

Open Ended Lab Report

Artificial Intelligence

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N-Queens Problem Using Genetic Algorithms

Introduction

The **N-Queens Problem** is a classic combinatorial problem that involves placing N queens on an $N \times N$ chessboard so that no two queens attack each other. A queen can attack another queen if they are on the same row, column, or diagonal.

This project utilizes the **Genetic Algorithm (GA)**, a heuristic search and optimization technique inspired by natural selection, to find solutions to the N-Queens problem. The implementation includes a graphical user interface (GUI) for visualizing solutions and navigating through them interactively.

Objective

To develop a program that:

1. Solves the N-Queens problem using the **Genetic Algorithm**.
2. Provides an interactive GUI for inputting the board size ($N \times N$) and visualizing solutions.

Genetic Algorithm Overview

The **Genetic Algorithm (GA)** is a metaheuristic search algorithm that mimics the process of natural evolution. It operates using the following principles:

- **Population:** A set of candidate solutions (chromosomes).
- **Fitness Function:** Evaluates how "fit" a solution is.
- **Selection:** Chooses the best solutions for reproduction.
- **Crossover:** Combines parts of two parent solutions to produce offspring.
- **Mutation:** Randomly alters a solution to introduce variety.

Problem Representation

Chromosome Encoding

Each chromosome represents a potential solution to the N-Queens problem:

- A chromosome is a permutation of numbers, where the i -th value represents the column position of the queen in row i .
- Example: `[0, 4, 7, 5, 2, 6, 1, 3]` means:
 - Queen in row 0 is at column 0.
 - Queen in row 1 is at column 4, and so on.

Implementation

Key Functions in the Code

1. Generating Initial Population

```
def generate_initial_population(self):
    return [random.sample(range(self.N), self.N) for _ in range(self.population_size)]
```

- **Purpose:** Creates an initial random population of potential solutions.
- **How It Works:** Each solution is a random permutation of numbers from 0 to $N-1$.

2. Fitness Function

```
def fitness(self, chromosome):
    non_attacking = 0
    for i in range(self.N):
        for j in range(i + 1, self.N):
            if abs(chromosome[i] - chromosome[j]) != abs(i - j): # Check diagonal attack
                non_attacking += 1
    return non_attacking
```

- **Purpose:** Measures the quality of a solution by counting non-attacking queen pairs.
- **Key Logic:**
 - Checks for diagonal conflicts using the condition $|x_1 - x_2| \neq |y_1 - y_2|$
 - Higher fitness means fewer attacks.

3. Selection

```
def selection(self):
    weighted_population = [(self.fitness(ch), ch) for ch in self.population]
    weighted_population.sort(reverse=True, key=lambda x: x[0])
    return [ch for _, ch in weighted_population[:self.population_size // 2]]
```

- **Purpose:** Selects the top 50% of the population based on fitness for reproduction.
- **How It Works:**
 - Sorts the population by fitness in descending order.
 - Retains the most fit individuals for crossover.

4. Crossover

```
def crossover(self, parent1, parent2):
    split = random.randint(1, self.N - 2)
    child = parent1[:split] + [g for g in parent2 if g not in parent1[:split]]
    return child
```

- **Purpose:** Combines genetic material from two parents to create a new solution (child).
- **How It Works:**
 - Chooses a random split point.
 - The child inherits the first part of genes from one parent and fills the rest with genes from the other parent, avoiding duplicates.

5. Mutation

```
def mutate(self, chromosome):
    if random.random() < self.mutation_rate:
        i, j = random.sample(range(self.N), 2)
        chromosome[i], chromosome[j] = chromosome[j], chromosome[i]
    return chromosome
```

- **Purpose:** Introduces random changes to a chromosome to maintain diversity in the population.
- **How It Works:**
 - Swaps two random positions in the chromosome with a probability defined by `mutation_rate`.

6. Evolution

```
def evolve(self):
    new_population = []
    selected = self.selection()
    for _ in range(self.population_size):
        parent1, parent2 = random.sample(selected, 2)
        child = self.crossover(parent1, parent2)
        child = self.mutate(child)
        new_population.append(child)
    self.population = new_population
```

- **Purpose:** Evolves the population to the next generation.

- **Key Steps:**

1. Selects the best individuals.
2. Produces new solutions via crossover and mutation.
3. Replaces the old population with the new one.

7. Finding Solutions

```
def find_all_solutions(self):
    seen = set()
    for _ in range(self.generations):
        for chromosome in self.population:
            if self.fitness(chromosome) == self.N * (self.N - 1) // 2: # Max fitness
                solution = tuple(chromosome)
                if solution not in seen:
                    self.solutions.append(solution)
                    seen.add(solution)
    self.evolve()
```

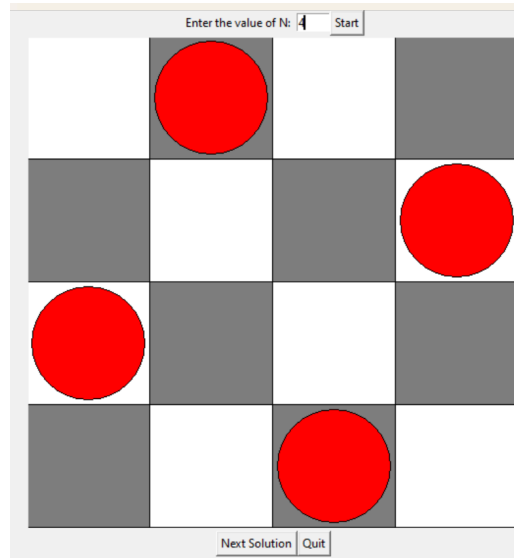
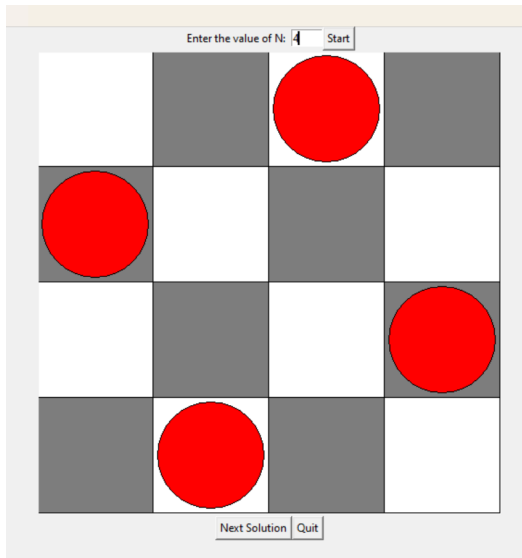
- **Purpose:** Finds all unique solutions by repeatedly evolving the population.
- **How It Works:**
 - Checks if a solution has maximum fitness.
 - Stores unique solutions in the `solutions` list.

User Interface (UI)

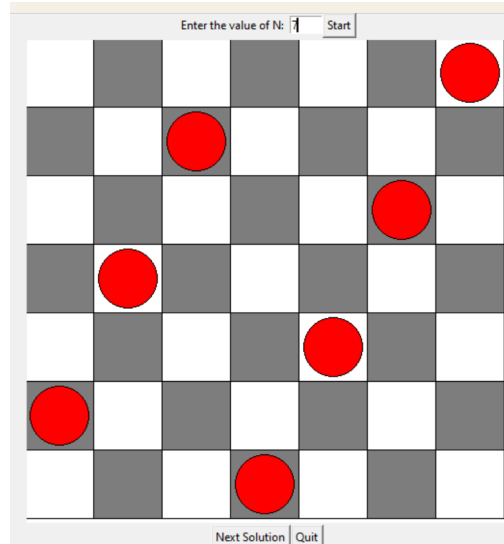
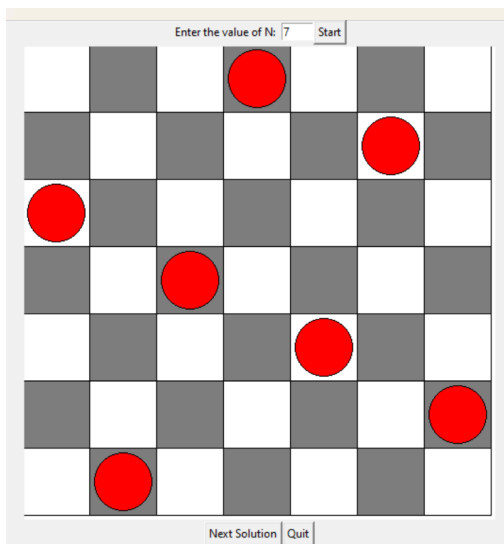
The **Tkinter UI** interacts with the genetic algorithm and provides:

- **Input Field:** Allows the user to enter the board size (N).
- **Chessboard Visualization:** Displays solutions with queens as red circles.
- **Navigation Buttons:** Lets the user view the next solution or quit the program.

//for N=4



//for N=7



Advantages of Genetic Algorithm for N-Queens

1. Scalability:

- Efficiently handles larger N compared to exhaustive search methods like backtracking.

2. Randomized Approach:

- Does not rely on pre-defined patterns or rules, making it flexible.

3. Fast Approximation:

- Quickly finds valid solutions even for larger problem sizes.

Limitations

1. Non-Guaranteed Completeness:

- May not find all solutions due to randomness.

2. Performance:

- Requires fine-tuning of parameters like `population_size` and `mutation_rate`.

3. Computational Overhead:

- For very large N, the algorithm may take longer to converge.