

A GENETIC ALGORITHM WITH A MULTI-LAYERED GENOTYPE-PHENOTYPE MAPPING

Seamus Hill and Colm O’Riordan

Computational Intelligence Research Group, National University of Ireland, Galway, Ireland

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Abstract: In this paper we investigate the introduction of a multiple-layer *genotype-phenotype* mapping to a Genetic Algorithm (GA) which attempts to mimic more closely, the effects of nature. The motivation for introducing multiple-layers into the *genotype-phenotype* mapping is to create a many-to-one *genotype-phenotype* mapping. The paper compares a traditional GA with a GA containing a multi-layered *genotype-phenotype* mapping using a number of well understood problems in an attempt to illustrate the potential benefits of including the multi-layered mapping. Initial findings suggest that the multi-layered mapping between the *genotype-phenotype* used in conjunction with a binary representation outperforms existing traditional GA approaches on well known problems, while still allowing the use well understood genetic operators.

1 INTRODUCTION

Genetic Algorithms (GAs) (Holland, 1975) are search algorithms based on the mechanics of natural selection and natural genetics and are often used to solve complex optimisation problems. However in nature there exists an intermediate layer between the *genotype* and the *phenotype*, which differs from the mapping found in a traditional GA. By introducing multiple-layers, which attempts to mimic more closely the effects of nature, into the *genotype-phenotype* mapping we aim to and to maintain diversity in the population and introduce a many-to-one *genotype-phenotype* mapping. The paper compares a GA with a multi-layered *genotype-phenotype* mapping to a traditional GA using a number of well understood problems using different levels of deception in an attempt to create a proof of concept. The paper is organised as follows: section 2, a review of a selection of previous work. Section 3 outlines the multiple mapping GA proposed by the authors. Section 4 describes the experiments conducted, while section 5 discusses the findings. Finally, section 6 concludes and outlines future work.

2 BACKGROUND

The motivation for attempting to create a GA which includes a number of biologically plausible concepts and yet provides a framework based on a binary

string, stems from the desire to allow a GA to provide a many-to-one representation and to test how traditional GA operators operate within this environment. Another motivation is that for a simple GA, solving deceptive problems can be difficult because of premature convergence. By using intermediate layers in the *genotype* space, it is hoped to overcome this failing while still using a binary representation.

3 MULTI-LAYERED MAPPING GENETIC ALGORITHM (MMGA)

The mmGA operates using a binary representation which allows the use of standard genetic operators such as crossover and mutation. The difference between the multi-layered mapping GA (mmGA) and the standard representation found in a GA lies in the fact that it contains multiple layers of mappings between the *genotype* to the *phenotype*. To discuss the mapping, consider the collections of genes 11100001, 11101100, 10110001 and 11000110. The mmGA randomly creates the genome string and the strings are converted to a *DNA template* using the following mapping 00 → A; 01 → C; 10 → G and 11 → T. This maps the strings 11100001 → TGAC; 11101100 → TGTA; 10110001 → GTAC and 11000110 → TACC. The next step in the mmGA mapping is a conversion

from the template to DNA coding using the mapping $G \rightarrow C$; $A \rightarrow T$; $T \rightarrow A$ and $C \rightarrow G$. The templates TGAC becomes ACTG, TGTA becomes ACAT, GTAC becomes CATG and TACG maps to ATGC.

The final stage of *transcription* are now carried out and the DNA coding strings are now converted into RNA where the only change maps $T \rightarrow U$. The final stage of *transcription* are now carried out and the DNA coding strings are now converted into RNA. The strings ACTG, ACAT, CATG and ATGC now become the *proteins* ACUG, ACAU, CAUG and AUGC respectively. These *proteins* are in turn then converted into *traits* i.e. 0 or 1 based upon the mapping generated by the mmGA, which are then combined to create the *phenotype*. The mappings created by the mmGA allow for a many-to-one relationship between the *genotype* and the *phenotype*. The mapping can be summarised as follows; 11100001 \rightarrow TGAC \rightarrow ACTG \rightarrow ACUG \rightarrow 0; 11101100 \rightarrow TGTA \rightarrow ACAT \rightarrow ACAU \rightarrow 1; 10110001 \rightarrow GTAC \rightarrow CATG \rightarrow CAUG \rightarrow 0 and 11000110 \rightarrow TACG \rightarrow ATGC \rightarrow AUGC \rightarrow 1. The pseudocode for the process is outlined in Algorithm 1.

Algorithm 1: Pseudocode - multi-layered mapping Genetic Algorithm.

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initialize mmGA;
r=1; (Number of runs);
for Number of runs do
  Initialise Individual Genomes P(g);
  Transcribe Genome to Amino Acids P(g);
  Translate Amino Acids to Phenotype P(g);
  Evaluate P(g); (Phenotype fitness);
  for Number of Generations do
    g=0; (generations);
    for All members of P do
      Select P(g) from P(g-1);
      Crossover P(g); genotype level;
      Mutation P(g); genotype level;
      Transcribe Genome to Amino Acids P(g);
      Translate Amino Acids to Phenotype P(g);
      Evaluate P(g); (phenotype fitness);
    end
    g+=1;
  end
  r+=1;
end
end mmGA;

```

4 EXPERIMENTS

In this paper the authors choose four types of problem to examine the effects of extending the representation; a two-bit minimal deceptive problem, a three-bit fully deceptive problem, a 100-bit One Max problem and a fully deceptive 30 bit problem.

4.1 Minimal Deceptive Problem (MDP)

A problem which causes a GA to diverge from the global optimum. can be viewed as a deceptive problem. By using short low-order building block to lead the search away from the global optimal to a sub-optimal point in the search space we are deceiving the GA. The MDF exhibits the characteristics of an epistatic problem and as it can be shown that one-bit problems cannot be deceptive the MDP is the smallest deceptive problem possible and by using a MDP one can carry out analysis into the workings of GAs.

4.2 Three Bit Fully Deceptive Problem

A fully deceptive problem of order-N can be viewed as being deceptive when all of the lower-order hyperplanes lead away from the global optimum and towards a deceptive attractor (Whitley, 1991). We use a fully deceptive order-3 problem as outlined by (Goldberg et al., 1990) in this paper.

4.3 One-max Problem

The One-Max problem (Ackley, 1987) can be described formally as having a string $\bar{x} = \{x_1, x_2, \dots, x_N\}$, with $x_i \in \{0, 1\}$, which attempts to maximise the following:

$$f(\bar{x}) = \sum_{i=1}^N x_i$$

In this paper the authors have defined $N = 100$.

4.4 Thirty-bit Fully Deceptive Problem

One failing of the 3-bit fully deceptive problem is that it is too small to really demonstrate a search strategy. The thirty-bit problem as outlined in (Goldberg et al., 1990) expands the three-bit problem into ten three-bit deceptive order-three subfunctions. To increase the level of difficulty we include a *loose ordering*, which makes the problem fully deceptive. This is achieved by increasing the defining length to twenty, where the defining length is the maximum distance between two defining symbols in a schema.

5 FINDINGS

We compare both the traditional GA and the mmGA over a suite of problems containing a two-bit minimal deceptive problem, a three-bit fully deceptive problem, a one hundred-bit One-Max problem and a fully deceptive thirty-bit problem. For the first three problems, each set of experiments has a population size of one hundred and is executed for two hundred generations. For the final problem, the thirty-bit fully deceptive the population size was increased to two hundred and the algorithm run for three thousand generations. Each experiment is executed over one hundred runs.

Figure 1 illustrates that the performance of the mmGA compares with that of a traditional GA., possibly because the MDP problem is easy for both forms of a GA to solve. Which may indicate that there is little to be gained by adopting the more complex multi-layered GA on easy problems. Figure 2 shows the performance of both GAs on a three-bit fully deceptive problem, which is considered *GA hard*. Although both GAs are successful in locating the *global optimum*, the GA using a traditional representation converges on the *global optimum*, while the mmGA converges for a number of generations but due to the multi-layered representation and many-to-one representation continues to explore the search space.

When comparing the results in Figure 2 to those of Figure 1, we see that although both forms of GA converge on the *global optimum*, it appears that the more difficult landscapes offer the GAs a more challenging task and that there may be an advantage in using a multi-layered mapping particularly on difficult problems. In Figure 3 we see the performance of both GAs on the One-Max problem. However, the One-Max problem can be viewed as a landscape that is relatively easy for a GA to search. The results indicate that both forms of a GA were successful at converging on the *global optimum*. The standard GA appears to have located the *global optimum* in a shorter time frame than the mmGA. The authors believe that the reason for this is due to the difference in the *genotype-phenotype* mapping and that the extra time take for the mmGA relates to the increased search space. This we believe should not be viewed negatively as it reduces the prospect of premature convergence for the mmGA.

The thirty-bit fully deceptive problem makes the problem far more difficult to solve particularly as the defining length has increased. Figure 4 show that the average fitness level per generation for the traditional GA is higher than that of the mmGA, as the traditional GA has converged prematurely on the deceptive attractor. However, figure 5 shows the average

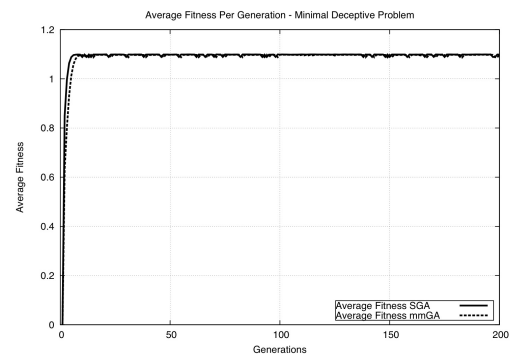


Figure 1: Two-bit minimal deceptive problem - Average fitness per generation.

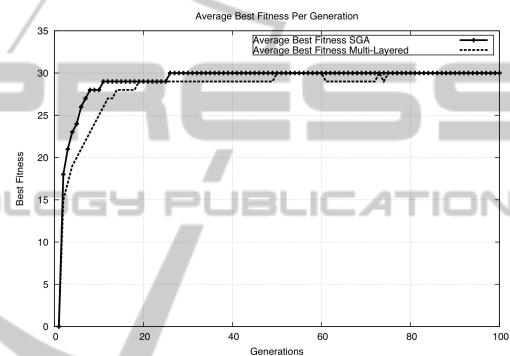


Figure 2: Fully deceptive problem - Average best fitness per generation.

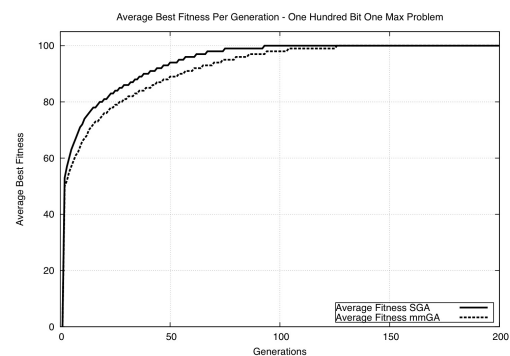


Figure 3: One hundred bit one max problem - Average best fitness per generation.

best fitness per generation and results show that the SGA failed to locate the global optimum as it converged on the deceptive attractor. The mmGA however, did manage to locate the optimum. This appears to indicate that the *genotype-phenotype* mapping contained in the mmGA exhibits the ability to avoid premature convergence and continue the search to locate the global optimum. Figure 6 allows us to see the number of subfunctions discovered per gen-

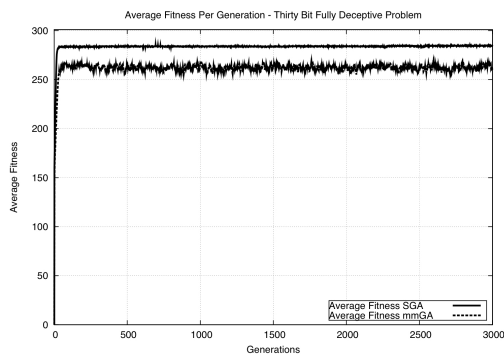


Figure 4: Thirty-bit fully deceptive problem - Average fitness per generation.

eration. We can see from the plot that mmGA outperforms the traditional GA in locating the ten fully deceptive subfunctions.

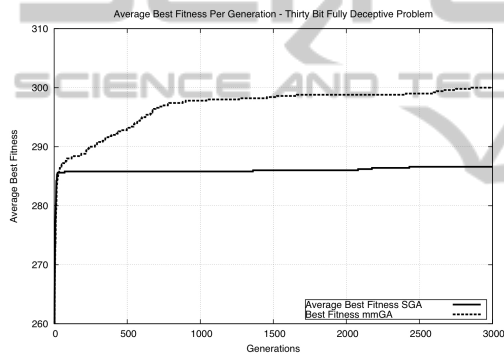


Figure 5: Thirty-bit fully deceptive problem - Average best fitness generation.

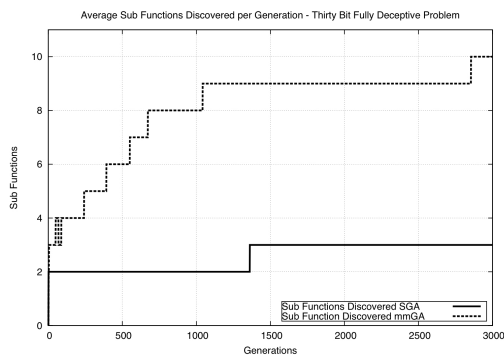


Figure 6: Thirty-bit fully deceptive problem - Sub functions discovered 3000 generations.

6 CONCLUSIONS

Initial results indicate that the mmGA has the ability to solve a number of problems over varying fitness landscapes. By introducing a multi-layered relationship between the *genotype* and the *phenotype*, the mmGA offers the ability to introduce a many-to-one *genotype-phenotype* mapping. This implies that identical *phenotypes* may be created from different *genomes*. From the results outlined above, it appears that the extended *genotype-phenotype* representation exhibits the ability to avoid premature convergence, particularly on more difficult problems.

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