# Exploration of Genetic Algorithms Through the Iterative Prisoner's Dilemma

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#### Abstract

Genetic algorithms are used for many optimization problems to find a near-optimal solution when finding the optimal solution would be too time-consuming. Although unable to tell when it has found the optimal solution, a genetic algorithm continues until the probability of having found the optimal solution is sufficiently high. There are many methods used to perform each step of a genetic algorithm, but it is not easy to identify which will work best for a specific problem. The goal of this project is to compare these different methods through the iterative prisoner's dilemma, and to hopefully find which methods work best generically.

#### 1 Introduction

The prisoner's dilemma is a problem involving two players. The players each must decide whether to cooperate with the other or to defect. These decisions must be made without knowledge of the other player's decision. Points are given to each player based on the moves they made. If both players cooperate, they are each given R points. If one cooperates and the other defects, the cooperating player receives S points and the defecting player receives T points. If they both defect, they are each given P points. In order for it to be a prisoner's dilemma, the values must follow the inequality T > R > P > S.

In the iterative prisoner's dilemma, the additional inequality 2R > T + S must be satisfied. In this scenario, the same thing happens, except that the players are against each other many times with memories of the past. In this scenario the best outcome is for both players to cooperate each time, because the total points given when both cooperate (2R) is greater than the number of points given if one defects (S + T) and greater than the number of points if both defect (2P). The problem associated with the iterative prisoner's dilemma is to find the rule that should be followed in order to maximize the number of points received when it participates in this scenario with a variety of other players.

This is a good problem on which to use a genetic algorithm because there is no algorithm faster than brute force that has been proven to find the optimal rule. In my genetic algorithm, I made each solution a collection of bits that represent whether the player should cooperate or defect given a past collection of turns. The fitness value for each possible solution is the number of points it accumulates after going through a set number of turns with each other possible solution in the population. The methods by which the each part of the genetic algorithm is done can be changed easily because each possible solution is a simple string of bits.

### 2 Background

The iterative prisoner's dilemma has been studied extensively in the past. Because the best rule is agreed upon, it is a good case with which to test genetic algorithms. It has been shown that the best rule is to cooperate on the first turn, and then do the same thing that the opposing player did on the previous turn for the rest of the turns. The only exception is if the opposing player defected the previous turn, there should be a 3% chance for the rule to tell the player to cooperate instead of following the opposing player. This rule was found by Robert Axelrod through a series of tournements in which he invited colleagues to devise rules for the iterative prisoner's dilemma, and then had them all play against each other. Many people have written a genetic algorithm that solves the iterative prisoner's dilemma, but I have not found a case in which this problem was used to study genetic algorithms.

#### **3** Development Sections

My program can run a genetic algorithm to find a solution to the iterative prisoner's dilemma using many different genetic algorithms methods. It can take user input to tell it which method to use for each part of the algorithm: the initial population creation, recombination, mutation, and natural selection. It can also set essential constants: the mutation rate, the number of generation, the size of the population, and the number of iterations of "memory" that the solutions have when running the prisoner's dilemma. As the program runs, it displays a graph showing the fitness of each of the current solutions. It also shows a graph that shows the average fitness of each previous generation. After the genetic algorithm completes its run, another method looks at the average fitness values and determines at which generation the fitness level stabilized. This represents how long it took the genetic algorithm to find the optimal solution. After many genetic algorithm runs with different methods and constant values it should become obvious which is most efficient.

Finally, it outputs a file with these average fitness values so that they can again be graphed after the program has completed the run. I used averages of the number of generations each method takes to get to the optimal solution in order to judge the usefulness of each method. The program automatically runs the genetic algorithm with each set of parameters and then determines when it found the correct answer (i.e. how many generations it took). The output of the program is made up of a file for each configuration of the genetic algorithm. Each file contains the results of 10 runs and an average of them. The analysis was done with a program that graphs based on attributes chosen by the user. This will hopefully aid in the deciding of which genetic algorithm methods to use by future programmers.

#### 4 Results

Based on my analysis, I was able to determine a number of things about the genetic algorithm that I was using.

First of all, I was unable to determine which mutation and initial population selection types were best. This is because the number of possible solutions was smaller than the number of solutions in the population. This means that it was very likely to have one of each possibility in the population with any of the initial population types. Additionally, mutation was unnecessary since all of the possibilities were present from the beginning.

For the recombination algorithms, I found that the double-point algorithm was slightly better than the single-point algorithm. Using the fitness-based algorithm for eliminating parts of the population was better than the static algorithm. Larger percentage of the best score required was better than smaller percentages, although I only tested up to 95%.

## 5 Bibliography

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