

Altering the Granularity of Neutrality in a Multi-Layered Genetic Algorithm

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Abstract: By adopting a basic interpretation of the biological processes of *transcription* and *translation*, the multi-layered GA (MGA) introduces a genotype-phenotype mapping for a haploid genotype, which allows the granularity of the representation to be tuned. The paper examines the impact of altering the level of neutrality through changes in the granularity of the representation and compares the performance of a standard GA (SGA) to that of a number of multi-layered GAs, each with a different level of neutrality, over both static and changing environments. Initial results indicate that it appears advantageous to include a multi-layered, biologically motivated genotype-phenotype encoding over more difficult landscapes. The paper also introduces an interpretation of *missense* mutation, which operates within the genotype-phenotype map (GP-map). Results also suggest that this mutation strategy can assist in tracking the optimum over various landscapes

1 INTRODUCTION

In standard GAs (SGA) variation is applied solely to genetic structures and not to phenotypic structures and each phenotype is represented by a distinct genotype. However, in nature we find a layered mapping between the genotype and the phenotype and that one phenotype can be represented by a number of different genotypes. Also, in nature, although most variation happens at the DNA level, mutation can also occur at the RNA level. By introducing a many-to-one GP-map into a GA you can represent these phenomena. Furthermore, this map introduces a level of neutrality into the representation.

The paper examines the impact of varying the level of neutrality by altering the granularity of the representation and examining to see whether or not, there are benefits associated with the inclusion of variation within the layers of the GP-map. Variation within the layers of the GP-map is achieved through an implementation of a form of variation operator found in biology, known as *missense* mutation. In biology, a point mutation that changes a codon that normally specifies a particular amino acid into one that codes for a different amino acid is known as a *missense mutation*. In a simplified summary of the biological process the pathway from DNA to a protein includes a transcription stage, which maps DNA to RNA and a translation stage which maps RNA to pro-

teins. Our intention is to include traditional variation operators (crossover/mutation) at the DNA level and to incorporate an interpretation of missense mutation at the RNA level. The motivation is to develop a robust GA which includes a tunable GP-map and resists premature convergence by incorporating redundancy through the GP-map, while allowing the use of a haploid binary representation. The contribution comes firstly, from examining the impact of varying the level of neutrality in the representation and secondly, from the analysis of a mutation strategy which operates within the GP-map.

The paper is organised as follows: Section 2 briefly examines previous research, Section 3 outlines the workings of the multi-layered GA (MGA). Section 4 gives an overview of the experiments, while Section 5 describes the results and finally, Section 6, discusses the conclusions.

2 Background

A population’s ability to survive in various environmental conditions often requires a level of diversity to be maintained within the population. As a GA’s search involves a mapping between the genotype and the phenotype, with a SGA, diversity is quickly eliminated from the population through its selection policy and low mutation rates. Generally, two strate-

gies are used to modify a SGA in order to improve its adaptiveness in a changing landscape (Grefenstette and Cobb, 1993): introduce increased memory to store responses for a environmental changes or use a method to increase diversity within the population i.e. increase mutation (Grefenstette and Cobb, 1993). However, by implicitly introducing neutrality into the GP-map, the level of diversity within the population can be maintained and can assist in tracking the optimum over a number of different landscapes (Hill and O’Riordan, 2011).

Deception is often used in testing GAs and implies that the search strategy can be misled (Whitley, 1991). As noted in (Morrison and DeJong, 2002), diversity is critical for GAs, particularly when the landscape is evolving as recombining a homogeneous population will not enable the GA to locate the new optimum. Hamming difference is used as a measure of diversity both for the genotypic and phenotypic diversity. In this paper we use a 30-bit and a 90-bit One Max max problem to examine the performance of the GAs over static landscapes and to examine how the GAs perform over a changing landscape we use a three-bit problem, introduced by (Whitley, 1991), which we expand into a thirty-bit problem, as outlined in (Goldberg et al., 1990).

The MGA introduces a tunable multi-layered GP-map, which allows a haploid GA to exhibit, some of the characteristics normally associated with a diploid i.e. a mechanism for allowing alleles or combinations of alleles which proved useful in previous generations (Goldberg and Smith, 1987) and thus maintaining a form of long term memory, without the need to develop a dominance scheme. The MGA population consists of a population of haploid individuals, which allows for the use of traditional crossover and mutation variation operators on the genotype. This differs from the approach used by diploid GAs (DGAs) i.e. (Yang, 2006), where each individual has two chromosomes and crossover is divided into two steps and mutation is viewed as being neutral. Another difference between the MGA mapping and that of a DGA, is that in the DGA, a phenotype allele is made up from a single genotype allele which is expressed. In the MGA a single phenotype allele is made from the cardinality incorporated in the genotype i.e. in this paper 4-bits, 6-bits or 8-bits. Although the MGA’s GP-map is non-deterministic, the approach differs from that of real-coded binary representation, which incorporate a gene-strength adjustment mechanism (Kubalik, 2005). Real-coded binary representations can use standard crossover operators, but mutation is implicit due to the gene-strength adjustment mechanism (Kubalik, 2005).

MGA 4-2-1 Mapping

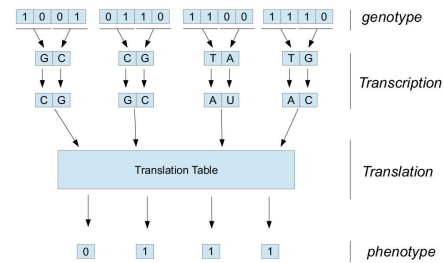


Figure 1: 4-2-1 MGA Representation Mapping

MGA 6-3-1 Mapping

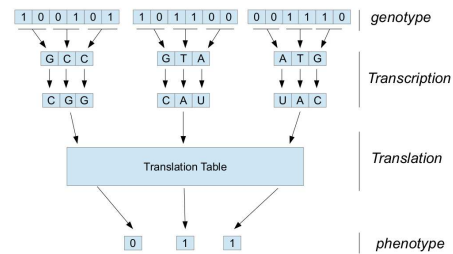


Figure 2: 6-3-1 MGA Representation Mapping

MGA 8-4-1 Mapping

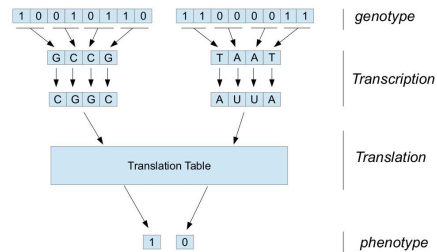


Figure 3: 8-4-1 MGA Representation Mapping

3 Multi-layered GA (MGA)

The transcription phase of the MGA maps, in two steps, the binary genotype to a string of characters taken from a four letter alphabet A, C, G and T . The mapping is carried out as follows $00 \rightarrow A$; $01 \rightarrow C$; $10 \rightarrow G$ and $11 \rightarrow T$. Once the initial mapping is complete, variation takes place amongst the characters, which may be viewed as a unique form of inversion. The mappings for this stage are $A \rightarrow U$; $C \rightarrow G$; $G \rightarrow C$ and $T \rightarrow A$. Following transcription, the translation phase takes place. Upon initialisation, the MGA creates a translation table, using characters taken from a four letter alphabet A, C, G and U , based on the granularity of the neutrality selected by the

user. T and U are used for biological plausibility and are interchangeable. In this paper we tune the MGA to use three different representations and examine the impact of altering the granularity: a 4-bit MGA representation; a 6-bit MGA representation and an 8-bit MGA representation. The size of the translation table is determined by the representation chosen. For a 4-bit MGA representation, a translation table of 2^4 is created; with a 6-bit MGA representation a 2^6 translation table is needed and with an 8-bit MGA representation a 2^8 translation table is required. The size of the translation table represents the granularity of neutrality which exists within the representation. A 4-bit MGA representation requires 4 bits for each individual element of the phenotype, which we refer to as a *phene*; a 6-bit MGA representation requires 6 bits for each phene and an 8-bit representation requires 8 bits for each phene. A form of neighbourhood equivalence is used in the translation phase, to map a predefined number of characters to a given phene, in this paper either a 0 or a 1. Figures 1, 2 and 3 represent a 4-bit MGA representation, a 6-bit MGA representation and an 8-bit MGA representation respectively. For a more detailed explanation of the MGA see (Hill and O’Riordan, 2011). Missense mutation takes place between the transcription and translation phases. This form of mutation flips a character from the four letter alphabet A, C, G and U , i.e. A can be flipped to either C, G or U etc.

4 Experiments

Experiments were carried out over both a One Max problem static landscape and a deceptive changing landscape. As we intend to examine the relationship between altering the granularity of the MGA representation (which determines the level of neutrality) and problem difficulty, experiments are carried out over increasingly difficult landscapes. The One Max problem experiments include 30-bit and 90-bit problems, while the deceptive changing landscape experiments use a ten 3-bit fully deceptive changing landscape and a more difficult thirty 3-bit fully deceptive changing landscape. The parameters for the experiments are as follows: one-point crossover is used at a rate of 0.7, single-point mutation is used at a rate of $1/l$, where l is the length of the chromosome and missense mutation is at a rate of $5/r$ where r is the length of the RNA string. The population consisted of 200 individuals, with the experiment results being averaged over 10 runs.

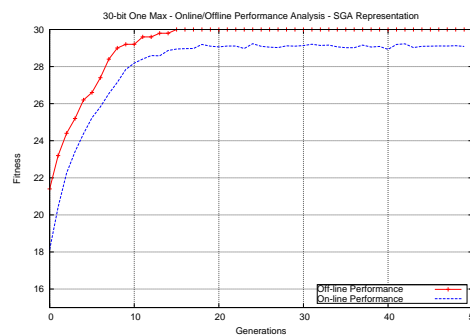


Figure 4: SGA - 30-Bit One Max Static Landscape

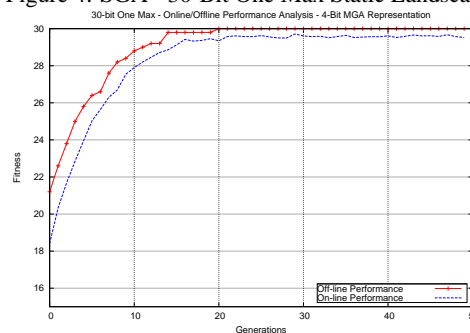


Figure 5: MGA 4-2-1 Representation 30-Bit One Max Static Landscape

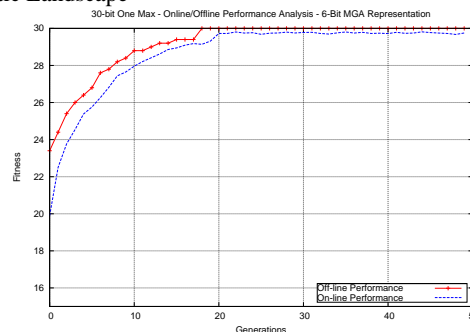


Figure 6: MGA 6-3-1 Representation 30-Bit One Max Static Landscape

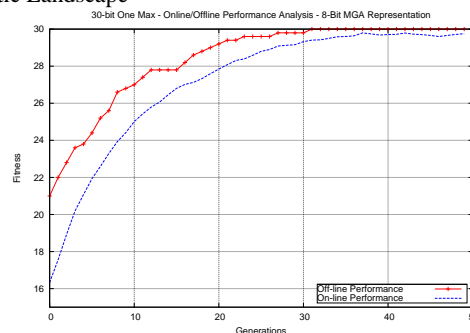


Figure 7: MGA 8-4-1 Representation 30-Bit One Max Static Landscape

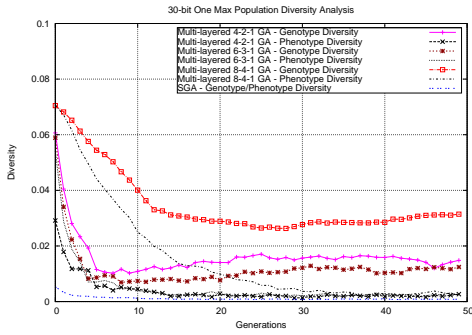


Figure 8: Diversity 30-Bit One Max SGA & MGA

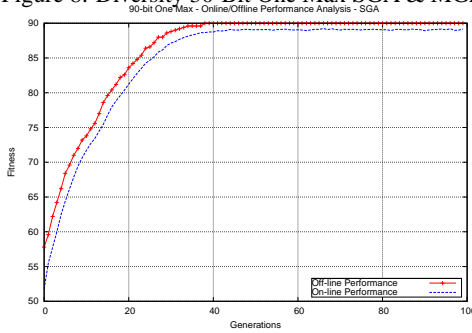


Figure 9: SGA 90-Bit One Max Static Landscape

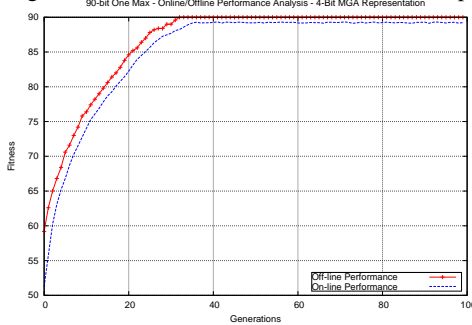


Figure 10: MGA 4-2-1 Representation 90-Bit One Max Static Landscape

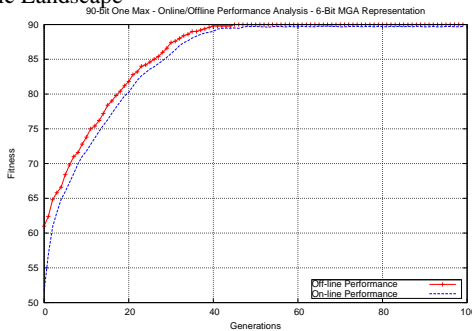


Figure 11: MGA 6-3-1 Representation 90-Bit One Max Static Landscape

5 Results

5.1 30-bit One Max Problem

Figures 4, 5, 6 and 7 compare the average-best (off-line) and the average (on-line) performance of an

SGA against a number of different MGA representations over a 30-bit One Max problem. Overall, although there is little difference in the performance of the SGA against that of the variously tuned MGAs, with the SGA performing marginally better in the off-line results and marginally worse in the on-line performance. Of the three MGA representations, the 6 – 3 – 1 representation produces a relatively similar off-line performance to the SGA, but returns a marginally better on-line performance. Figure 8 compares the level of diversity in the population between the SGA and the MGA representations. The SGA phenotypic diversity is similar to that of the MGA, however there is a significant difference between the genotypic diversity of the MGA representations with that of the SGA. The above results indicate that although there is little difference in the off-line results of the various GA's, the MGA's on-line performances are marginally better than those of the SGA, this may be a result of a greater level genotypic diversity associated with the MGA.

5.2 90-bit One Max Problem

As we increased the level of difficulty to a 90-bit One Max problem (illustrated in Figures 9, 10, 11 and 12), the performance of both the SGA and the various MGA's are again quite similar in terms of off-line performance. Although there may be a very marginal improvement in the on-line performance of the MGA over the SGA. Population diversity for the SGA and MGA representations is shown in Figure 13. As with diversity in the 30-bit One Max experiments, there is little difference at the phenotypic level, which is what we would expect, however there is a difference at the genotypic level, with the MGA variations maintaining a far higher level of diversity than that of the SGA, which has converged.

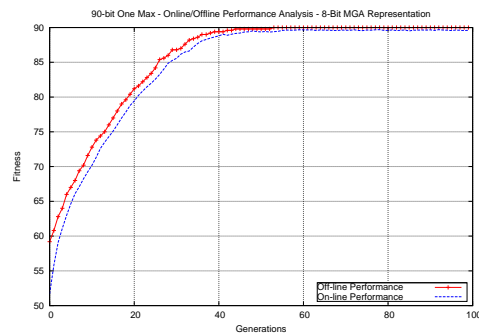


Figure 12: MGA 8-4-1 Representation 90-Bit One Max Static Landscape

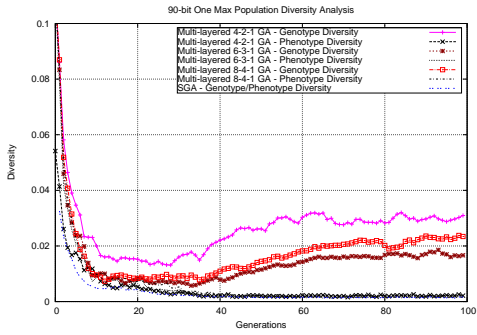


Figure 13: Diversity 90-Bit One Max SGA & MGA

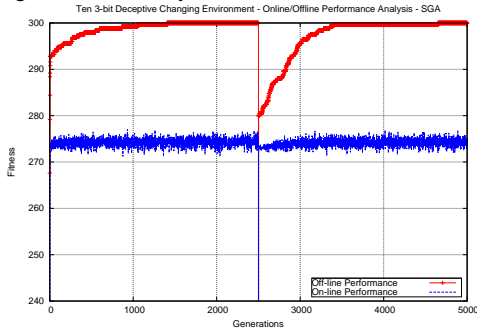


Figure 14: SGA - Ten 3-Bit Fully Deceptive Changing Landscape

5.3 Ten 3-bit Fully Deceptive Changing Landscape Problem

Figures 14, 15, 16 and 17 show the results of the SGA and the various MGA representations. As the problem landscape is comparatively easy, both the SGA and the various MGA representations, found the global optimum before and after the landscape changed, illustrating that they both managed to adapt. However, the results indicate that all of the MGA representations out performed the SGA in terms of locating the global optimum earlier in the search. With the performance improving as the level of granularity decreased. Examining the population diversity, we can see in Figure 18 that there is a significant difference between the SGA and MGA in relation to the maintenance of diversity within the population. Although the phenotypic diversities are relatively similar, there is a greater level of genotypic diversity maintained in the MGA representations. What's interesting here is that the level of genotypic diversity maintained by the 8-4-1 MGA representation was less than the MGA variations, this may account for its relatively poorer performance.

To develop an understanding of the impact of the transcription phase on the MGA, we ran an experiment using a 6-bit MGA representation without transcription. The results in Figure 19 show that although the search locates the global optimum, the performance

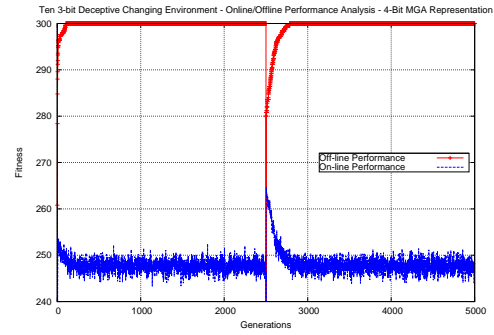


Figure 15: 4-2-1 MGA Representation - Ten 3-Bit Fully Deceptive Changing Landscape

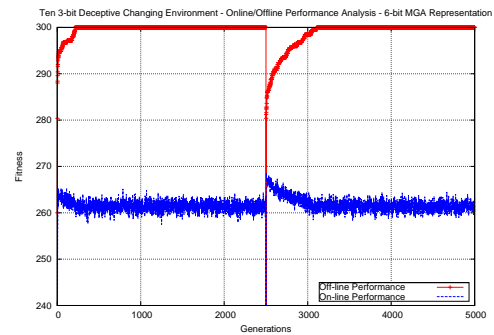


Figure 16: 6-3-1 MGA Representation - Ten 3-Bit Fully Deceptive Changing Landscape

was similar to the results achieved using transcription. However, this may indicate that the transcription phase has a minor impact prior to the translation phase, but further experiments over more difficult landscapes may assist in understanding the impact of transcription.

Figure 20 shows the results of not including the missense mutation operator. Even on the relatively easy ten three-bit landscape, the 6-bit MGA representation fails in its search for the global optimum, indicating that the missense mutation strategy assists in tracking the optimum over a changing landscape and without it the search struggles to escape from a deceptive attractor. To examine whether the results found by using the multi-layered MGA can be replicated by increasing the level of mutation in the SGA, we ran another set of experiments where we increased the level of mutation, as a method to increasing diversity (Grefenstette and Cobb, 1993), in the SGA to $2/l$. The results shown in Figure 21 illustrate that in comparison to Figure 14, the extra mutation has improved the performance of the SGA. However, it is worth noting that the off-line performance is lower, which is a result of the increased level of mutation and may prove problematic in a more challenging environment.

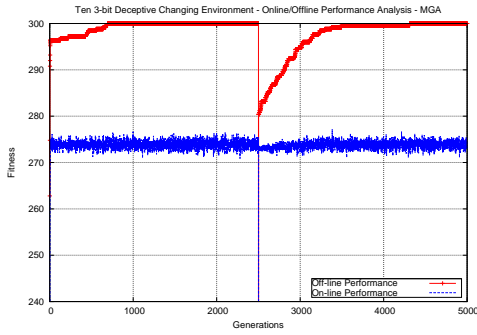


Figure 17: 8-4-1 MGA Representation - Ten 3-Bit Fully Deceptive Changing Landscape

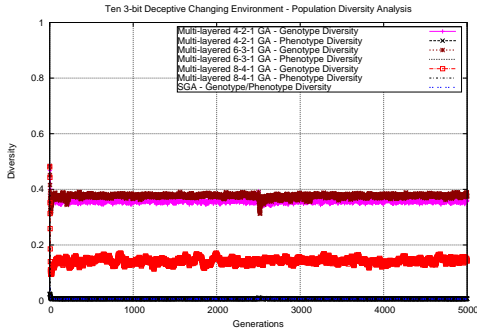


Figure 18: Diversity Ten 3-bit Fully Deceptive Changing Landscape Problem SGA & MGA

5.4 Thirty 3-bit Fully Deceptive Changing Landscape Experiments

Over the more difficult fully deceptive landscape the SGA attempts to locate the global optimum of 900, but fails, both before and after the landscape changes, see Figure 22. In contrast to this Figures 23, 24 and 25 show an improved performance from the MGA representations and although the 6-3-1 and 8-4-1 representations were successful, both before and after the landscape changed, the 4-2-1 MGA representation, however, failed to recover sufficiently after the landscape changed, indicating that the adaptive quality of the MGA may lessen as the level of granularity in the representation decreases over more difficult changing landscapes.

As with previous results the level of phenotypic diversity maintained by the various GAs remains quite similar (see Figure 26). However, the level of genotypic diversity maintained by the MGA representations is again greater than that of the SGA. Over this deceptive changing landscape, the levels of genotypic diversity maintained by the MGA representations are quite similar to one another. This may explain their relatively similar performances over the landscapes.

To understand the impact of transcription over a more difficult landscape, we ran an experiment using a 6-bit MGA representation, without the transcription phase,

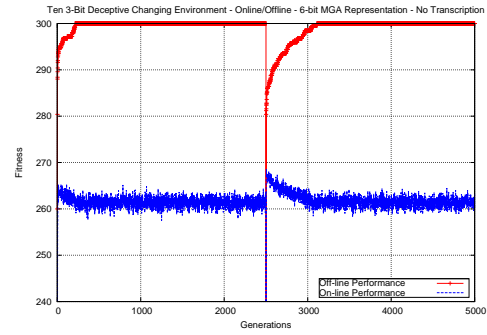


Figure 19: 6-3-1 MGA Representation - Ten 3-Bit Deceptive Problem - No Transcription

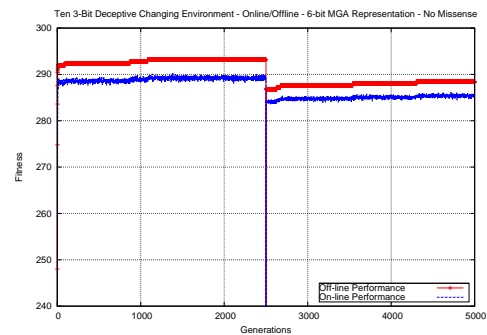


Figure 20: 6-3-1 MGA Representation - Ten 3-Bit Deceptive Problem - No Missense

illustrated in Figure 27. The results indicate that the 6-3-1 MGA representation fails to locate the global optimum. This result differs from our previous experiment, shown in Figure 19, suggesting over the more difficult landscape the transcription phase assists in the search strategy. One reason for this maybe that the transcription phase provides a form of inversion, which assists the search strategy over more difficult changing landscape.

By removing missense mutation, we hope to develop a better understanding of its impact over a more difficult environment. Figure 28 shows that the 6-3-1 MGA representation fails to escape the deceptive attractor throughout the search. In a similar way to the results shown in Figure 20, there appears to be too little variation, probably due to quite low mutation rates, in the population.

In order to establish whether the SGA may improve its performance with a higher level of diversity we increased the level of mutation. However, over the more difficult landscape, the SGA fails to locate the global optimum, shown in Figure 29. This appears to indicate that even with a higher level of diversity associated with higher mutation, the search strategy of the SGA fails over the given landscape.

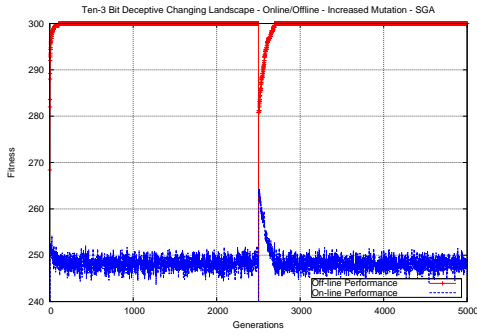


Figure 21: Ten 3-Bit Deceptive Problem - SGA - Increased Mutation

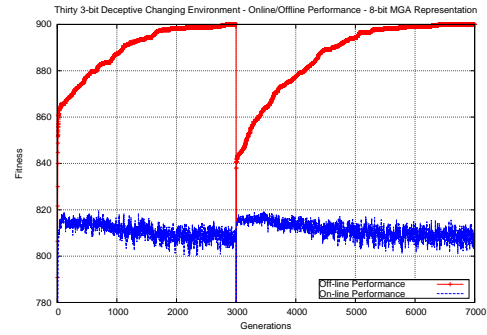


Figure 25: Thirty 3-Bit Fully Deceptive Changing Landscape 8-4-1 MGA Representation



Figure 22: Thirty 3-Bit Fully Deceptive Changing Landscape SGA

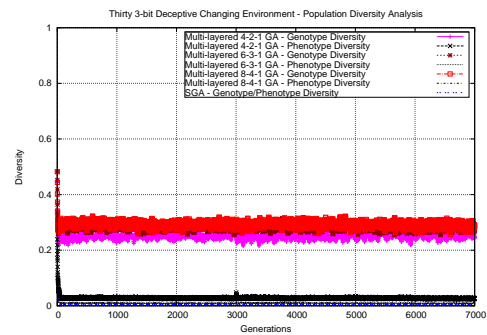


Figure 26: Thirty 3-bit Fully Deceptive Changing Landscape Problem SGA & MGA Representation Population Diversity

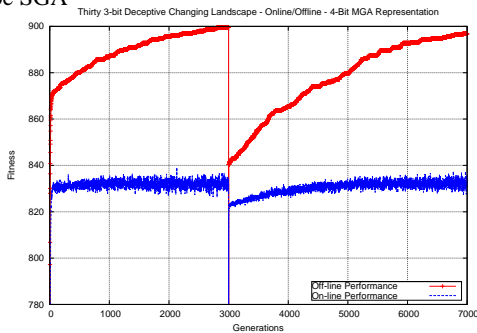


Figure 23: Thirty 3-Bit Fully Deceptive Changing Landscape 4-2-1 MGA Representation -

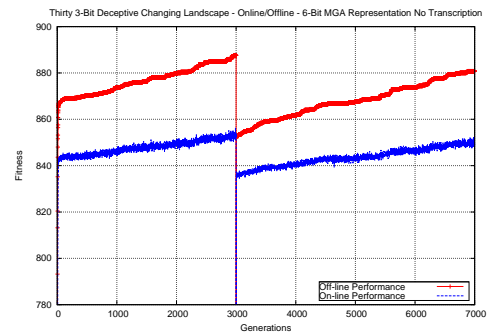


Figure 27: 6-3-1 MGA Representation - Thirty 3-Bit Deceptive Problem - No Transcription

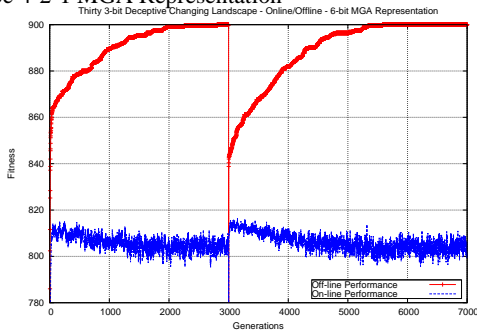


Figure 24: Thirty 3-Bit Fully Deceptive Changing Landscape 6-3-1 MGA Representation -

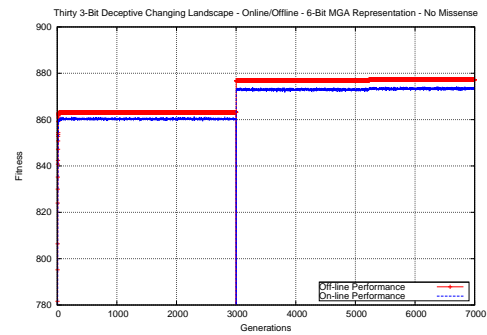


Figure 28: 6-3-1 MGA Representation - Thirty 3-Bit Deceptive Problem - No Missense

5.5 Statistical Significance

A two-sided paired Wilcoxon signed rank test was carried out on the results of each experiment and were

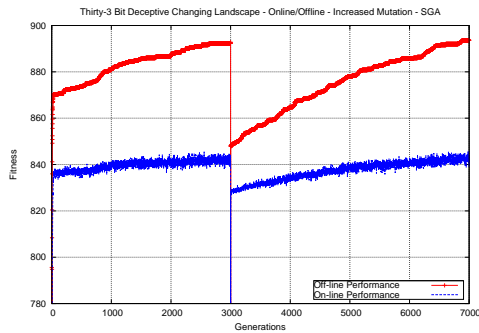


Figure 29: Thirty 3-Bit Deceptive Problem - SGA - Increased Mutation

shown to be highly significant with a P - value of $p < 2.2e - 16$ recorded for each of the experiments illustrated in the paper.

6 Conclusion

From the above experiments the results indicate that overall the MGA appears robust over both static and changing landscapes. By incorporating a tunable GP-map, the MGA offers the ability to vary the granularity of the representation, which appears beneficial over various landscapes and the performance of the MGA varies slightly depending on the level of granularity in the representation. Over the static One Max landscapes, the MGAs off-line and on-line performances were quite similar to those of the SGA over both the easier 30-bit problem and the more difficult 90-bit problem. On reason for this is that the problem, by its nature, is relatively easy for a GA to solve. What is of interest is that the MGA, in terms of robustness, performed as well as the SGA. However, over the deceptive changing landscape experiments the MGA outperformed the SGA, both on-line and off-line, particularly over the more difficult ten 3-bit deceptive problem. In relation to the use of Transcription, it appear from the results that the advantage of this phase is not apparent over less difficult landscapes, but once the level of difficulty increases, as with the more difficult deceptive problem, Transcription proved useful. one possible reason for this is that the Transcription phase performs a form variation, quite similar to inversion.

The inclusion of a missense mutation operator, which operated at the RNA level, results indicated that it assisted in the search strategy, over both difficult and less difficult landscapes. To examine whether this was a result of additional mutation, we carried out experiments which increased the mutation rate for the SGA and results indicated that over the more challenging deceptive landscape additional mutations failed to as-

sist the search strategy. This we feel, indicates that the use of missense mutation, which operates at levels within the GP-map may be beneficial.

To conclude, the inclusion of a biologically inspired tunable GP-map which allows the granularity of the GP-map to be altered appears to assist in searching various landscapes and further research into the levels of diversity maintained and population dynamics by the various MGA representations is required to gain a greater understanding.

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