

A benchmark for cooperative coevolution

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Abstract Cooperative co-evolution algorithms (CCEA) are a thriving sub-field of evolutionary computation. This class of algorithms makes it possible to exploit more efficiently the artificial Darwinist scheme, as soon as an optimisation problem can be turned into a co-evolution of interdependent sub-parts of the searched solution. Testing the efficiency of new CCEA concepts, however, it is not straightforward: while there is a rich literature of benchmarks for more traditional evolutionary techniques, the same does not hold true for this relatively new paradigm. We present a benchmark problem designed to study the behavior and performance of CCEAs, modeling a search for the optimal placement of a set of lamps inside a room. The relative complexity of the problem can be adjusted by operating on a single parameter. The fitness function is a trade-off between conflicting objectives, so the performance of an algorithm can be examined by making use of different metrics. We show how three different cooperative strategies, Parisian Evolution, Group Evolution and Allopatric Group Evolution, can be applied to the problem. Using a Classical Evolution approach as comparison, we analyse the behavior of each algorithm in detail, with respect to the size of the problem.

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1 Introduction

Cooperative co-evolution algorithms (CCEAs) share common characteristics with standard artificial Darwinist methods, i.e. evolutionary algorithms (EAs), but with additional components that aim at implementing collective capabilities. For optimisation purpose, CCEAs are based on a specific formulation of the problem where various inter- or intra-population interaction mechanisms occur. Usually, these techniques are efficient as optimisers when the problem can be split into smaller interdependent subproblems. The computational effort is then distributed onto the evolution of smaller elements of similar or different nature, that aggregates to build a global solution.

Besides precursory research lines, focused on co-evolution as a complex phenomenon leading to stable or unstable equilibria (complex systems, multi-agent systems) [3], first attempts to exploit a cooperative co-evolution paradigm for optimisation purposes have been made by Husbands and Mill [14], Potter and De Jong [22] and Ahluwalia and Bull [1]. These approaches were heterogeneous, in the sense that co-evolution is implemented between separated subpopulations.

A few years later, inspired by the Michigan approach (John Holland's classifier systems [13]), a single-population co-evolution scheme, first called “Individual-GP” [9], then “Parisian Scheme”, was developed in France for complex optimisation tasks [10].

Cooperative co-evolution is increasingly becoming the basis of successful applications [2, 8, 11, 21, 27], including

learning [5] and scheduling problems [14]. These approaches can be shared into two main categories: heterogeneous co-evolution that happens between a fixed number of separate populations [7, 19, 20], and homogeneous co-evolution, that occurs within a single population [16, 26, 28].

The design and fine tuning of such algorithms remain however difficult and strongly problem dependent. A critical question is the design of simple test problem for CCEAs, for benchmarking purpose. A first test-problem based on Royal Road Functions has been proposed in [18]. A second example is the NK-landscapes variant redesigned for CCEAs, known as NKC-landscapes [15]: however, Kauffman's work was more focused on the study of the dynamics of coevolution than on evolution as an optimization process.

We propose another simple problem, the *lamps problem*. The main difference with previous benchmarks is the absence of a priori information on the optimal number of individuals/partial solutions that should compose the global solution. As for many real-world problems, the number of individuals in the best group is unknown. Moreover, various instances of the problem, of increasing complexity can be generated by acting on a single ratio parameter. For these reasons we feel that the proposed benchmark, while simple to implement and calibrate, can still pose a challenge to cooperative coevolution algorithms. We show below how three CCEAs can be designed and compared against a classical approach, with a special focus on their behaviour with respect to the size of the problem. Preliminary results on this topic have been presented in [25].

The paper is organised as follows: the lamps problem is described in Sect. 2, then the design of three cooperative coevolution strategies, Parisian Evolution, Group Evolution, and Allopatric Group Evolution, is detailed in Sects. 3, 4 and 5, respectively. The experimental setup is described in Sect. 6: four strategies are tested, a classical genetic programming approach (CE for Classical Evolution), the Parisian Evolution (PE), the Group Evolution (GE) and the Allopatric Group Evolution (AGE). All methods are implemented using the μ GP toolkit [23]. Results are presented and analysed in Sect. 7, and conclusions and future work are given in Sect. 8.

2 The lamps problem

The optimisation problem chosen to test cooperative coevolution algorithms requires to find the best placement for a set of lamps, so that a target area is fully brightened with light. The minimal number of lamps needed is unknown, and heavily depends on the topology of the area. Lamps are all alike, modeled as circles, and each one may be evaluated separately with respect to the final goal. In the example, the optimal solution requires four lamps (Fig. 1, left): interestingly, when examined independently, all lamps in the solution

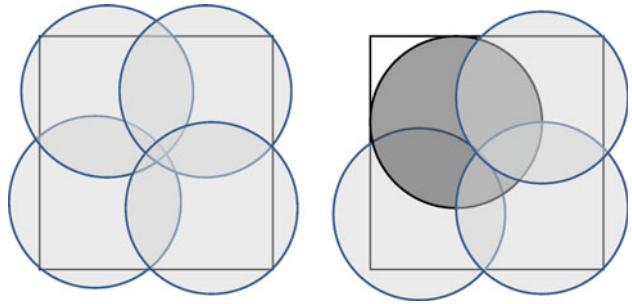


Fig. 1 Placement of a set of lamps. The aim is to enlighten all the square area. It is interesting to notice how a solution where some of the light of each lamp is wasted outside the area (*left*) overall performs better than a solution where the grayed lamp maximises its own performance (*right*)

waste a certain amount of light outside the target area. However, if one of the lamps is positioned to avoid this undesired effect, it becomes impossible to lighten the remaining area with the three lamps left (Fig. 1, right). Since lamps are simply modeled as circles, the problem may also be seen as using the circles to completely cover the underlying area, as efficiently as possible.

This apparently simple benchmark exemplifies a common situation in real-world applications: many problems have an optimal solution composed of a set of homogeneous elements, whose individual contribution to the main solution can be evaluated separately. Note that, in this context, *homogeneous* is used to label elements sharing the same base structure. Not only, but often the optimal solution is composed of not-optimal sub-solutions.

A similar toy problem has been sketched in [24]. Here the structure of the benchmark is improved and parametrised, and a modified fitness function increases the complexity and the number of local optima on the fitness landscape.

2.1 Size of the problem

It is intuitive that the task of enlightening a room with a set of lamps can be more or less difficult, depending on the size of the room and the cone of light of each lamp. If small sources of light are used to brighten a large room, surely a greater number of lamps will be required, and the number of possible combinations will increase dramatically.

With these premises, the complexity of the problem can thus be expressed by the ratio between the surface to be enlightened and the maximum area enlightened by a single lamp:

$$\text{problem_size} = \frac{\text{area_room}}{\text{area_lamp}}$$

as this ratio increases, finding an optimal solution for the problem will become harder.

It is interesting to notice how variations in the shape of the room could also influence the complexity of the task: rooms with an irregular architecture may require more intricate lamp placements. However, finding a dependency between the shape of the room and the difficulty of the problem is not trivial, and results might be less intuitive to analyze. Also, the cooperative coevolution algorithms do not search for or benefit from eventual symmetries in the problem. For all these reasons, the following experiments will feature square rooms only.

2.2 Fitness value

Comparing different methodologies on the same problem requires a common fitness function, to be able to numerically evaluate the efficiency of each approach.

Intuitively, the fitness of a candidate solution should be directly proportional to the area fully brightened by the lamps and inversely proportional to the number of lamps used, favoring solutions that cover more surface with light using the minimal number of lamps. The first term in the fitness value will thus be proportional to the ratio of the area enlightened by the lamps,

$$\frac{\text{area_enlightened}}{\text{total_area}}$$

A further contribution is added, to help experts in evaluating the goodness of the grouping performed by the algorithms: the area brightened by more than one lamp is to be minimised, in order to have as little overlapping as possible. The second term will be then proportional to:

$$-\frac{\text{area_overlap}}{\text{total_area}}$$

It is interesting to note that minimising the overlap also implies an optimisation of the number of lamps used, since using a greater number would lead to more overlapping areas.

The final fitness function will then be:

$$\begin{aligned} \text{fitness} &= \frac{\text{area_enlightened}}{\text{total_area}} - W \cdot \frac{\text{area_overlap}}{\text{total_area}} \\ &= \frac{\text{area_enlightened} - W \cdot \text{area_overlap}}{\text{total_area}} \end{aligned}$$

where W is a weight associated to the relative importance of the overlapping, set by the user.

Using this function with $W = 1$, fitness values will range between $(0, 1)$, but it is intuitive that it is impossible to reach the maximum value: by problem construction, overlapping and enlightenment are inversely correlated, and even good solutions will feature overlapping areas and/or parts not enlightened. This problem is actually multi-objective, the fitness function we propose corresponds to a compromise between the two objectives.

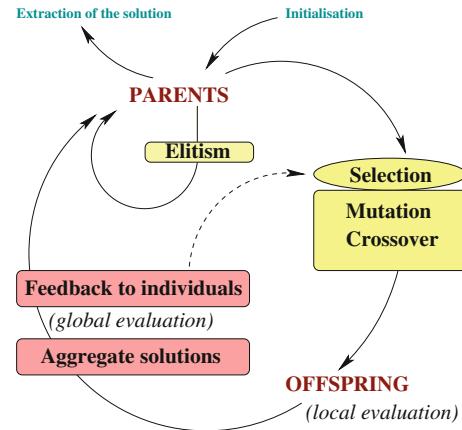


Fig. 2 A Parisian EA: a monopopulation cooperative-coevolution. Partial evaluation (local fitness) is applied to each individual, while global evaluation is performed once a generation

3 Parisian Evolution

Initially designed to address the inverse problem for Iterated Function System (IFS), a problem related to fractal image compression [10], this scheme has been successfully applied in various real world applications: in stereovision [6], in photogrammetry [12, 17], in medical imaging [26], for Bayesian Network Structure learning [4], in data retrieval [16].

Parisian Evolution (PE) is based on a two-level representation of the optimisation problem, meaning that an individual in a Parisian population represents only a part of the solution. An aggregation of multiple individuals must be built to complete a meaningful solution to the problem. This way, the co-evolution of the whole population (or of a major part of it) is favoured over the emergence of a single best individual, as in classical evolutionary schemes.

This scheme distributes the workload of solution evaluations at two levels. Light computations (e.g. existence conditions, partial or approximate fitness) can be done at the individual's level (local fitness), while the complete calculation (i.e. global fitness) is performed at the population level. The global fitness is then distributed as a bonus to individuals who participate to the global solution. A Parisian scheme has all the features of a classical EA (see Fig. 2) with the following additional components:

- A grouping stage at each generation, that selects individuals that are allowed to participate to the global solution.
- A redistribution step that rewards the individuals who participate to the global solution: their bonus is proportional to the global fitness.
- A sharing scheme, that avoids degenerate solutions where all individuals are identical.

Efficient implementations of the Parisian scheme are often based on partial redundancies between local and global

fitness, as well as clever exploitation of computational shortcuts. The motivation is to make a more efficient use of the evolution of the population, and reduce the computational cost. Successful applications of such a scheme usually rely on a lower cost evaluation of the partial solutions (i.e. the individuals of the population), while computing the full evaluation only once at each generation or at specified intervals.

3.1 Implementation of the lamps problem

For the lamps problem, the PE has been implemented as follows. An individual represents a lamp: its genome is its position, a pair of real values (x, y), plus a third element, e , that can assume values 0 or 1 (on/off switch). Lamps with $e = 1$ are “on” (expressed) and contribute to the global solution, while lamps with $e = 0$ do not.

Global fitness is computed as described in Sect. 2.2. In generation 0 the global solution is computed simply considering the individuals with $e = 1$ among the μ initial ones. Then, at each step, λ individuals are generated. For each new individual with $e = 1$, its potential contribution to the global solution is computed. Before evaluation, a random choice ($p = 0.5$) is performed: new individuals are either considered in addition to or in replacement of the existing ones.

If individuals are considered for addition, the contribution to the global solution of each one is computed. Otherwise, the less performing among the old individuals is removed, and only then the contribution to the global solution of each new individual is evaluated. If the addition or replacement of the new individuals leads to an improvement over the previous global fitness, the new individual selected is rewarded with a high local fitness value ($local_fitness = 2$), together with all the old individuals still contributing to the global solution. New expressed individuals ($e = 1$) that are not selected for the global solution are assigned a low fitness value ($local_fitness = 0$). Non-expressed individuals ($e = 0$) have an intermediate fitness value ($local_fitness = 1$).

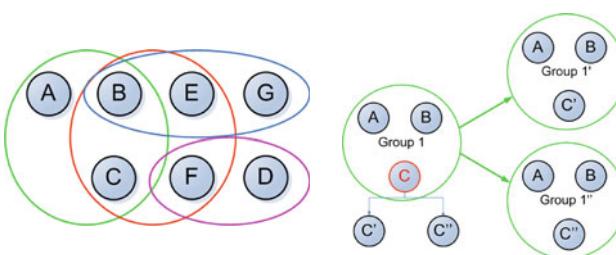


Fig. 3 *Left* individuals and groups in a sample population of eight individuals. While individual A is part of only one group, Individual B is part of three different groups. The effect of an individual genetic operator, applied to individual C. Since individual C is part of Group 1, two groups are created and added to the population. *Right* schema of Group

Sharing follows the simple formula

$$fitness_sharing(I_k) = \frac{local_fitness(I_k)}{\sum_{i=0}^{individuals} sharing(I_k, I_i \neq k)}$$

with

$$sharing(I_1, I_2)$$

$$= \begin{cases} 1 - \frac{d(I_1, I_2)}{2 \cdot lamp_radius} & d(I_1, I_2) < 2 \cdot lamp_radius \\ 0 & d(I_1, I_2) \geq 2 \cdot lamp_radius \end{cases}$$

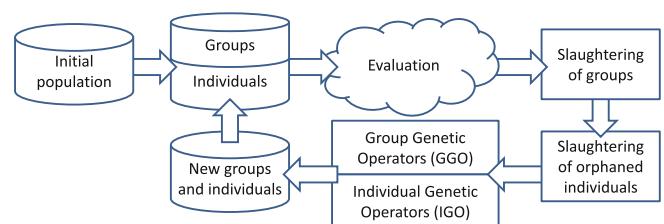
Lamps that have a relatively low number of neighbours will be preferred for selection over lamps with a bigger number of neighbours. In this implementation, sharing is computed only for expressed lamps ($e = 1$) and used only when selecting the less performing individual to be removed from the population.

4 Group Evolution

Group Evolution (GE) is a novel generational cooperative coevolution concept presented in [24]. The approach uses a population of partial solutions, and exploits non-fixed sets of individuals called *groups*. GE acts on individuals and groups, managing both in parallel. During the evolution, individuals are optimised as in a common EA, but concurrently groups are also evolved. The main peculiarity of GE is the absence of a priori information about the grouping of individuals.

At the beginning of the evolutionary process, an initial population of individuals is randomly created on the basis of a high-level description of a solution for the given problem. Groups at this stage are randomly determined, so that each individual can be included in any number of different groups, but all individuals are part of at least one group.

The number of groups μ_{groups} , the minimum and maximum size of the groups are set by the user before the evolution starts. The number of individuals in the population is a direct consequence of the specified values for groups. Figure 3 (left)



Evolution algorithm. Groups are sets of individuals. At each step, new groups and individuals are produced. The slaughtering is performed at the level of groups, and in a second step individuals that are not included in any group are also removed

shows a sample population where minimum group size is 2, and maximum group size is 4.

4.1 Generation of new individuals and groups

GE exploits a generational approach: at each evolutionary step, a number of genetic operators is applied to the population. Genetic operators can act on both individuals and groups, and produce a corresponding offspring, in form of individuals and groups.

The offspring creation phase comprehends two different actions at each generation step (see Fig. 3):

1. Application of *group genetic operators*.
2. Application of *individual genetic operators*.

Each time a genetic operator is applied to the population, parents are chosen and offspring is generated. The children are added to the population, while the original parents are unmodified. Offspring is then evaluated, while it is not compulsory to reconsider the fitness value of the parents again. It is important to notice that the number of children produced at each evolutionary step is not fixed: each genetic operator can have any number of parents as input and produce in output any number of new individuals and groups. The number and type of genetic operators applied at each step can be set by the user.

4.1.1 Group genetic operators

Group genetic operators (GGOs) work on the set of groups. Each operator needs a certain number of groups as parents and produces a certain number of groups as offspring that will be added to the population. GGOs implemented in our approach are:

1. *crossover*: generates offspring by selecting two individuals, one from parent group A and one from parent group B. Those individuals are switched, creating two new groups;
2. *add-mutation*: generates offspring by selecting one or more individuals from the population and a group. Chosen individuals are added (if possible) to the parent group, creating a single new group;
3. *removal-mutation*: generates offspring by selecting a group and one or more individuals inside it. Individuals are removed from the parent group.
4. *replacement-mutation*: generates offspring by selecting a group and one or more individuals inside it. Individuals are removed from the parent group, and replaced by other individuals selected from the population.

Parent groups are chosen via tournament selection.

4.1.2 Individual genetic operators

Individual genetic operators (IGOs) operate on the population of individuals, very much like they are exploited in usual GA. The novelty of GE is that for each individual produced as offspring, new groups are added to the group population. For each group the parent individual was part of, a copy is created, with the offspring taking the place of the parent.

This approach, however, could lead to an exponential increase in the number of groups, as the best individuals are selected by both GGOs and IGOs. To keep the number of groups under a strict control, we choose to create a copy only of the highest-fitness groups the individual was part of.

IGOs select individuals by a tournament selection in two parts: first, a group is picked out through a tournament selection with moderate selective pressure; then an individual in the group is chosen with low selective pressure. The actual group and the highest-fitness groups the individual is part of are cloned once for each child individual created: in each clone group the parent individual is replaced with a child. An example is given in Fig. 3 (right): an IGO, selects individual C as a parent. The chosen individual is part of only one group, Group 1. The IGO produces two children individuals: since the parent was part of a group, a new group is created for each new individual generated. The new groups (Group 1' and Group 1'') are identical to Group 1, except that individual C is replaced with one of its children, C' in Group 1' and C'' in Group 1'', respectively.

The aim of this process is to select individuals from well-performing groups to create new groups with a slightly changed individual, in order to explore the a near area in the solution space.

4.2 Evaluation

During the evaluation phase, a fitness value is associated to each group: the fitness value is a number that measures the goodness of the candidate solutions with respect to the given problem. When a group is evaluated, a fitness value is also assigned to all the individuals composing it. Those values reflect the goodness of the solution represented by the single individual and have the purpose to help discriminate during tournament selection for both IGOs and GGOs.

An important strength of the approach resides in the evaluation step: if there is already a fitness value for an individual that is part of a new group, it is possible to take it into account instead of re-evaluating all the individuals in the group. This feature can be exceptionally practical when facing a problem where the evaluation of a single individual can last several minutes and the fitness of a group can be computed without examining simultaneously the performance of the individuals

composing it. In that case, the time-wise cost of both IGOs and GGOs becomes very small.

4.3 Slaughtering

After each generation step, the group population is resized. The groups are ordered fitness-wise and the worst one is deleted until the desired population size is reached. Every individual keeps track of all the groups it belongs to in a set of references. Each time a group ceases to exist, all its individuals remove it from their set of references. At the end of the group slaughtering step, each individual that has an empty set of references, and is therefore not included in any group, is deleted as well.

4.4 Implementation of the lamps problem

For the lamps problem, GE has been implemented as follows. One individual represents a lamp: its genome is its position, a pair of real values (x, y). The fitness of a single individual, which must be independent from all groups it is part of, is simply the area of the room it enlightens. The group fitness is computed as described in Sect. 2.2.

5 Allopatric Group Evolution

The third proposed cooperative coevolution algorithm stems from the Group Evolution presented in the previous section. While the flow of the algorithm is the same, however, two major improvements distinguish the Allopatric Group Evolution from the previously described methodology: allopatric group slaughtering and heuristic group genetic operators.

5.1 Allopatric group slaughtering

Allopatric speciation (from the ancient Greek *allos*, other, and *patra*, fatherland) is a speciation that occurs when biological populations of the same species become isolated, usually due to geographical changes, such as major changes in the landscape.

Taking inspiration from this natural phenomenon, the allopatric group slaughtering temporarily isolates from the main population all the offspring of a group. Subsequently, the offspring compete between themselves, and only the best resulting group is added back into the main population. A summary of the technique is presented in Fig. 4.

This mechanism is aimed at improving diversity, trying to avoid the invasion of the main group population by the offspring of a particularly promising group, an event that could drive the evolution deep into a local optima.

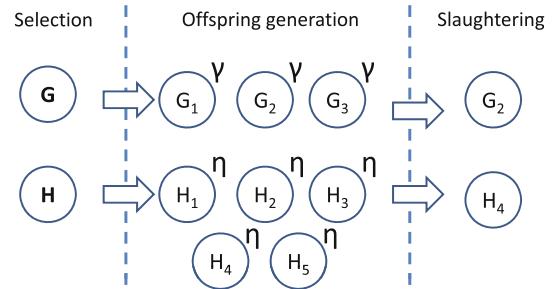


Fig. 4 Allopatric slaughtering of groups. Group G and group H are chosen for reproduction in the same iteration. The operator applied to group G creates three children groups, with the allopatric tag γ , while the operator applied to group H creates five, with tag η . Among all the offspring that share the same allopatric tag, only the best group is added to the population: in the example, the fitness value of offspring H_3 might be higher than the value of G_2 ; nevertheless, G_2 is the best group among the offspring of G

5.2 Heuristic group operators

The second important difference between AGE and GE, is the presence of heuristic group operators. The aim of the operators is to exploit expert knowledge of the problem that cannot be incorporated inside the group or individual fitness function. In the current implementation, two heuristic operators substitute removal-mutation and add-mutation presented in Sect. 4.1.1. When an individual is chosen to be removed from or added to a group, instead of a simple random selection, an external heuristic selector chooses the best candidate, according to a metric specified by the user.

Heuristic operators might prove beneficial to the evolution, especially when a run of the selector is not as computationally expensive as the evaluation of a group.

5.3 Implementation of the lamp problem

The setup is the same as in GE, with the only necessary addition of a heuristic to add or remove individuals from a group. For the lamp problem, we exploited the distance between individuals as an intuitive measurement of possible overlap. In particular, candidate individuals to be added to a group are evaluated on their distance from the nearest component of a group; while candidate individuals to be removed from a group are evaluated on the sum of the distances from the other components of the group.

6 Experimental setup

Before starting the experiments on the cooperative coevolution algorithms, a series of ten runs for each *problem_size* of a classical evolutionary algorithm is performed, to better understand the characteristics of the problem and to set parameters leading to a fair comparison. The genome of a single

individual is a set of lamps, modeled as an array of N pairs of real values $((x_1, y_1), (x_2, y_2), \dots, (x_N, y_N))$, where each (x_i, y_i) describes the position of individual i in the room. The algorithm uses a classical $(\mu + \lambda)$ evolutionary paradigm, with $\mu = 20$ and $\lambda = 10$, probability of crossover 0.2 and probability of mutation 0.8. Each run lasts 100 generations.

By examining these first experimental results, it is possible to reach the following conclusions: good solutions for the problem use a number of lamps in the range $(problem_size, 3 \cdot problem_size)$; and, as expected, as the ratio grows, the fitness value of the best individual at the end of the evolution tends to be lower.

In order to perform a comparison with PE, GE and AGE as fair as possible, the stop condition for each algorithm will be set as the average number of single lamp evaluations performed by the classical evolutionary algorithm for each $problem_size$. In this way, even if the algorithms involved have significantly different structures, the computational effort is commensurable. The number of eval-

uation ranges is 3,500 for $problem_size = 3$; 5,000 for $problem_size = 5$; 11,000 for $problem_size = 10$; 22,000 for $problem_size = 20$; and 120,000 for $problem_size = 100$.

6.1 PE setup

Due to the observation of the CE runs, $\mu = 3 \cdot problem_size$, while $\lambda = \mu/2$. The probability of mutation is 0.8 and the probability of crossover is 0.2.

6.2 GE and AGE setup

The number of groups in the population is fixed, $\mu_{groups} = 20$, as is the number of genetic operators selected at each step $\lambda = 10$. The number of individuals in each group is set to vary in the range $(problem_size, 3 \cdot problem_size)$. The probability of mutation is 0.8, while the probability of crossover is 0.2, for both individuals and groups.

Table 1 Average results for 100 runs, for each evolutionary approach

Problem size	Evolution	Avg. fitness	SD	Avg. enlight.	SD	Avg. lamps	SD	Avg. overlap	SD	Avg. lamps before acceptable	SD
3	CE	0.861	0.0149	0.8933	0.0212	4	0	0.0323	0.0168	313.2	126
	PE	0.8355	0.064	0.8945	0.0439	4.02	0.3344	0.059	0.0502	316.9	262.2
	GE	0.8764	0.0498	0.8963	0.0533	3.75	0.435	0.0198	0.0103	267.32	138.1
	AGE	0.8507	0.0263	0.8998	0.0271	4.2	0.324	0.049	0.0266	349.65	179.21
5	CE	0.7802	0.023	0.8574	0.04	6.2	0.64	0.0772	0.0278	572.7	215.38
	PE	0.7825	0.03	0.8803	0.0335	6.96	0.6936	0.0978	0.0395	511.35	373.85
	GE	0.8136	0.0241	0.8537	0.0349	6.03	0.7372	0.0401	0.0166	741.36	221.86
	AGE	0.79	0.0234	0.8875	0.0303	7.24	0.6624	0.0975	0.03189	608.32	255.75
10	CE	0.7487	0.0149	0.834	0.0235	11.3	0.62	0.0853	0.0216	1,779.8	407.4
	PE	0.7791	0.0221	0.8847	0.0207	12.84	0.8184	0.1055	0.0274	1,018.47	546.16
	GE	0.7532	0.0178	0.8132	0.0255	10.66	0.6336	0.0599	0.0215	1,836.87	412.08
	AGE	0.75	0.017	0.866	0.0229	13.09	0.8352	0.1159	0.0251	1,715.85	637.91
20	CE	0.6804	0.0117	0.7749	0.0148	20.6	0.72	0.0946	0.0123	3,934.7	702.24
	PE	0.7624	0.0147	0.8762	0.0168	23.57	1.053	0.1138	0.0177	2,759.28	965.45
	GE	0.697	0.0127	0.7837	0.0147	20.49	0.5978	0.0867	0.0142	4,602.1	1,156.5
	AGE	0.7086	0.0182	0.8523	0.0269	25.14	1.514	0.1437	0.0293	3,151.31	1,067.89
100 ^a	CE	0.558	0.0049	0.7334	0.0062	102.9	1.88	0.1755	0.0093	29,567.3	4,782.96
	PE	0.6867	0.0073	0.8708	0.0078	117.2	3.2	0.1841	0.0134	5,318.9	332.72
	GE	0.5647	0.0057	0.7309	0.0073	101.5	1.7	0.1662	0.0072	30,048	8,876.8
	AGE	0.6095	0.021	0.792	0.0314	111.4	9.2	0.1824	0.04	12,094.4	3,686.28

For $problem_size = 100$, the distribution of overlap for AGE presents such a vast standard deviation that it is not distinguishable from the other three approaches ($p > 0.05$). Taking into account the remaining distributions (GE, PE, GE), GE is the best ($p < 0.05$) for the overlap: the corresponding value in the table is reported in bold and italic to highlight this peculiarity

^a Due to the times involved, data for problem size 100 is computed on 10 runs

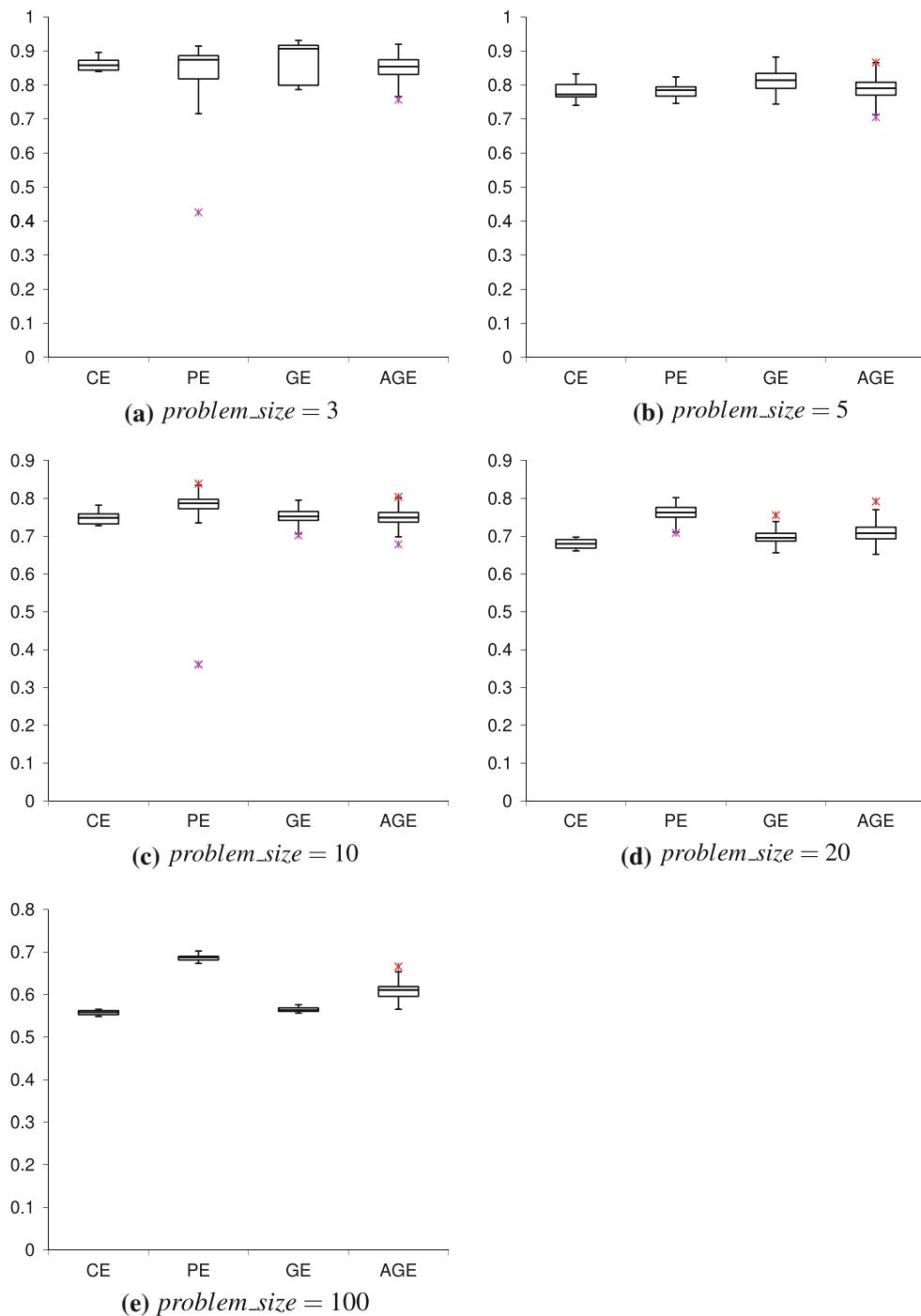


Fig. 5 Box plots of fitness values for each *problem_size*

6.3 Implementation in μ GP

The two CCEAs used in the experience have been implemented using μ GP [23], an evolutionary toolkit developed by CAD group of Politecnico di Torino. Exploiting μ GP's flexible structure, it is possible to replicate the behavior of very different EAs.

In particular, to implement PE, it is sufficient to operate on the fitness evaluator, setting the environment to evaluate the whole population and the offspring at each step. Obtaining GE behavior is slightly more complex, and requires the addition of new classes to manage groups. CE is simply μ GP standard operation mode.

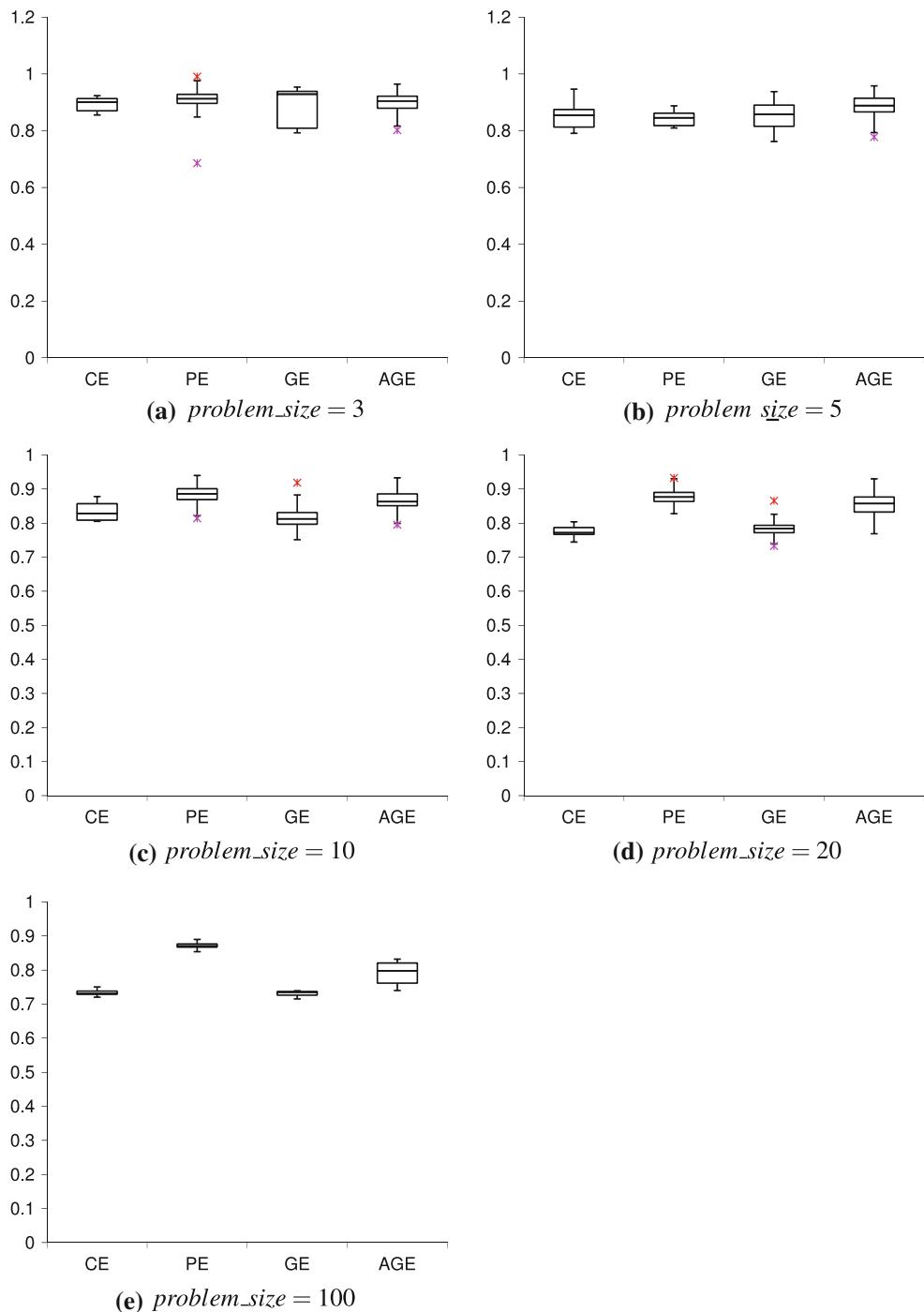


Fig. 6 Box plots of enlightenment percentage for each *problem_size*

The individual-level operators chosen for the experience are `singleParameterAlterationMutation` and `onePointCrossover`. The mutation operator acts on one or both coordinates of a single individual, modifying them according to a gaussian distribution and producing a single child. The crossover operator, on the other hand, starting from two individuals $A = (x_a, y_a)$ and $B = (x_b, y_b)$, creates

two children individuals $AB = (x_a, y_b)$ and $BA = (x_b, y_a)$. For further information, see [23].

7 Results and analysis

In a series of experiments, 100 runs of each evolutionary approach are executed, for a set of meaningful values of

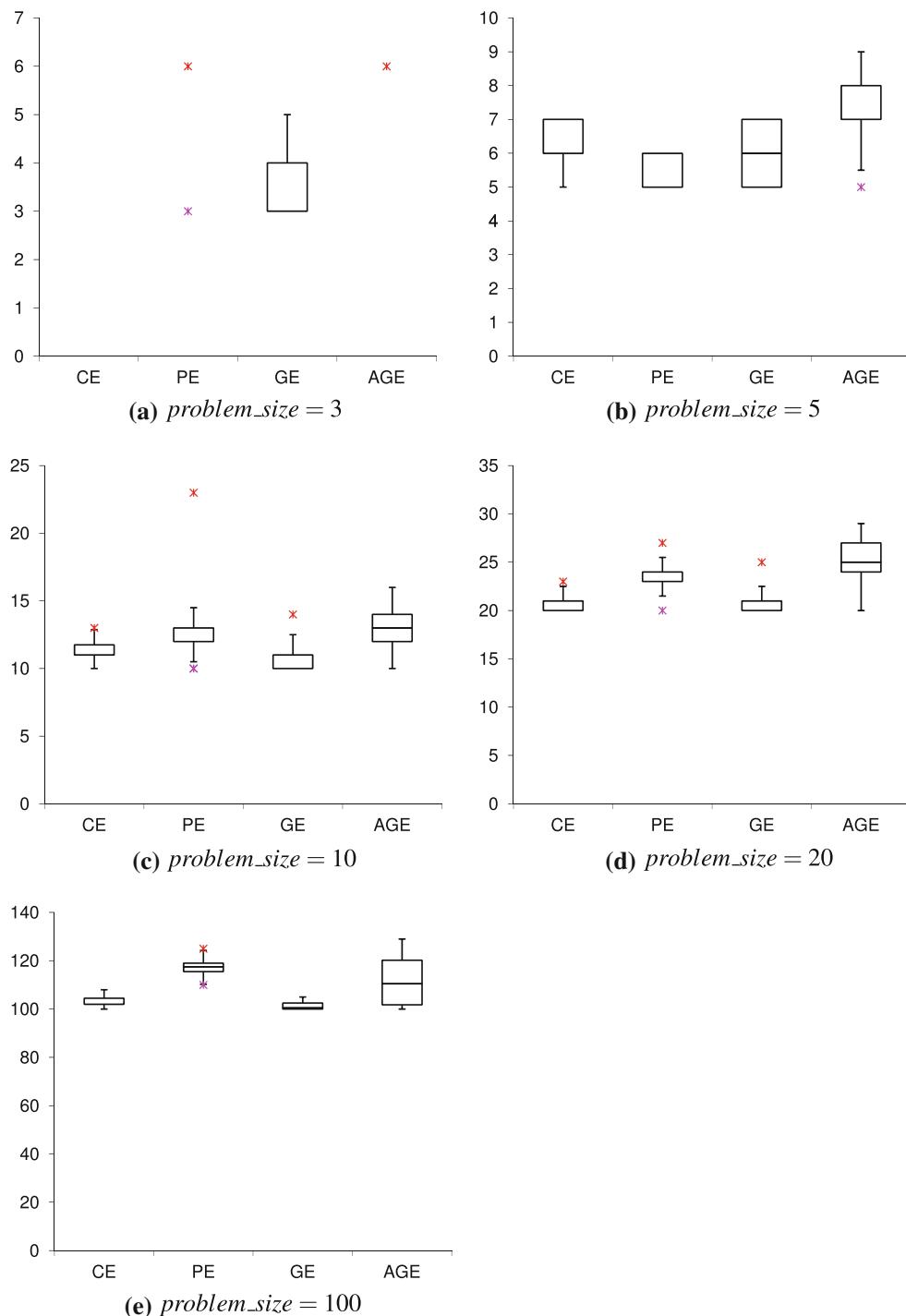


Fig. 7 Box plots of number of lamps used for each *problem_size*

problem_size. To exploit a further measurement of comparison, the first occurrence of an acceptable solution also appears in the results: here an *acceptable solution* is defined as a global solution with at least 80 % of the final fitness value obtained by the CE. Table 1 summarises the results for significant values of *problem_size*. For each evolution-

ary algorithm are reported the results reached at the end of the evolution: average fitness value, average enlightenment percentage of the room, average number of lamps, and average number of lamps evaluated before finding an acceptable solution (along with the standard deviation for each value).

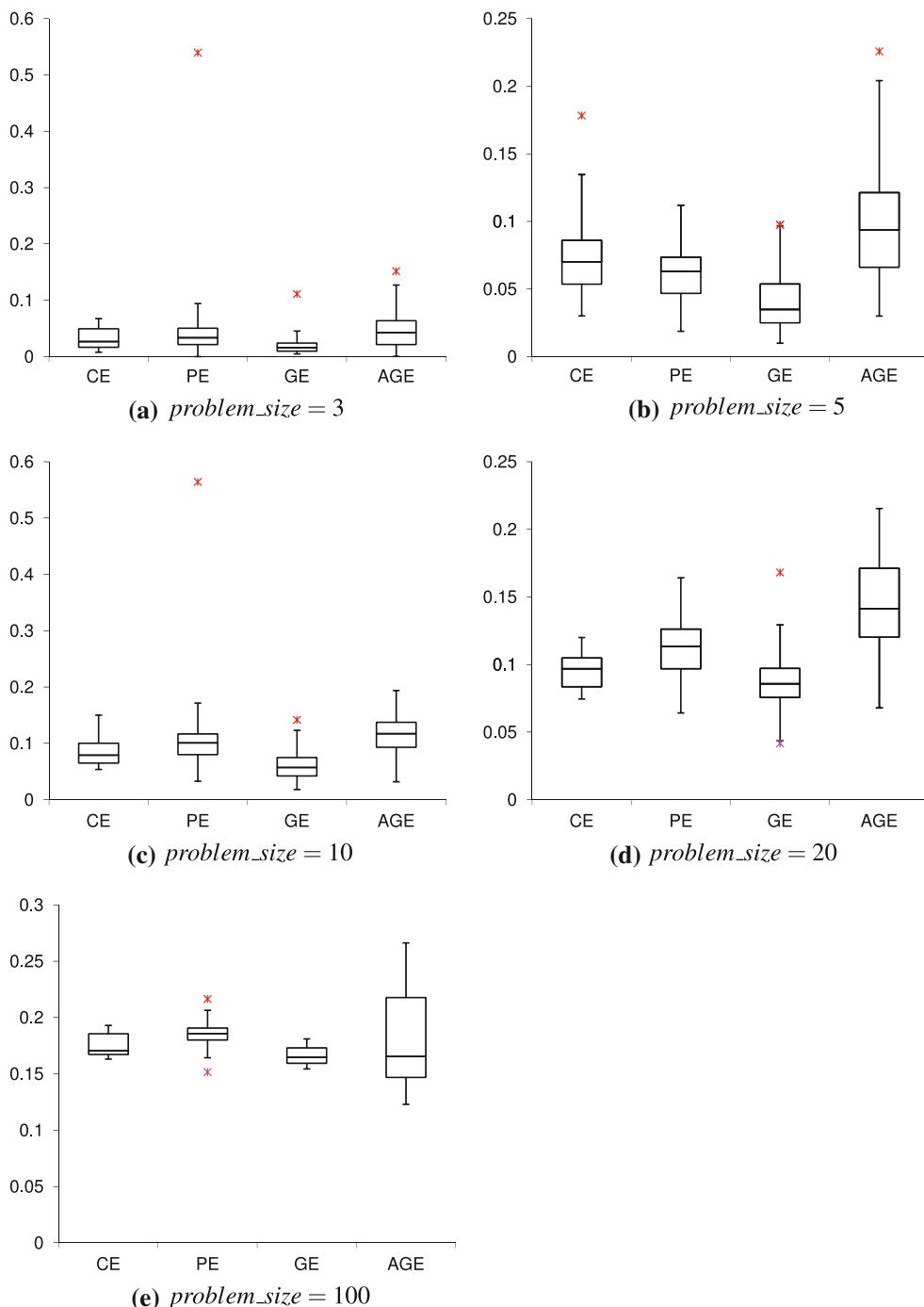


Fig. 8 Box plots of overlap percentage for each *problem_size*

Results distributions are compared with a standard Student's *t* test, with no assumptions on the homogeneity of the standard deviations. Entries in bold in Table 1 are significantly better ($p < 0.05$) than others in the same column, for the same *problem_size*.

For each *problem_size*, a box plot is provided in Fig. 5. It is noticeable how the performance of each evolutionary algorithm is very close for small values of *problem_size*, while

PE, GE and AGE gain the upper hand when the complexity increases.

In particular, PE obtains the best performance from *problem_size* = 10 onwards. The extra information inserted allows the approach to obtain high enlightenment percentages even when the complexity of the task increases, as shown in Fig. 6.

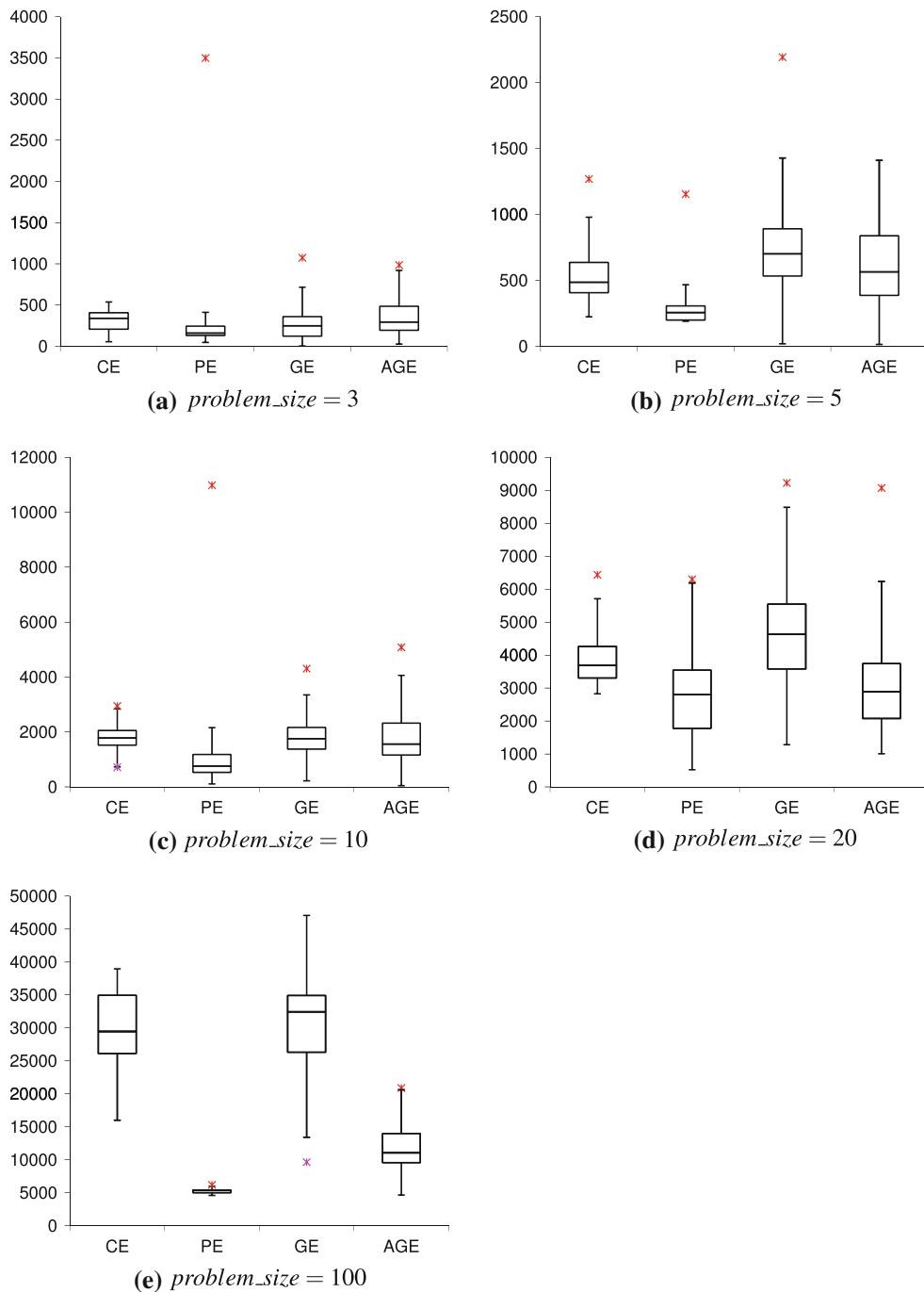


Fig. 9 Box plots of number of lamps evaluated before reaching an acceptable solution, for each *problem_size*

On the other hand, GE obtains enlightenment percentages close to CE, but on the average it uses a lower number of lamps, that leads to a lower overlap, as it is noticeable in Figs. 7 and 8.

When dealing with the number of lamp evaluations needed before reaching what is defined an *acceptable* solution, PE is again in the lead, see Fig. 9.

In Fig. 10, a profile of the best run for *problem_size* = 100 for each algorithm is reported. PE enjoys a rapid growth in the first stages of the evolution, thanks to the extra information it can make use of, while GE proves more efficient than CE in the last part of the evolution, where exploitation becomes more prevalent. AGE shows several steep improvements, in correspondence of

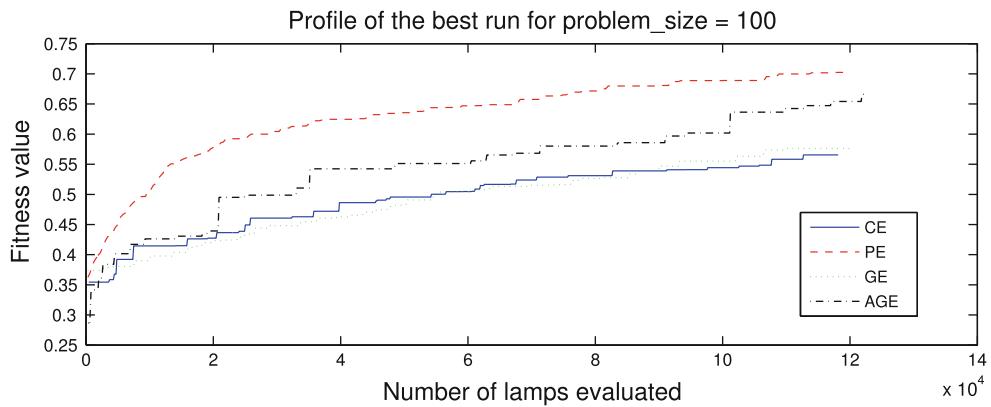


Fig. 10 Profile of the best run for each evolutionary algorithm at *problem_size* = 100

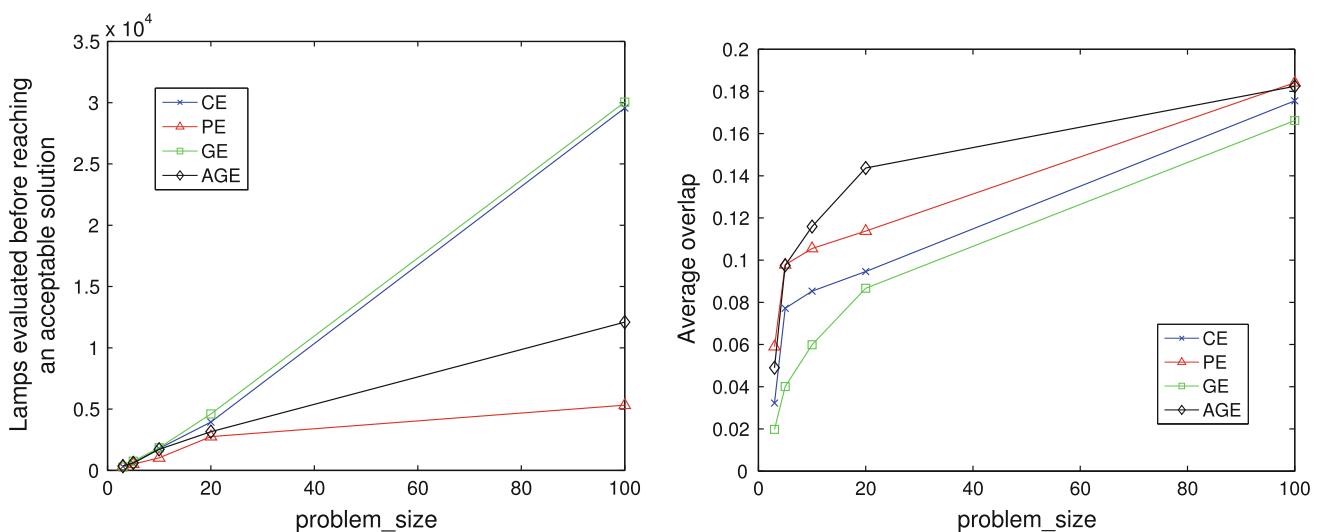


Fig. 11 Left average number of lamps evaluated before reaching an acceptable solution, for different values of *problem_size*. Right average overlap of the final solution, for different values of *problem_size*

a particularly successful activation of the heuristic group operators.

As it is noticeable in Fig. 11 (left), while the number of lamps evaluated before reaching an acceptable solution grows more than linearly for CE and GE, AGE shows a less steep growth and PE outperforms all the other algorithms. On the other hand, GE presents the lowest overlap for all values of *problem_size*, see Fig. 11 (right).

8 Conclusion and future work

The lamps benchmark has the major advantage to provide a set of toy problems that are simple, and for which the complexity can be characterised with a single real value (the surface ratio between room and lamp sizes). This formulation

is very convenient to get some insight on the behaviour of algorithms with respect to the size of the problem.

The intuition that guided the development of Parisian Evolution, Group Evolution and Allopatric Group Evolution has been confronted to experimental analysis, to yield the following conclusions: Parisian Evolution is the most efficient approach in terms of computational expense, and scalability; Group Evolution yields better and more precise results, in a computational time that is similar to Classical Evolution; Allopatric Group Evolution shows a behavior that is intermediate between the two, obtaining some advantage over GE for higher complexities of the problem, but failing to meet Parisian Evolution's performance. In general, AGE solutions also present a considerable standard deviation, probably because the combination of heuristics and evolutionary operators drives the evolution in very different niches of the search space, depending on the single run.

These differences can be explained by the nature of a priori information that has been injected into the algorithms. Parisian Evolution relies on a deterministic algorithm for selecting the group of lamps that are used as the current global solution at each generation, while Group Evolution does not make any assumption on it and let evolution decide which lamps can be grouped to build various solutions. Allopatric Group Evolution is a first attempt to make use of expert knowledge while avoiding a strong algorithmic choice, by adding information at the level of its heuristic genetic operators: the chosen metrics, however, show a variable performance, dependent on the size of the problem.

In some sense Parisian Evolution is making a strong algorithmic choice for the grouping stage, that acts as a constraint on the evolution of the population of lamps. It has the major advantage to reduce the complexity of the problem by providing solutions from the evolution of simpler structures (lamps instead of groups of lamps for both Allopatric and standard Group Evolution, or for Classical Evolution). It may be considered as a “quick and dirty” approach.

Future work on this topic will investigate other possible compromises between heuristics and evolution: a promising option is an hybridisations between Parisian and Group Evolution, i.e. running a population of elements in the Parisian mode to rapidly get a good rough solution, and using the Group scheme to refine it. A more realistic modeling of the lamp’s light could also be used, taking into account the gradual fading from the source; this approach would shift the problem from enlightening the room to ensure that each point of it receives at least a certain amount of light.

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