bits (multi-bit mutation). The `max_mutation_size` hyperparameter is set, and the number of bits

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mutated for every child is a random number between 0 and `max_mutation_size`.

#### **4. Keeping mutated children:**

Instead of mutating children with some “probability”, I **always** mutated all children formed. This was done as mentioned in the previous point because mutation improves diversity and may help to steer us out of a path leading to local optima.

So, each call to `mutate()` would return 2 children: mutated versions of the children formed after crossover. This way, I retained important and “good” genetic material, and also improved diversity in the population

#### **5. Include a hyperparameter “elitism”:**

Elitism was defined as a hyperparameter (0 - 1 range) that defines how much fraction of the population (ordered descending by fitness function value) would be transferred onto the next generation. Hence,  $(1 - \text{elitism\_rate}) * \text{population\_size}$  members (the last members with low fitness function values) are removed at every iteration, hence eliminating unfavourable genes.

#### **6. Early-Stopping after a certain number of iterations:**

To improve the time taken by the program, I decided to terminate the program if the fitness doesn't improve over several generations (iterations). The value of this certain number of generations is a hyperparameter, “`iterations_to_terminate`”, and its value depends on `m`. Because for larger `m`, even after 800-1000 generations of constant

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fitness%, I found the fitness may improve. For smaller  $m$ , the value required was smaller, for example, for  $m \leq 120$ , I used the value 350.

## **Approaches that failed to improve GA:**

### 1. Increasing population size:

I tried to increase population size to large values like 100, 200, 400 etc. However, it didn't have a great impact on the fitness%, at the same time, it considerably slowed down the program, because there are multiple sorting operations involved in the program. After experimentation, it was best to choose population size = 20, this value gave best fitness% values, at the same time, it took a moderate-low amount of time

### 2. Modifying elitism rate:

- Making the elitism rate small:

For elitism rate  $< 0.4$ , the model failed to converge quickly

- Making the elitism rate big:

For elitism rate  $> 0.9$ , the model again, did not perform well

After multiple experiments, I found that elitism rate = 0.6-0.7 works best. I finally implemented GA with the rate = 0.7

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### 3. Different forms of mutation:

I tried various forms of mutation: Single point, Multiple point, FlipGA (iterate left to right, flipping only if fitness% would increase, then another iteration right to left), etc.

However, after multiple tries in each of these methods, a simple Random-multiple point mutation worked best. This was explained previously under “Extensive multi-bit mutation”

### 4. Different forms of crossover:

- Single point crossover: As mentioned in the Textbook
- 2-point crossover: As explained under the section with the same name. This is the final method implemented
- Copy-parent crossover: Only one child is created here. Those genes where  $\text{parent1}[i] == \text{parent2}[i]$  are directly copied, i.e  $\text{child}[i] = \text{parent1}[i]$ , and the other bits, where  $\text{parent1}[i] != \text{parent2}[i]$  are randomly set to be either 0 or 1

After multiple tries in all these methods, 2 point crossover worked best for me, in terms of both fitness% and efficiency in time



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#### Q4 Where GA might find it difficult to find a good solution:

- Genetic algorithms are highly sensitive to the initial population used. As I noticed during the experimentation, even for lower values of  $m$ , sometimes the algorithm gets stuck at  $((\text{number of clauses}) - 1)$ , and fails to converge correctly within 45 seconds. Most other times, it would solve and find a correct variable assignment within 2 seconds, hence the solution is random and dependent on the initial population used. Since the initial population is created in a random fashion, the GA may fail sometimes, or perform extremely slowly.
- Usually GA and different evolutionary algorithms, like particle swarm optimization (PSO), have a big stochastic component. This means that one needs to find a statistical convergent solution after multiple simulations/experiments.
- **Local Optima:** It is possible that GA gets stuck in a local optimum, and even mutations and crossovers are unable to remove the algorithm from the local optima. In this case, it usually gets stuck here for a very large number of iterations, sometimes more than 1000, and fails to converge to the global maximum. From my experience, a lot of local optima occur quite close to 100% satisfiability, even for lower values of  $m$ .

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**Q5 What does the above graphs tell you about the difficulty of satisfying a 3-CNF sentence in  $n$  variables. When does a 3-CNF sentence become difficult to satisfy?**

For given  $n = 50$ , GA was able to satisfy sentences with clauses upto around  $m \approx 200$ . After  $m = 220$ , GA was finding difficulties in converging quickly to 100% clauses satisfied. So, I believe, the difficulty of satisfying 3-CNF depends on the **ratio  $m/n$** : The ratio of clauses to variables. It is easier to satisfy 3-CNF if this ratio is smaller, and vice versa

So, if  $m/n$  is high, in general, if  $m/n > \sim(3 - 4)$ , it becomes more difficult to satisfy the 3-CNF problem