# Multi-level selection genetic algorithm applied to CEC '09 test instances

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Abstract— Genetic algorithms (GAs) are population-based optimisation tools inspired by evolution and natural selection. They are applied in many areas of engineering and industry, on increasingly complex problems. To improve the performance, the new algorithms have a tendency to be derived from sophisticated mathematical and computational mechanisms, where many biological and evolutionary advances have been neglected. One such mechanism is multi-level selection theory which has been proposed as being necessary for evolution. Previously, an algorithm developed using this theory as its inspiration has shown promising performance on simple test problems. It proposes the addition of a collective reproduction mechanism alongside the standard individual one. Here the algorithm, Multi-Level Selection Genetic Algorithm (MLSGA), is benchmarked on more sophisticated test instances from CEC '09 and compared to the final rankings. In this instance a simple genetic algorithm is used at the individual level. The developed algorithm cannot compete with top algorithms on complex unconstrained problems, however it shows interesting results and behaviour, and better performance on constrained test functions. The approach provides promise for further

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investigation, especially in integrating state-of-the-art

individual reproduction methods to improve the performance

and improving the novel collective mechanism.

## I. INTRODUCTION

A number of evolutionary algorithms have been developed since the original work of Holland in the 1960's [1]-[19]. Each of them introduced novel mechanisms and improvements, with many of the leading algorithms benchmarked on complex multi-objective optimisation problems as part of CEC'09 [20]. Current trends are towards more complex algorithms based on mathematical and computational concepts, and advanced evolutionary-based concepts are often pushed aside. Evolutionary theory has made much progress since the original GA was developed and the authors feel that exploration of these modern evolutionary concepts can lead to an increase in performance of the modern genetic algorithms. One of them is multi-level selection originally proposed by Sober and Wilson [21].

In nature individuals often group together, into packs or tribes, in order to cooperate and increase their chances of Adam J. Sobey
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survival. During the life cycle, collectives are competing with each other, and similar to individuals, fitter groups survive and create offspring, while those with a lower fitness produce fewer or no offspring. This process can be compared to insect colonies, or human tribes, competing for food sources, where fitter colonies are able to collect enough to sustain their population and the less fit die from starvation [22]. According to Sober and Wilson [21] and Okasha [22] development of many skills of organisms such as, social skills and altruism are difficult to explain if selection only occurs at one level, as these traits are usually disadvantageous to the individual. A good example is self-sacrifice of the individual for the 'greater good" of the group, reducing the chances of the individual passing on its genetic material. Multi-level selection (MLS) theory states that fitness can be defined, and thus selection and competition can occur, separately on more than one level. In multi-level selection theory two main variants are proposed MLS1 and MLS2. In multilevel selection 1, both collective and individual levels of selections depend on a single evolutionary function and thus the fitness of the collective is simply the aggregate of the fitness of individuals inside of it. This means that both collective and each individual contribute to the improvement of this single function. In MLS2 the fitness function differs on each level and therefore the selection is independent; which generates an emergent selection property. Thus, it is possible that the evolution of collectives will be dependent on different key aspects than the evolution of individuals, leading to competition between them [22].

There have already been attempts to incorporate the levels of selection theory into evolutionary computation, however in these attempts key aspects have been ignored. Lenaert et al. [23] presented an algorithm with intradermic group selection. However, no direct selection method, with separate fitness, can be found at the higher level, which is required for a true multi-level selection approach. Akbari et al. [24][25] proposed an algorithm with three separate between-group dynamics: colonization, regrouping and migration. The colonization module partially reproduces MLS1, and after each generation the worst collective is eliminated with one of the remaining collectives reproducing and creating offspring. However, in this algorithm, only the best individual in the collective is considered to represent the fitness for the whole

collective, and the collective with one strong individual can survive, when other stronger in average, but with weaker the best solution will extinct, which results in the loss of many good solutions. In addition, only one collective takes part in the reproduction step, which decreases the possible gene pool. Wu and Banzhaf [26] introduced a genetic algorithm using a many level hierarchical model. In this algorithm reproduction occurs separately on group level and on the individual level. However, in this approach, groups from different levels can interact with each other during the reproduction step and basic concepts of MLS, such as separate selection mechanisms on each level, are not obviously introduced.

In this paper both MLS1 and MLS2 variants of MLSGA are benchmarked on a number of complex two-objectives constrained and unconstrained test instances, taken from CEC'09, in order to evaluate their performance on more demanding functions.

## II. METHODOLOGY

MLSGA splits the whole population into subgroups, collectives, which operate on the same search space, and introduces the additional level of selection. After each generation, the worst collectives, defined by low fitness, are eliminated and replaced by an equal number of new groups. generated from the remaining population. The size of the new collectives is equal to the size of the old ones in order to maintain a constant size for the whole population. Two variants of multi-level selection (MLS) are defined in MLSGA. In MLS1 the fitness function is defined as the same at both levels, collective and individual, and is the average of the f1 and f2 objective functions (1). In MLS2 both levels have separate fitness definitions; the f1 objective function is assigned to individual level (2) and f2 is assigned to collective level (3). The assignment of the fitness function at each level, individual or collective, has a large impact on the results as the both selection levels have a different selection pressure; where previous research implies that the individual level is stronger. In order to continue to investigate the sensitivity of the results to the way in which the objective is defined, MLS2R is defined here as the opposite of MLS2. Table I demonstrates the specific mechanisms used in this paper.

Fitness = 
$$\frac{f_1(x)+f_2(x)}{2}$$
 (1)  
Individual Fitness =  $f_1(x)$  (2)

Individual Fitness = 
$$f_1(x)$$
 (2)

Collective Fitness = 
$$f_2(x)$$
 (3)

TABLE I. MLSGA PARAMETERS

Step	Parameter	Value	
	Type	Random	
1. Initialisation	Engoding	Real values	
1. IIIItiansation	Encoding	16 decimals	
	Pop. Size	600	
2.	Method	SVM	
Classification	N. Collectives	6	
3. Individual Reproduction			

Step	Step Parameter	
3.1 Fitness Evaluation	Туре	MLS1 or MLS2
3.2 Selection	Туре	Roulette wheel
	Crossover type Crossover rate	Real variable SBX 0.7
3.3 Mating	Mutation type  Mutation rate	Polynomial 0.08
3.4 Elitism	Rate	0.1
4. Collective Repr	roduction	1
4.1 Fitness Evaluation	Туре	MLS1 or MLS2 or MLS2R
4.2 Elimination	Number of eliminated collectives	2
4.3 Replacement	Number of new collectives	2
5.Termination	Criterion	Reaching 500 generation

TABLE II. UNCONSTRAINED TEST CASES

Test	Objective functions
case	Objective functions
UF1 [28]	$f_{1} = x_{1} + \frac{2}{ J_{1} } \sum_{j \in J_{1}} y_{j}^{2}$ $f_{2} = 1 - \sqrt{x_{1}} + \frac{2}{ J_{2} } \sum_{j \in J_{2}} y_{j}^{2}$ $y_{j} = x_{j} - sin\left(6\pi x_{1} + \frac{j\pi}{n}\right), j = 2,, n$ For all UF functions: $J_{1} = \{j   j \text{ is odd and } 2 \leq j \leq n\}$ $J_{2} = \{j   j \text{ is even and } 2 \leq j \leq n\}$ $Search space is [0, 1] \times [-1, 1]^{n-1}$
UF2 [28]	$f_{1} = x_{1} + \frac{2}{ J_{1} } \sum_{j \in J_{1}} y_{j}^{2}$ $f_{2} = 1 - \sqrt{x_{1}} + \frac{2}{ J_{2} } \sum_{j \in J_{2}} y_{j}^{2}$ $y_{j} = \begin{cases} x_{j} - h(t)cos\left(6\pi x_{1} + \frac{j\pi}{n}\right)j \in J_{1} \\ x_{j} - h(t)sin\left(6\pi x_{1} + \frac{j\pi}{n}\right)j \in J_{1} \end{cases}$ $h(t) = 0.3x_{1}^{2}cos\left(24\pi x_{1} + \frac{4t\pi}{n}\right) + 0.6x_{1}$ Search space is $[0, 1] \times [-1, 1]^{n-1}$
UF3 [28]	$f_{1} = x_{1} + \frac{2}{ J_{1} } \left( 4 \sum_{j \in J_{1}} y_{j}^{2} - 2 \prod_{j \in J_{1}} \cos\left(\frac{20y_{j}\pi}{\sqrt{j}}\right) + 2 \right)$ $f_{2} = 1 - \sqrt{x_{1}} + \frac{2}{ J_{2} } \left( 4 \sum_{j \in J_{2}} y_{j}^{2} - 2 \prod_{j \in J_{2}} \cos\left(\frac{20y_{j}\pi}{\sqrt{j}}\right) + 2 \right)$ $y_{j} = x_{j} - x_{1}^{0.5 \left(1 + \frac{3(j-2)}{n-2}\right)}, j = 2,, n$ $Search space is [0, 1]^{n}$
UF4	$f_1 = x_1 + \frac{2}{ J_1 } \sum_{i \in J} h(y_i)$
[28]	$J_1 \mid J_1 \mid \sum_{j \in J_1} J_j \mid J_j $

Test case	Objective functions		
	$f_2 = 1 - x_1^2 + \frac{2}{ J_2 } \sum_{j \in J_2} h(y_j)$		
	$y_j = x_j - \sin\left(6\pi x_1 + \frac{j\pi}{n}\right), j = 2, \dots, n$		
	$h(t) = \frac{ t }{1 + e^{2 t }}$		
	Search space is $[0,1] \times [-2,2]^{n-1}$		
	$f_1 = x_1 + \left(\frac{1}{2N} + \varepsilon\right)  sin(2N\pi x_1)  + \frac{2}{ J_1 } \sum_{j \in J_1} h(y_j)$		
UF5	$f_2 = 1 - x_1 + \left(\frac{1}{2N} + \varepsilon\right)  sin(2N\pi x_1)  + \frac{2}{ J_2 } \sum_{j \in J_2} h(y_j)$		
[20]	$y_j = x_j - \sin\left(6\pi x_1 + \frac{j\pi}{n}\right), j = 2, \dots, n$		
	$h(t) = 2t^2 - \cos(4\pi t) + 1$		
	Search space is $[0,1] \times [-1,1]^{n-1}$		
	$f_1 = \sqrt[5]{x_1} + \frac{2}{ J_1 } \sum_{j \in J_1} y_j^2$		
UF7 [28]	$f_2 = 1 - \sqrt[5]{x_1} + \frac{2}{ J_2 } \sum_{j \in J_2} y_j^2$		
	$y_j = x_j - \sin\left(6\pi x_1 + \frac{j\pi}{n}\right), j = 2, \dots, n$		
	Search space is $[0,1] \times [-1,1]^{n-1}$		

In MLSGA the Pareto front is stored externally, and is used for illustrative purposes and to calculate the Inverted Generational Distance (IGD), but is not integral to the reproduction mechanisms. The biggest advantage of the approach is the fact that the mechanism allows for the use of any standard genetic algorithm mechanism at the individual level, though a simple mechanism is selected for early benchmarking. IGD is the performance measure function of Pareto front, which shows the average distance between all points in the true Pareto front and the closest solution from the achieved set and is calculated by (4);

$$IGD(A, P^*) = \frac{\sum_{v \in P^*} d(v, A)}{|P^*|};$$
 (4)

where P\* is a set of uniformly distributed points along the Pareto front, in the objective space, A is an approximate set to the Pareto front and d(v, A) is the minimum Euclidean distance between v and the points in A.

The test cases used are the unconstrained UF and constrained CF functions taken from the CEC '09 competition [28] and are shown in the Table II and Table III. The results are compared with other algorithms following the CEC '09 competition rules [20], based on the calculated IGD values.

TABLE III. CONSTRAINED TEST CASES

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Test case	Objective functions
CF1 [28]	$f_{1} = x_{1} + \frac{2}{ J_{1} } \sum_{j \in J_{1}} \left( x_{j} - x_{1}^{0.5 \left( 1 + \frac{3(j-2)}{n-2} \right)} \right)^{2}$ $f_{2} = 1 - x_{1} + \frac{2}{ J_{2} } \sum_{j \in J_{2}} \left( x_{j} - x_{1}^{0.5 \left( 1 + \frac{3(j-2)}{n-2} \right)} \right)^{2}$ For all CF functions: $J_{1} = \{ j   j \text{ is odd and } 2 \leq j \leq n \}$ $J_{2} = \{ j   j \text{ is even and } 2 \leq j \leq n \}$ Constrain: $f_{1} + f_{2} - a  \sin[N\pi(f_{1} - f_{2} + 1)]  - 1 \geq 0$ Search space is $[0, 1]^{n}$
CF2 [28]	$f_{1} = x_{1} + \frac{2}{ J_{1} } \sum_{j \in J_{1}} \left( x_{j} - \sin\left(6\pi x_{1} + \frac{j\pi}{n}\right) \right)^{2}$ $f_{2} = 1 - \sqrt{x_{1}} + \frac{2}{ J_{2} } \sum_{j \in J_{2}} \left( x_{j} - \sin\left(6\pi x_{1} + \frac{j\pi}{n}\right) \right)^{2}$ $\text{Constrain:}$ $\frac{t}{1 + e^{4 t }} \ge 0$ $t = \sqrt{f_{1}} + f_{2} - a \sin\left[N\pi\left(\sqrt{f_{1}} - f_{2} + 1\right)\right] - 1$ $\text{Search space is } [0, 1] \times [-1, 1]^{n-1}$
CF3 [28]	$f_{1} = x_{1} + \frac{2}{ J_{1} } \left( 4 \sum_{j \in J_{1}} y_{j}^{2} - 2 \prod_{j \in J_{1}} \cos\left(\frac{20y_{j}\pi}{\sqrt{j}}\right) + 2 \right)$ $f_{2} = 1 - x_{1}^{2} + \frac{2}{ J_{2} } \left( 4 \sum_{j \in J_{2}} y_{j}^{2} - 2 \prod_{j \in J_{2}} \cos\left(\frac{20y_{j}\pi}{\sqrt{j}}\right) + 2 \right)$ $y_{j} = x_{j} - \sin\left(6\pi x_{1} + \frac{j\pi}{n}\right), j = 2,, n$ $Constrain:$ $f_{1}^{2} + f_{2} - a \sin[N\pi(f_{1}^{2} - f_{2} + 1)] - 1 \ge 0$ $Search space is [0, 1] \times [-2, 2]^{n-1}$
CF4 [28]	$f_{1} = x_{1} + \sum_{j \in J_{1}} h_{j}(y_{j})$ $f_{2} = 1 - x_{1} + \sum_{j \in J_{2}} h_{j}(y_{j})$ $y_{j} = x_{j} - sin\left(6\pi x_{1} + \frac{j\pi}{n}\right), j = 2,, n$ $h_{2}(t) = \begin{cases}  t  & \text{if } t < \frac{3}{2}\left(1 - \frac{\sqrt{2}}{2}\right) \\ 0.125 + (t - 1)^{2} & \text{otherwise} \end{cases}$ and $h_{j}(t) = t^{2}$ $\text{Constrain:}$ $\frac{t}{1 + e^{4 t }} \ge 0$ $t = x_{2} - sin\left(6\pi x_{1} + \frac{2\pi}{n}\right) - 0.5x_{1} + 0.25$ $Search space is [0, 1] \times [-2, 2]^{n-1}$
CF5 [28]	$f_1 = x_1 + \sum_{j \in J_1} h_j(y_j)$

Test case	Objective functions		
	$f_2 = 1 - x_1 + \sum_{i \in I} h_i(y_i)$		
	$(x_i - 0.8x_1 \cos(6\pi x_1 + \frac{j\pi}{2})) \in I_1$		
	$y_{j} = \begin{cases} x_{j} - 0.8x_{1}cos\left(6\pi x_{1} + \frac{j\pi}{n}\right)j \in J_{1} \\ x_{j} - 0.8x_{1}sin\left(6\pi x_{1} + \frac{j\pi}{n}\right)j \in J_{1} \end{cases}$		
	_		
	$h_2(t) = \begin{cases}  t  & \text{if } t < \frac{3}{2} \left( 1 - \frac{\sqrt{2}}{2} \right) \\ 0.125 + (t - 1)^2 & \text{otherwise} \end{cases}$		
	$(0.125 + (t - 1)^2 \text{ otherwise})$ and $h_j(t) = 2t^2 - \cos(4\pi t) + 1$		
	Constrain:		
	$x_2 - 0.8x_1 \sin\left(6\pi x_1 + \frac{2\pi}{n}\right) - 0.5x_1 + 0.25 \ge 0$		
	Search space is $[0,1] \times [-2,2]^{n-1}$		
	$f_1 = x_1 + \sum_{j \in J_1} y_j^2$		
	$f_2 = (1 - x_1)^2 + \sum_{i \in I} y_i^2$		
	J=J <sub>2</sub>		
	$y_{j} = \begin{cases} x_{j} - 0.8x_{1}cos\left(6\pi x_{1} + \frac{j\pi}{n}\right)j \in J_{1} \\ x_{j} - 0.8x_{1}sin\left(6\pi x_{1} + \frac{j\pi}{n}\right)j \in J_{1} \end{cases}$		
CF6 [28]	$\left(x_j - 0.8x_1 \sin\left(6\pi x_1 + \frac{1}{n}\right) j \in J_1\right)$ Constrains:		
[20]	$x_2 - 0.8x_1 sin\left(6\pi x_1 + \frac{2\pi}{n}\right) - sign(k_1)\sqrt{ k_1 } \ge 0$		
	$x_{2} = 0.0x_{1}sin\left(0.0x_{1} + \frac{1}{n}\right) = sign(k_{1})\sqrt{ k_{1} } \ge 0$ $x_{4} = 0.8x_{1}sin\left(6\pi x_{1} + \frac{4\pi}{n}\right) - sign(k_{2})\sqrt{ k_{2} } \ge 0$		
	$k_1 = 0.5(1 - x_1) - (1 - x_1)^2$ $k_2 = 0.5(1 - x_1) - (1 - x_1)^2$		
	$k_1 = 0.25\sqrt{1 - x_1} - 0.5(1 - x_1)$ $k_2 = 0.25\sqrt{1 - x_1} - 0.5(1 - x_1)$		
	Search space is $[0,1] \times [-2,2]^{n-1}$		
	$f_1 = x_1 + \sum_{i=1}^n h_j(y_j)$		
	$f_2 = (1 - x_1)^2 + \sum_{i \in I_1} h_i(y_i)$		
	J C J 2		
	$y_{i} = \begin{cases} x_{j} - \cos\left(6\pi x_{1} + \frac{J\pi}{n}\right) j \in J_{1} \\ \vdots \end{cases}$		
	$y_j = \left\{ x_j - \sin\left(6\pi x_1 + \frac{j\pi}{n}\right) j \in J_1 \right\}$		
CF7 [28]	$h_2(t) = h_4(t) = t^2$ and $h_i(t) = 2t^2 - \cos(4\pi t) + 1$		
	Constrains:		
	$x_2 - \sin\left(6\pi x_1 + \frac{2\pi}{n}\right) - \operatorname{sign}(k_1)\sqrt{ k_1 } \ge 0$		
	$x_4 - \sin\left(6\pi x_1 + \frac{4\pi}{n}\right) - \operatorname{sign}(k_2)\sqrt{ k_2 } \ge 0$		
	$k_1 = 0.5(1 - x_1) - (1 - x_1)^2$		
	$k_2 = 0.25\sqrt{1 - x_1} - 0.5(1 - x_1)$		
	Search space is $[0,1] \times [-2,2]^{n-1}$		

# III. RESULTS

The benchmarking is performed for different unconstrained two-objective functions, UF1-UF5 and UF7 shown in Table II, with 30 variables each and for constrained

problems, CF1-5 with 1 constraint and CF6-7 with 2 constraints shown in Table III, with 10 variables each. For each function 30 separate runs have been performed, the stopping criteria is 300,000 function evaluations for each run, and the 100 best solutions are considered for the purpose of the IGD calculation; the average IGD value is presented in order to follow CEC'09 rules [20].

The results for UF1-5 and 7 are found and the Pareto fronts are illustrated for the UF1, Fig 1, and for the UF2 function Fig 2. UF1 and UF2 functions are chosen as they provide results representative of the worst and the best cases for the unconstrained functions.

For UF1 function the Pareto fronts show that all the variants were able to reach the true Pareto front but were unable to find a range of values. MLS2 and MLS2R found a wider variety of points along the true Pareto front than MLS1 although most of them were far from it.

Better performance can be observed on the UF2 problem, Fig. 2. In this benchmark all variants, were able to reach the front and to develop a diverse range of points. However, even in the best case of MLS2R, only half of the front is explored.

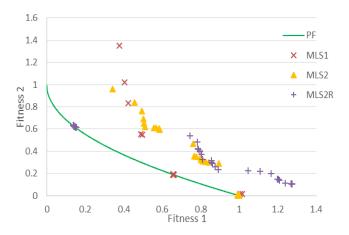


Fig. 1. Pareto front of MLSGA variants on UF1 function

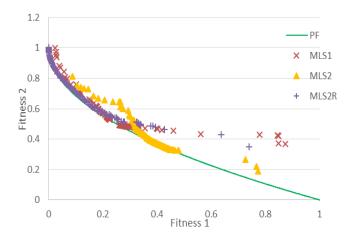


Fig. 2. Pareto front of MLSGA variants on UF2 function

The process is repeated for the constrained CF1-7 functions. CF1 Fig 3, and CF3, Fig 4, are chosen as they illustrate the worst and the best cases for the constrained test set.

The obtained results show that all MLSGA variants perform similarly; they were all able to reach the true Pareto front, and find a diverse range of points. However, the algorithm lacks accuracy for the border values of Pareto front, and concentrates its search on the middle.

The worst performance is observed on the CF3 test instance, Fig 4, where MLSGA was not able to reach the front, and only gets close to it in the best case. For this function, as for the other CF functions, all variants perform similarly.

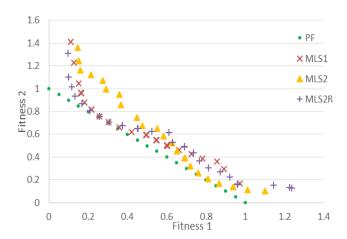


Fig. 3. Pareto front of MLSGA variants on CF1 function

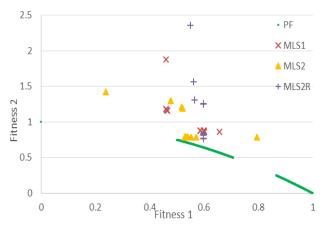


Fig. 4. Pareto front of MLSGA variants on CF3 function

The results of the three variants of MLSGA are compared to each other and to the other algorithms from CEC '09 [20]. The rankings are presented for unconstrained test cases in Table IV for UF1-3, in Table V for UF 4,5 and 7, and for constrained functions CF1-4, Table VI, and CF5-7, Table VII.

TABLE IV. MLSGA IN CEC'09 RANKING ON TWO-OBJECTIVE UNCONSTRAINED UF1-3 PROBLEMS

	Name/Average IGD			
Rank	UF1	UF2	UF3	
1	MOEA/D	MTS	MOEA/D	
	0.00435	0.00615	0.00742	
2	GDE3	MOEA/D-GM	LiuLi	
	0.00534	0.0064	0.01497	
3	MOEA/D-GM	DMOEA-DD	DMOEA-DD	
	0.0062	0.00679	0.03337	
4	MTS	MOEA/D	MOEA/D-GM	
	0.00646	0.00679	0.049	
5	LiuLi	OW-MOSaDE	MTS	
	0.00785	0.0081	0.0531	
6	DMOEA-DD 0.01038	GDE3 0.01195	Clustering MOEA 0.0549	
7	NSGA II LS	LiuLi	AMGA	
	0.01153	0.0123	0.06998	
8	OW-MOSaDE	NSGA II LS	DECMOSA-SQP	
	0.0122	0.01237	0.0935	
9	Clustering MOEA 0.0299	AMGA 0.01623	MOEP 0.099	
10	AMGA	MOEP	OW-MOSaDE	
	0.03588	0.0189	0.103	
11	MOEP 0.0596	Clustering MOEA 0.0228	NSGA II LS 0.10603	
12	DECMOSA-SQP	DECMOSA-SQP	GDE3	
	0.07702	0.02834	0.10639	
13	OMOEA II	OMOEA II	OMOEA II	
	0.08564	0.03057	0.27141	
14	MLS2R	MLS2	MLS1	
	0.205	0.105	0.401	
15	MLS1	MLS1	MLS2R	
	0.213	0.156	0.438	
16	MLS2	MLS2R	MLS2	
	0.223	0.176	0.447	

TABLE V. MLSGA IN CEC'09 RANKING ON TWO-OBJECTIVE UNCONSTRAINED UF4-7 PROBLEMS

Rank	Name/Average IGD			
капк	UF4	UF5	UF7	
1	MTS	MTS	MOEA/D	
	0.02356	0.01489	0.00587	
2	GDE3	GDE3	LiuLi	
	0.0265	0.03928	0.0073	
3	DECMOSA-SQP	AMGA	MOEA/D-GM	
	0.03392	0.09405	0.0076	
4	AMGA	LiuLi	DMOEA-DD	
	0.04062	0.16186	0.01032	
5	DMOEA-DD	DECMOSA-SQP	MOEP	

Rank	Name/Average IGD			
Kank	UF4	UF5	UF7	
	0.04268	0.16713	0.0197	
6	MOEP	OMOEA II	NSGA II LS	
	0.0427	0.1692	0.02132	
7	LiuLi 0.0435	MOEA/D 0.18071	Clustering MOEA 0.0223	
8	OMOEA II	MOEP	DECMOSA-SQP	
	0.04624	0.2245	0.02416	
9	MOEA/D-GM 0.0476	Clustering MOEA 0.2473	GDE3 0.02522	
10	OW-MOSaDE	DMOEA-DD	OMOEA II	
	0.0513	0.31454	0.03354	
11	NSGA II LS	OW-MOSaDE	MTS	
	0.0584	0.4303	0.04079	
12	Clustering MOEA 0.0585	NSGA II LS 0.5657	AMGA 0.05707	
13	MOEA/D	MLS2	OW-MOSaDE	
	0.06385	0.696	0.0585	
14	MLS2R	MLS1	MLS1	
	0.0881	0.733	0.398	
15	MLS1	MLS2R	MLS2	
	0.0882	0.767	0.410	
16	MLS2	MOEA/D-GM	MLS2R	
	0.103	1.7919	0.412	

The unconstrained rankings show that all the MLSGA variants are outperformed by all other algorithms on almost all of the functions, except UF5 where MOEA/D-GM is the worst; additionaly in this case the MLS2 variant is not far from the NSGA II LS. Interesingly all MLS variants show similar performance in all tests, and it's hard to distinguish which one is better, which is in contradiction with previous research on the simpler ZDT functions, where on these simpler functions MLS2R variant strongly outperforms the others and MLS1 provides poor diversity of points.

For the constrained ranking, the results are more promising. In this case MLSGA variants are outperformed by all other algorithms only on the CF1 and CF4 benchmark. However, MLS2R is better than both DECMOSA and MOEA/D-GM on CF3 and CF5; MLS1 outperforms these algorithms on CF5 and CF6. Again all the MLSGA variants have similar performance, but MLS1 is slightly better, which is surprising in comparison to the unconstrained ZDT functions, where the MLS1 algorithm lacks the diversity of points to get a low IGD value.

The detailed Pareto front results show that MLSGA is unable to properly find the whole Pareto front, however it is able to reach it and explore it partially, providing promising results at this stage, especially for constrained problems. The current algorithm only implements simple genetic algorithm mechanics at the individual level and the collective level

mechanics are at an early stage of development but is already outperforming some leading algorithms on a range of stateof-the-art problems.

TABLE VI. MLSGA IN CEC'09 RANKING ON TWO-OBJECTIVE CONSTRAINED CF1-4 PROBLEMS

Rank	Name/Average IGD			
Kank	CF1	CF2	CF3	CF4
	LiuLi	DMOEADD	DMOEADD	DMOEADD
1	0.00085	0.0021	0.056305	0.00699
2	NSGAIILS	LiuLi	MTS	GDE3
2	0.00692	0.0042	0.10446	0.00799
3	MEOADGM	MEOADGM	GDE3	MTS
3	0.0108	0.008	0.127506	0.01109
4	DMOEADD	NSGAIILS	LiuLi	LiuLi
4	0.01131	0.01183	0.182905	0.01423
5	MTS	GDE3	NSGAIILS	NSGAIILS
3	0.01918	0.01597	0.23994	0.01576
-	GDE3	MTS	MLS2R	MEOADGM
6	0.0294	0.02677	0.435	0.0707
-	MLS2R	DECMOSA	MEOADGM	DECMOSA
7	0.0872	0.0946	0.5134	0.15265
8	MLS1	MLS1	MLS1	MLS1
8	0.0933	0.160	0.522	0.332
9	MLS2	MLS2R	MLS2	MLS2R
	0.0996	0.164	0.633	0.348
1.0	DECMOSA	MLS2	DECMOSA	MLS2
10	0.10773	0.1829	1000000	0.387

TABLE VII. MLSGA IN CEC'09 RANKING ON TWO-OBJECTIVE CONSTRAINED CF5-7PROBLEMS

Rank	Name/Average IGD		
Kank	CF5	CF6	UF7
1	DMOEADD	LiuLi	DMOEADD
1	0.01577	0.013948	0.01905
2	MTS	DMOEADD	MTS
2	0.02077	0.01502	0.02469
2	GDE3	MTS	GDE3
3	0.06799	0.01616	0.04169
4	LiuLi	NSGAIILS	LiuLi
4	0.10973	0.02013	0.10446
5	NSGAIILS	GDE3	NSGAIILS
3	0.1842	0.06199	0.23345
	MLS1	MLS1	DECMOSA
6	0.390	0.141	0.26049
7	MLS2R	DECMOSA	MLS2R
	0.393	0.14782	0.522116
0	DECMOSA	MEOADGM	MEOADGM
8	0.41275	0.2071	0.5356
9	MLS2	MLS2	MLS1
	0.426	0.502	0.538741
10	MEOADGM	MLS2R	MLS2
	0.5446	2.746	0.571981

#### IV. DISCUSSION

The results show that MLSGA provides promising results on complex two-objective problems, representing the current state-of-art. On all the test cases, each MLSGA variant has a similar performance and is able to reach the true Pareto front, in some cases with a good diversity of points. This increases the performance of the classic genetic algorithm, on which the current version of MLSGA is based, to compete with current state-of-the-art genetic algorithms. Lack of a substantial difference between MLS2 and MLS2R was predictable as all UF and CF test cases have similar complexity for both  $f_1$  and  $f_2$ . This is different from the ZDT functions tested on previously where performance between the variants showed a large difference, but these functions are less well balanced between objectives. More interesting is the similarity in performance between MLS1 and MLS2 variants, which also differs from the previous results. The authors suggest that the reason behind this is that the UF test problems are too complex for the current, simple version of MLSGA and both variants are struggling in finding the proper solutions. MLS1 results have previously shown poor diversity of solutions whereas MLS2 finds a diverse range of solutions once the front is found. In these cases, the front is more difficult to find so the MLS2 variant is not able to create a spread of results, showing similar performance to MLS1.

MLSGA is based on simple principles and thus further work is necessary. Possible methods to improve the performance are to develop the new collective evolutionary mechanism, which is underdeveloped compared to individual level mechanisms, and employ better performing methods than the simple classic GA mechanisms. The authors suggest to incorporate top individual mechanisms, such as MOEA/D, NSGAII and DMOEA-DD into MLSGA for even greater gains in performance. Other approaches could be implementation of local search methods such as MTS into collective reproduction mechanisms, and the introduction of different MLS types and fitness definitions into each collective, where some collectives in one run can utilise MLS1 and others MLS2 or MLS2R, for wider area searches. Additionally, the authors suggest that the MLSGA approach may be more successful on other kind of problems including: multi-level functions, noisy and dynamic problems, where the split in fitness function and population may lead to increased performance.

## V. CONCLUSION

There are a number of different approaches to improve performance of genetic algorithms. This paper implements a novel bio-inspired mechanism, based on modern theories of evolution, multi-level selection. A simple genetic algorithm is used at the individual level and an initial, novel, collective level mechanism is implemented; these are benchmarked on CEC'09 test problems and compared with current state-of-art competitors. Interestingly all variants are able to reach the true Pareto front and show similar performance to each other.

The developed algorithm needs further improvements, however the current mechanisms are simple and it is proposed to introduce more complex individual reproduction from the current state-of-art to improve performance.

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