Submission By - Venkatesh Duraiarasan

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 ${\bf pseudo-code}$

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Algorithm 1: MaxSatGA Class Structure

ı class MaxSatGA

 $(clauses, num_vars, weights, k, population_size, pc, pm1, max_generations, time_budget);$

 ${\bf Input} \quad {\bf :} \ {\bf Clauses, \ number \ of \ variables, \ weights, \ number \ of \ constraints \ to}$

 $learn,\,population\,\,size,\,crossover\,\,probability,\,mutation\,\,probabilities,$

maximum generations, time budget, penalty, and debug flag

Output: Best solution, best fitness, and runtime

2 Initialize population using

 $initialize_population(population_size, num_vars);$

3 Optimize using optimize();

Algorithm 2: Optimize

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13

```
1 function optimize
    (clauses, num\_vars, weights, k, population\_size, pc, pm1, max\_generations, time\_budget);
   Input: Clauses, number of variables, weights, number of constraints to
            learn, population size, crossover probability, mutation probabilities,
            maximum generations, time budget, penalty, and debug flag
   Output: Best solution, best fitness, and runtime
 2 Initialize best_solution, best_fitness, and start_time;
 3 for each generation up to max_generations do
      if time elapsed exceeds time_budget then
         Break:
 5
      end
 6
      Evaluate population fitness using
       evaluate_population(population, clauses, weights, penalty);
      Update best solution and best fitness if necessary;
 8
      Create a new population through crossover and mutation;
      Update population;
10
      if debug mode then
         Print generation details;
12
      end
14 end
15 Return best solution, best fitness, and runtime;
```

Algorithm 3: Initialize Population

```
1 function initialize population (population size, num vars);
 Input: Population size and number of variables
```

Output: A random binary population matrix

2 Return a random binary matrix of size population $size \times num \ vars;$

Algorithm 4: Evaluate Population

```
1 function evaluate population (population, clauses, weights, penalty);
  Input: Population, clauses, weights, and penalty
  Output: List of fitness values for the population
2 forall individuals in population do
     Evaluate each individual using
      evaluate_individual(individual, clauses, weights, penalty);
4 end
5 Return a list of fitness values:
```

Algorithm 5: Evaluate Individual

Algorithm 6: Crowding Selection

```
function crowding_selection (fitness_values, population_size);

Input: Fitness values and population size

Output: Selected individuals

Initialize an empty list selected;

for i = 1 to population_size/2 do

Randomly select two parents p1 and p2;

Choose the parent with higher fitness and add it to selected;

end

Return selected individuals;
```

Algorithm 7: Clause Crossover

function clause_crossover (parent1, parent2, num_vars);
 Input : Two parent individuals and number of variables
 Output: Child individual

- 2 Convert parents to strings;
- 3 Randomly select a crossover point;
- 4 Create a child by combining parts of parents;
- 5 Convert child back to a binary array;
- 6 Return child;

Algorithm 8: Mutate Hardness

 ${\tt 1} \ \ {\tt function} \ \ {\tt mutate_hardness} \ \ (individual, pm1, k);$

Input: Individual, mutation probability, and number of constraints

Output: Mutated individual

- ${f 2}$ if $random\ probability < pm1$ then
- Randomly select an index idx;
- 4 Flip the bit at idx in the individual;
- 5 end
- 6 Return individual;

Exercise 5. Experimental Analysis of Essential Parameters in a MAXSAT Algorithm

The parameters chosen for analysis were population size, crossover probability, and mutation probability, as these are crucial in genetic algorithms for MAXSAT problems.

Experimental Setup Problem Instances: A subset of the smallest instances from the MSE17 benchmarks was selected to ensure computational feasibility.

- scpcyc06_maxsat.wcnf
- t4pm3-6666.spn.wcnf

Parameter Settings:

• **Population Sizes:** [10, 25, 50, 75, 100]

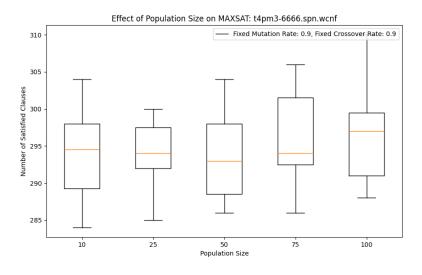
• Mutation Rates: [0.1, 0.25, 0.5, 0.75, 0.9]

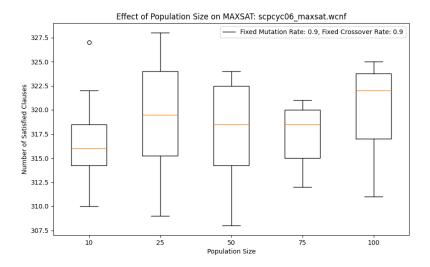
• Crossover Rates: [0.1, 0.25, 0.5, 0.75, 0.9]

• Timeout: 6 seconds per repetition

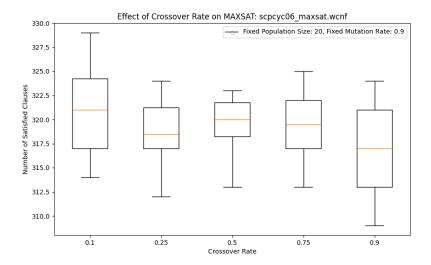
• Repetitions: 100 runs per parameter setting and problem instance

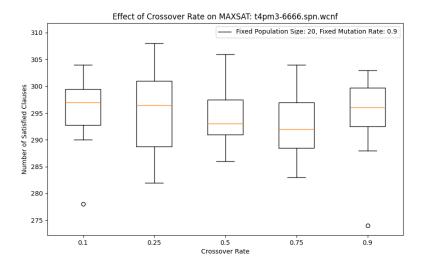
Results & Analysis Population Size: Larger population sizes improved solution quality by enhancing search space diversity. Increasing from 10 to 100 significantly improved results but increased computational time.





Crossover Probability: A probability of 0.1 to 0.25 balanced exploration and exploitation effectively. Higher values (e.g., 0.9) led to instability due to excessive genetic variation.





Mutation Probability: Lower rates (e.g., 0.1) limited diversity, while higher rates (e.g., 0.9) caused population instability.

