Real-Valued Evolutionary Multi-Modal Optimization driven by Hill-Valley Clustering

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ABSTRACT

Model-based evolutionary algorithms (EAs) adapt an underlying search model to features of the problem at hand, such as the linkage between problem variables. The performance of EAs often deteriorates as multiple modes in the fitness landscape are modelled with a unimodal search model. The number of modes is however often unknown a priori, especially in a black-box setting, which complicates adaptation of the search model. In this work, we focus on models that can adapt to the multi-modality of the fitness landscape. Specifically, we introduce Hill-Valley Clustering, a remarkably simple approach to adaptively cluster the search space in niches, such that a single mode resides in each niche. In each of the located niches, a core search algorithm is initialized to optimize that niche. Combined with an EA and a restart scheme, the resulting Hill-Valley EA (HillVallEA) is compared to current state-of-the-art niching methods on a standard benchmark suite for multi-modal optimization. Numerical results in terms of the detected number of global optima show that, in spite of its simplicity, HillVallEA is competitive within the limited budget of the benchmark suite, and shows superior performance in the long run.

CCS CONCEPTS

• Mathematics of computing \rightarrow Evolutionary algorithms; Continuous optimization;

KEYWORDS

continuous optimization, black box, niching, clustering, multi-modal optimization, evolutionary algorithm, model based

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1 INTRODUCTION

Model-based evolutionary algorithms (EAs) adapt an underlying search model based on certain features of the fitness landscape. Classically, these features are related to linkage, or dependence, of problem variables. In this work, we focus on adaptive methods to exploit multi-modality of the fitness landscape.

In this work, In optimization, a *niche* is a subset of the search space where only one mode resides. When the fitness landscape is multi-modal, the performance of many EAs deteriorates as high-quality solutions can be found in different parts of the search space, which may prevent narrowing down the search to a specific region. Being able to explicitly deal with multi-modality may therefore be beneficial to many EAs. In addition, exploring multiple niches can provide additional insight into the structure of the problem at hand. Real-world problems are often not unimodal, and by providing the decision maker with multiple high-quality solutions, the final solution can be chosen based on external factors that are best considered once the set of interesting alternatives is known.

Niching methods originated as a tool for improving population diversity in EAs [10], but are now generally designed for multi-modal optimization (MMO). MMO aims at locating multiple high-quality locally optimal solutions in a single run. One of the difficulties with niching methods is that they often introduce additional *niching parameters* such as a minimal niche size or the number of niches [10]. The number of niches is however generally unknown a priori, and niching methods should therefore make few assumptions on their size, shape, or number. In this work, we therefore introduce a niching method based on a remarkably simple clustering method that is adaptive to the fitness landscape and investigate to what extent it can improve the multi-modal optimization power of existing well-known EAs.

The remainder of the paper is organized as follows. In Section 2, we discuss different niching concepts of related methods. In Section 3, we introduce the clustering method Hill-Valley Clustering. In Section 4, we describe how we combined clustering with a core search algorithm and a restart scheme into the Hill-Valley Evolutionary Algorithm (HillVallEA). In Section 5, the performance of HillVallEA is numerically evaluated on the CEC2013 niching benchmark suite [9], that was also used for the GECCO'17 niching

competition. Performance is compared to state-of-the-art niching methods. We finally summarize and discuss the results in Section 6 and conclude in Section 7.

2 RELATED WORK

Two-stage niching methods are a class of niching methods where the first stage is aimed at locating different niches in the search space. In the second stage, core search algorithms are initialized in the niches. The Nearest-better EA (NEA2+) [12, 13] is a realvalued two-phase MMO EA, which uses nearest better clustering (NBC) in the first phase to cluster the initial population. In NBC, a nearest better tree is constructed, i.e., a spanning tree that connects each solution to its nearest solution that has better fitness. By the definition of a local optimum, all nearby solutions are of worse fitness, and the outgoing edge of a local optimum is therefore expected to be longer than average. By removing long edges from the spanning tree, niches can be detected, and a core search algorithm then performs optimization in each niche. To determine edges that are longer than expected, two cutting rules were tuned to a set of benchmark problems. When sampling few solutions with respect to the size of the search space, which occurs especially in high-dimensional problems, the distance between neighbouring solutions is heavily subject to randomness, and distinguishing long edges becomes hard. To reduce the dependency of the spread of the initial sample on the performance of the algorithm, the initial samples can be spread out more evenly over the search space compared to the commonly used uniform sampling, by means of rejection sampling [15]. This is used in NEA2+ for a sharper tuning of the cutting rules, improving its performance.

In [11], an EA is introduced where the same nearest better tree is constructed as in NBC, but instead of cutting rules, the correlation between the search space and fitness values is exploited to determine the edges that need to be cut. This correlation is computed based on a set of fitted Gaussian mixture models, but the performance of this approach deteriorates in situations where Gaussian mixture models are ill-fitted.

Winner of the GECCO'17 niching competition is the repellingsubpopulations (RS-CMSA) algorithm [1, 2]. In RS-CMSA, instances of the core search algorithm CMSA are randomly initialized and maintain a minimum distance to each other, using rejection sampling. When all core search algorithms are terminated, the method is restarted, and the core search algorithms are furthermore kept away from previously located global optima. In this way, easy optima are found first, and the search is later pushed to unexplored regions of the search space. Rejection regions from located optima, referred to as taboo regions, are considered to be (hyper-)spheres around the located optimum. The radius of the taboo region grows adaptively if the corresponding optimum is located multiple times, and shrinks again if rejection sampling fails too often. These adaptation parameters and the number of simultaneously run core search algorithms are then validated on the same set of benchmark problems used in this work.

One of the performance measures in the niching competition is *precision*, meant as the fraction of relevant detected solutions compared to the overall number of detected solutions. In order to filter out the local optima, RS-CMSA applies a post-processing

step based on the *Hill-Valley* test [14]. The Hill-Valley test can be used to detect whether two solutions belong to the same niche, see Algorithm 1. To achieve this, an edge is drawn between the two solutions in the search space, and the fitness is evaluated on N_t equidistantly located test points along the edge. If the fitness on any of the test points is worse than the fitness of both solutions, there is a *hill* between the two solutions, and the two solutions presumably belong to different *valleys* (niches).

```
Algorithm 1: Hill-Valley test [14]function: [B] =Hill-Valley(\mathbf{x}_{left}, \mathbf{x}_{right}, N_t)input :Solutions \mathbf{x}_{left}, \mathbf{x}_{right}, Number of test points N_t, Fitness function f (to be minimized)output :\mathbf{x}_{left} and \mathbf{x}_{right} belong to the same niche? (boolean)for k = 1, \ldots, N_t do\mathbf{x}_{test} = \mathbf{x}_{right} + \frac{k}{N_t+1}(\mathbf{x}_{left} - \mathbf{x}_{right});if \max\{f(\mathbf{x}_{right}), f(\mathbf{x}_{left})\} < f(\mathbf{x}_{test}) then| return false;endendreturn true;
```

The Hill-Valley test was applied in the multi-national EA [14] to each newly-sampled solution to determine if it belongs to an existing population, if two populations need to be merged, or if a new population needs to be formed. This approach was tested on noise-free two-dimensional problems while using five test points ($N_t = 5$). In RS-CMSA, the equidistant test points are replaced by a golden section search, with a maximum of $N_t = 10$ test points. More test points could be used due to its limited application only as post-processing step to filter out indistinct global optima [2].

3 HILL-VALLEY CLUSTERING

In this work, we define multi-modal optimization (MMO) as the search for only global optima. In the presence of noise, or in real-world applications, one might however also be interested in locating high-quality local optima. Guided by the problems in the CEC2013 niching benchmark suite, the niching method we present in this work discards these local optima, but this can be easily adapted.

Given a real-valued search space $X \subseteq \mathbb{R}^d$ and objective function $f: X \to \mathbb{R}$, that, without loss of generality, needs to be minimized, the aim of MMO is to find all $\mathbf{x}^* \in X$ that obtain the minimum possible objective value, i.e.,

$$f(\mathbf{x}^*) \le f(\mathbf{x}), \quad \forall \mathbf{x} \in X \subseteq \mathbb{R}^d.$$
 (1)

The core of Hill-Valley Clustering is the Hill-Valley test, combined with the concept of the nearest better tree. The idea is that solutions belonging to the same niche should be in the same cluster.

Hill-Valley Clustering is initialized with a population \mathcal{P} of $|\mathcal{P}| = N$ solutions, and these solutions are sorted such that the best solution is first. This best solution \mathbf{x}_0 forms the first cluster $C_0 = \{\mathbf{x}_0\}$. Then, we consider the second-best solution \mathbf{x}_1 , and the Hill-Valley test in Algorithm 1 is used to test whether it belongs to the same

niche as \mathbf{x}_0 . When it does, it is added to the cluster of \mathbf{x}_0 , denoted by $C_{(\mathbf{x}_0)} = C_{(\mathbf{x}_0)} \cup {\mathbf{x}_1}$. Otherwise, as there are no other solutions with higher fitness to check against, a new cluster is formed, i.e., $C_1 = {\mathbf{x}_1}$.

Next, the third-best solution \mathbf{x}_2 is tested against the nearest solution that has better fitness, which can either be \mathbf{x}_0 or \mathbf{x}_1 , depending on which one is nearer. If \mathbf{x}_2 does not belong to the same niche as its nearest better solution, we test it against the second-nearest better solution. If it also does not belong to that niche, we create a new cluster from \mathbf{x}_2 . Multiple neighbours are checked as the nearest better solution could belong to a different niche, but a second-nearest better solution, which is located slightly further, but in a different direction, might belong to the same niche. For each solution, we check its $\gamma = d+1$ nearest better neighbours, where d is the problem dimensionality. In this way, even for the one-dimensional problems, at least two neighbours are considered.

If a solution does not belong to any of the niches of its nearest neighbour, it forms a new cluster. This procedure is repeated for all solutions in the selection.

The output is a set of clusters, each consisting of at least one solution. With these clusters, core search algorithms are initialized.

To slightly reduce the number of function evaluations, we only check nearest better solutions from each cluster once. If two nearest better solutions belong to the same cluster, we only check the first, and if it is rejected, we reject the second without checking.

3.1 Number of test points N_t

We base the number of test points N_t on the (Euclidean) length of the edge connecting the two solutions, and distribute the points equidistantly over the edge. We compute the expected edge length (EEL) as the distance between two solutions when they would have been scattered equidistantly in the search space, by,

$$EEL = \sqrt[d]{V_X/|\mathcal{S}|},\tag{2}$$

where V_X is the volume of the search space X. The longer an edge is, compared to the expected edge length, the more test points are sampled, according to,

$$N_t = N_t(\mathbf{x}_{\text{left}}, \mathbf{x}_{\text{right}}) = 1 + \left\lfloor \frac{\|\mathbf{x}_{\text{left}} - \mathbf{x}_{\text{right}}\|}{\text{EEL}} \right\rfloor, \tag{3}$$

where the fraction is the current Euclidean edge length divided by the expected edge length. Note that N_t is the maximum number of test points that is evaluated for a single call of the Hill-Valley test. When two solutions belong to the same niche, all points will be evaluated. If not, the Hill-Valley test rejects the edge before all N_t points are evaluated.

Alternatively, if the volume of the search space cannot be easily determined, the expected edge length can be replaced by the average edge length. For each solution, find its nearest better solution and the corresponding edge length. Then, compute the average edge length over all edge lengths. If the nearest better distances are stored, the computational overhead is a negligible O(|S|), compared to a complexity of $O(d|S|^2)$ of Algorithm 2.

```
Algorithm 2: Hill-Valley Clustering
 function: [K] = HillValleyClustering(S)
              :Set of solutions S,
 input
                search space volume V_X,
                problem dimension d
 output :Set of clusters \mathcal{K} = \{C_0, C_1, \dots, C_{K-1}\}
 Sort solutions x_i \in S on fitness value, fittest first;
 for i = 1, ..., |S| - 1 do
      for j = 0, ..., i - 1 do
       \delta_i \leftarrow \|\mathbf{x}_i - \mathbf{x}_i\|; // Euclidean distances to better \mathbf{x}_i
      end
      for j = 0, ..., \min\{i - 1, d\} do
           k = index of the j-th nearest better solution in \{\delta_j\};
           N_t = 1 + \left| \delta_k / \sqrt[d]{V_X/|\mathcal{S}|} \right|;
           if not already checked cluster C_{(\mathbf{x}_k)} then
                if Hill-Valley(\mathbf{x}_k, \mathbf{x}_i, N_t) then
                     C_{(\mathbf{x}_k)} = C_{(\mathbf{x}_k)} \cup \{\mathbf{x}_i\}; // add to cluster of \mathbf{x}_k
                end
           end
      end
      if x_i was not added to any cluster then
           C := \{\mathbf{x}_i\};
                                          // new cluster containing \mathbf{x}_i
           \mathcal{K} = \mathcal{K} \cup C:
                                 // add new cluster to the cluster set
      end
```

4 HILL-VALLEY EVOLUTIONARY ALGORITHM

end

return K;

After clustering the initial population using Hill-Valley Clustering described in Algorithm 2, for each cluster, a core search algorithm is initialized. The niching methods discussed in this work all used different core search algorithms, which disguises the added value of a specific niching approach. We therefore equip HillVallEA with different core search algorithms.

4.1 Core search algorithms

We test HillVallEA with the core search algorithms $\mathcal A$ listed in Table 1, which are CMSA [4] and different versions of AMaLGaM [5]. These algorithms are based on a Gaussian distribution, initialized by a mean μ and covariance matrix Σ , population size N_c , and return a single solution, i.e.,

$$[\mathbf{x}] = \mathcal{A}(\mu, \Sigma, N_c) \tag{4}$$

// return cluster set

AMaLGaM is an estimation of distribution algorithm, where a Gaussian is fitted to selected solutions with maximum likelihood. CMSA was used instead of the more common CMA-ES [7], as it was suggested to perform better when adding elitism [2]. CMSA fits a Gaussian using the population mean of the previous generation.

The methods have different recommended population sizes, see Table 1. Likely, a smaller core population size is better, since the

Table 1: Different core search algorithms $\mathcal A$ with corresponding recommended cluster (population) size $N_c^{\rm rec}$ as taken from literature, for problem dimensionality d.

Core search algorithm \mathcal{A}	abbr.	Nrec
CMSA [4]	CMSA	$\frac{3 \log d}{}$
AMaLGaM [5]	AM	$17 + 3 \cdot d\sqrt{d}$
AMaLGaM-Univariate [5]	AMu	$10\sqrt{d}$
iAMaLGaM [5]	iAM	$10\sqrt{d}$
iAMaLGaM-Univariate[5]	iAMu	$4\sqrt{d}$

complexity of the problem within each niche is expected to be smaller than the complexity of the full problem. Therefore, a small population might be sufficient to converge to the optimum within a niche, using fewer function evaluations. This would allow for generations of the core search algorithms to be run, which is beneficial especially when many clusters are detected.

Core search algorithms are initialized from a cluster by setting the μ as the cluster mean, and Σ as the sample covariance matrix of the solutions in the cluster with respect to μ . If the cluster consists of fewer than d+1 solutions, only the diagonal of Σ is estimated and all other entries are set to be zero. If a cluster consists of only one solution, the covariance matrix is initialized by the identity matrix, multiplied by $0.01 \cdot \text{EEL}$, so that newly sampled solutions will be closer-by than its nearest better solutions in the initial population.

CMSA is terminated using the recommended criteria [7], and parameters set as in RS-CMSA [2], that is, if the improvement in fitness value over the last $10 + \lfloor 30d/N_c \rfloor$ generations is less than $TOL = 10^{-5}$. Tolerance TOL corresponds to the desired accuracy of optima in the CEC2013 niching benchmark suite, and should be adapted if a different accuracy is required for the problem at hand. Furthermore, fail-safe termination criteria are added to terminate the core search algorithms if the standard deviation of solutions in the search space reaches machine accuracy (10^{-15}) and if the condition number of the covariance matrix is larger than 10^{14} .

Termination is enforced for AMaLGaM-variant core search algorithms if the maximum standard deviation of solutions in the search space is too low (10^{-12}), or if the standard deviation of fitness values is too low (10^{-12}).

Finally, the whole optimization is terminated if the budget, in terms of a maximum number of function evaluations specified in the benchmark, is reached.

4.2 Restarts with an elitist archive

There are two population size parameters that need to be set. First, there is the initial population size N used as input for Hill-Valley Clustering. The second parameter is the population size of the core search algorithms N_c . Schemes that grow the population of an EA over time, i.e., with restarts, are common to overcome setting these parameters, such as used in I-POP-CMA [3], the parameter-free genetic algorithm [8], and the interleaved multistart scheme [6].

HillVallEA is initialized with a population \mathcal{P} of size N=16d, where d is the problem dimensionality. To enhance performance, truncation selection is performed with a selection fraction τ based on the core search algorithm used. This is $\tau=0.35$ for the versions

```
Algorithm 3: Hill-Valley Evolutionary Algorithm
 function: [\mathcal{E}] = HillVallEA(\mathcal{A})
 input
              :Core search algorithm \mathcal{A},
               problem dimension d
 output :Set of presumed global optima \mathcal{E}
 N = 16d;
                                              // initial population size
 N_c = N_c^{\rm rec}(\mathcal{A});
                                                            // cluster size
 \tau = \tau(\mathcal{A});
                             // 	au=0.5 for \mathcal{A}= CMSA or 	au=0.35 else
 \mathcal{E} = \{\};
                                                        // elitist archive
 while budget remaining do
      // First phase - locating niches
      \mathcal{P} = \text{uniformly sample}(N); // also evaluates solutions
      \mathcal{P} = \mathcal{P} \cup \mathcal{E};
                                             // add elites to population
      S = truncation\_selection(\mathcal{P}, \tau);
      \mathcal{K} = \text{HillValleyClustering}(\mathcal{S});
      // Second phase - niche optimization
      foreach C_i \in \mathcal{K} do
           if best solution in C is an elite then
                // skip already-optimized niches
                continue
           end
           // Run core search algorithm
           \mathbf{x}_i = \mathcal{A}(\mu(C_i), \Sigma(C_i), N_c)
           if x_i is a presumed distinct global optimum then
                if for any e \in \mathcal{E}, x_i + TOL < e then
                     // Empty the archive
                    \mathcal{E} = \{\}
                end
                // Add to elitist archive
                \mathcal{E} = \mathcal{E} \cup \mathbf{x}_i
           end
      end
      if no new solution was added to \mathcal{E} then
           N \leftarrow 2N;
                                   // increase initial population size
           N_c \leftarrow 1.2N_c;
                                                 // increase cluster size
      end
 end
```

of AMaLGaM and $\tau=0.5$ for CMSA. By performing selection, less effort is spent on low-fitness regions in the search space. On the other hand, small niches could be accidentally discarded when selection pressure is too high.

The selection S of $\lfloor \tau N \rfloor$ solutions is then clustered using Hill-Valley Clustering, of which pseudo code is given in Algorithm 2. From each of the resulting clusters, a core search algorithm is ran one after another. By construction of Hill-Valley Clustering, the resulting clusters are sorted on the fitness values of the best, and the cluster containing the overall best solution is ran first. The run order is of importance when more clusters are found than the computational budget allows to run until convergence.

After running all core search algorithms with a cluster size of $N_c = N_c^{\rm rec}$, the result is a set of $|\mathcal{K}|$ candidate optima. A post-processing step is performed to discard the local optima and the

duplicate global optima. First, all solutions that are more than TOL worse than the all-time best, are discarded. Then, all remaining optima are tested for being in a different niche by the Hill-Valley test. The expected edge length does not make sense now, so we use the originally proposed $N_t=5$ test points, similar to the post-processing step in [2]. All presumed distinct global optima are then added to the elitist archive \mathcal{E} .

If, after all core search algorithms are terminated, there is budget remaining, we restart the procedure. To prevent overhead, we add all elite solutions (the presumed distinct global optima) to the new population. If, after clustering, the cluster-best is any of these elites, we do not re-run that cluster, as that niche was presumably already optimized.

If no new global optima are found in a run, restarts are performed with increased population size to be able to detect smaller niches. Furthermore, the population size of the core search algorithms is also increased in order to find optima in more complex niches.

The initial population size is increased with a factor $N^{\rm inc}=2$ after each run in which no new global optima are detected. The core search algorithm starts with the recommended value in Table 1, and is incremented by a factor $N_c^{\rm inc}=1.2$. Pseudo code of HillVallEA is given in Algorithm 3.

5 EXPERIMENTS

We evaluate the performance of HillVallEA on the test problems in the CEC2013 niching benchmark suite [9]. The benchmark consists of 20 problems, described in Table 2, to be solved within the corresponding benchmark budget in terms of a limited number of function evaluations.

For each of the benchmark problems, the location of the optima and the corresponding fitness values are known. As performance measure, the peak ratio is used, which measures the fraction of global optima detected, computed according to the guidelines of the niching benchmark suite [9]. In contrast to the full benchmark suite, where a range of accuracies ε is used, and the final peak ratio is the mean of these results, we only use the highest accuracy of $\varepsilon=10^{-5}$. Due to the post-processing step in HillVallEA, all local optima and duplicate global optima are filtered out, making the results independent of the choice of $\varepsilon \geq 10^{-5}$.

All benchmark functions are defined on a bounded domain, making computation of the search space volume in Equation (2) straightforward.

All experiments in this work are repeated 50 times, and resulting performance measures are averaged over all repetitions. Results are tested for statistical significance with a two-sided 2-sample Wilcoxon rank sum test ($\alpha=0.05$). A non-parametric test is used because the peak ratio is discrete, making it non-normally distributed.

For comparison, RS-CMSA was re-implemented in C++ based on the on-line available MATLAB implementation. Performance, measured in the peak ratio, was comparable to the results published in [1].

In Section 5.1 we compare performance of different versions of the restart scheme. In Sections 5.2 and 5.3, an attempt was made to further improve HillVallEA by respectively tuning hyperparameters and by incorporating an advanced sampling scheme.

Table 2: Niching benchmark suite from the CEC2013 special session on multi-modal optimization [9]. For each problem the function name, problem dimensionality d, number of global optima #gopt, and local optima #lopt and budget in terms of function evaluations are given.

#	Function name	d	#gopt	#lopt	budget
1	Five-Uneven-Peak Trap	1	2	3	50K
2	Equal Maxima	1	5	0	50K
3	Uneven Decreasing Maxima	1	1	4	50K
4	Himmelblau	2	4	0	50K
5	Six-Hump Camel Back	2	2	5	50K
6	Shubert	2	18	many	200K
7	Vincent	2	36	0	200K
8	Shubert	3	81	many	400K
9	Vincent	3	216	0	400K
10	Modified Rastrigin	2	12	0	200K
11	Composition Function 1	2	6	many	200K
12	Composition Function 2	2	8	many	200K
13	Composition Function 3	2	6	many	200K
14	Composition Function 3	3	6	many	400K
15	Composition Function 4	3	8	many	400K
16	Composition Function 3	5	6	many	400K
17	Composition Function 4	5	8	many	400K
18	Composition Function 3	10	6	many	400K
19	Composition Function 4	10	8	many	400K
20	Composition Function 4	20	8	many	400K

Then, in Section 5.4, the performance of HillVallEA with multiple core search algorithms is compared. Finally, in Section 5.5, performance is investigated on the benchmark suite with extended computational budget.

5.1 Restart efficiency

In an attempt to reduce the computational overhead of the restart scheme, the located global optima are added to the population, as a representative for that entire niche. After clustering, these global niches are then skipped in the resulting local phase. Alternatively, both global and local optima could be added, which would prevent local niches from being re-explored as well. We compare these to the use of a restart scheme without addition of optima to the population for HillVallEA and the simplest core search algorithm, AMaLGaM-Univariate and run it on all of the benchmark problems limited to the benchmark budget. Achieved peak ratios are given in Table 3.

Adding all optima to the population deteriorates performance significantly, especially notable for problems 6, 8, 13–20, all functions with many local optima. By adding all local optima back to the population, the population grows faster, and more of the budget is required for clustering these local optima over and over after each restart. Problems 9 has no local optima, so there is no difference between adding all located optima or only the located global optima. Compared to not adding optima, a huge improvement in peak ratio from 0.503 to 0.875 can be observed.

Table 3: Peak ratios of HillVallEA-AMu with three forms of the restart scheme. Average (avg.) peak ratio is computed over all benchmark problems. The best achieved results per problem, and those results not statistically different from it, are typeset in boldface. The right three columns show the fraction of budget (in function evaluations) used by the 'only global' restart scheme of HillVallEA-AMU for initialization (init), Hill-Valley Clustering (HVC) and local optimization (lopt).

	Peak ra	Peak ratio of Restart Schemes			Budget usage		
#	all optima	only global	no optima	init	HVC	lopt	
1	1.000	1.000	1.000	0.04	0.05	0.91	
2	1.000	1.000	1.000	0.07	0.11	0.82	
3	1.000	1.000	1.000	0.04	0.06	0.90	
4	1.000	1.000	1.000	0.08	0.12	0.80	
5	1.000	1.000	1.000	0.03	0.04	0.93	
6	0.942	0.997	0.999	0.01	0.02	0.97	
7	1.000	1.000	0.969	0.23	0.36	0.41	
8	0.263	0.656	0.776	0.01	0.02	0.97	
9	0.875	0.875	0.503	0.10	0.26	0.64	
10	1.000	1.000	1.000	0.03	0.07	0.90	
11	1.000	1.000	1.000	0.04	0.07	0.89	
12	1.000	1.000	1.000	0.01	0.02	0.97	
13	0.857	0.973	0.953	0.02	0.03	0.95	
14	0.713	0.783	0.760	0.01	0.01	0.98	
15	0.723	0.750	0.750	0.01	0.01	0.98	
16	0.670	0.673	0.670	0.01	0.01	0.98	
17	0.703	0.745	0.743	0.01	0.01	0.98	
18	0.653	0.657	0.643	0.00	0.01	0.99	
19	0.512	0.512	0.490	0.01	0.01	0.98	
20	0.313	0.318	0.315	0.01	0.01	0.98	
avg.	0.811	0.847	0.829	0.04	0.07	0.90	

5.2 Parameter tuning of the restart scheme

The restart scheme has four parameters: the initial population size N, the increment factor $N_c^{\rm inc}$, the initial cluster size N_c and the increment factor $N_c^{\rm inc}$. These parameters have been tuned manually to intuitively make sense, and are found to perform well. To demonstrate robustness of the scheme, we tuned these parameters further in an automatic fashion. We set $N=2^{\xi_0}\cdot d$, $N_c^{\rm inc}=\xi_1, N_C=\xi_2\cdot N_C^{\rm rec}$, $N_c^{\rm inc}=\xi_3$, with $\xi_0\in[4,10]$ and $\xi_i\in[1,5]$ for i=1,2,3. The manually chosen parameters as stated in Algorithm 3 can then be written as $\xi=(4,2,1,1.2)$.

AMaLGaM-Univariate was used to maximize the performance of HillVallEA-AMu, measured by the average of achieved peak ratio over 50 repetitions of the 20 benchmark problems. The obtained average peak ratios are shown in Figure 1. The best obtained average peak ratio is 0.850, attained for the values $\xi = (7.872, 3.839, 1.365, 1.351)$. This is only a slight improvement over the manually tuned parameters (0.847), but with very different values for the parameters.

5.3 Advanced initial population sampling

Both NEA2+ and RLSIS sample the initial population using the MaxiMin Reconstruction (MMR) method, where, based on rejection sampling, the initial population is more evenly distributed

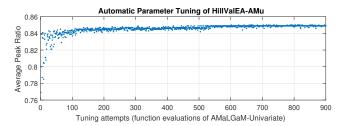


Figure 1: Automatic parameter tuning of the restart scheme of HillVallEA-AMu on the average peak ratio of 50 repetitions of the full benchmark suite.

over the search space compared to uniform sampling or Latin hypercube sampling [15]. We replaced $\mathcal{P}=$ uniformly_sample(N) in HillVallEA (Algorithm 3) by a sample generated with MMR. HillVallEA-AMu+MMR achieved an average peak ratio of 0.849 over the entire benchmark set. This is not statistically significantly different from the previously obtained 0.847 with uniform sampling (see Table 3). For individual problems, the differences in peak ratio were all less than 0.02, which is again not statistically significant. Standard uniform sampling is therefore preferred, as it is computationally cheaper.

5.4 Core search algorithms in HillVallEA

We equip HillVallEA with each of the core search algorithms in Table 1. The top three performing competitors in the GECCO'17 niching competition, based on the CEC2013 benchmark suite, are RS-CMSA, with an average peak ratio of 0.856, RLSIS [15] with 0.822, and NEA2+ with 0.810 over the entire benchmark set.

Obtained peak ratios for each of the problems are given in Table 4. For all core search algorithms, HillVallEA performs well, with the average peak ratio over all problems between 0.816–0.847, outperforming NEA2+ in all cases, and RLSIS for four different core search algorithms.

Problems 1–5 and 10 are easily solved within a fraction of the benchmark budget. HillVallEA also solves problems 7, 11, and 12 fully, but this requires almost the entire budget.

The largest differences can be observed for problems 8, 9, and 20. Problem 8 (Shubert function, 3D) has 81 global optima, all with a small niche. A strategy that allows many restarts will perform well here, which is the case with HillVallEA-AMu, which maintains a cluster size and requires fewer function evaluations to converge each local optimizer. Problem 9 (Vincent function, 3D) has largely varying niche sizes, but no local optima. Hill-Valley Clustering works very well here in detecting the large niches first, and the restart scheme prevents re-optimizing these niches, which significantly outperforms the peak ratio achieved by RS-CMSA. Problem 20 (CF4), has the largest dimensionality in the benchmark, d=20, and has one very large niche. RS-CMSA achieves a better success rate than any HillVallEA variant for this problem.

5.5 Maximum Achievable Peak Ratio

The benchmark budget is tight, as not all problems can be solved within the budget. We therefore increase the computational budget

Table 4: Peak ratio for RS-CMSA and HillVallEA equipped with different core search algorithms, for all benchmark problems and with budget limited to the benchmark budget. Average (avg.) peak ratio is computed over all problems. The best achieved results per problem, and those results not statistically different from it, are typeset in boldface.

	RS	HillVallEA				
#	CMSA	AM	AMu	CMSA	iAM	iAMu
1	1.000	1.000	1.000	1.000	1.000	1.000
2	1.000	1.000	1.000	1.000	1.000	1.000
3	1.000	1.000	1.000	1.000	1.000	1.000
4	1.000	1.000	1.000	0.999	1.000	1.000
5	1.000	1.000	1.000	1.000	1.000	1.000
6	1.000	0.976	0.997	0.980	0.989	0.976
7	0.996	1.000	1.000	1.000	1.000	1.000
8	0.881	0.410	0.656	0.437	0.465	0.501
9	0.721	0.782	0.875	0.882	0.851	0.922
10	1.000	1.000	1.000	1.000	1.000	1.000
11	1.000	1.000	1.000	1.000	1.000	1.000
12	1.000	1.000	1.000	1.000	1.000	1.000
13	1.000	0.923	0.973	0.973	0.957	0.963
14	0.780	0.763	0.783	0.793	0.750	0.783
15	0.748	0.740	0.750	0.728	0.740	0.735
16	0.667	0.670	0.673	0.670	0.677	0.677
17	0.688	0.720	0.745	0.670	0.720	0.720
18	0.667	0.623	0.657	0.637	0.640	0.667
19	0.505	0.465	0.512	0.512	0.490	0.505
20	0.465	0.238	0.318	0.338	0.275	0.315
avg.	0.856	0.816	0.847	0.831	0.829	0.838

to one billion (10⁹) function evaluations. Due to computational limits, experiments are repeated 20 times (instead of 50 in the other experiments). We consider the peak ratio for each individual problem in the benchmark versus the number of function evaluations. In Figure 2 the results for HillVallEA-AMu and RS-CMSA are given.

Problems 1–5 and 10 are solved easily within the original benchmark budget. On the Vincent function, (7 and 9), HillVallEA-AMu outperforms RS-CMSA, i.e., it achieves a peak ratio of 1.0 faster, while the reverse is true for the Shubert function, (6 and 8). All these problems are still fully solved by both algorithms.

Problems 13,14, 16, and 18 are all versions of Composition Function 3 (CF3), with increasing problem dimensionality, and contain two niches of Griewank, two of Weierstrass, and two of EF8F2 [9]. CF3-2D and 3D (problem 13 and 14) are fully solved for all repetitions. In 5D, RS-CMSA is unable to solve the Weierstrass niches, while HillVallEA-AMu occasionally solves them. In 10D, both algorithms achieve the same peak ratio, again being unable to solve the Weierstrass niches.

Problems 15, 17, 19 and 20 are all versions of CF4 with increasing problem dimensionality. For problem 15, the two Weierstrass niches are almost never solved. In problem 20, the Griewank function can furthermore not be solved by either of the methods.

Interestingly, in one repetition of problem 20, RS-CMSA accidentally discarded a previously detected global optima in the post-processing step. Overall, in the long run, HillVallEA-AMu outperforms RS-CMSA.

6 SUMMARY AND DISCUSSION

In this work, we introduced a principled approach of adaptively clustering the search space into promising niches based on Hill-Valley Clustering. Equipped with a restart scheme and various core search algorithms, Hill-Valley Evolutionary Algorithm (HillVallEA) is compared to state-of-the-art EAs designed for MMO.

HillVallEA performs very well on the CEC2013 niching benchmark. Combined with AMaLGaM-Univariate as the core search algorithm, performance is only slightly inferior to the best performing algorithm in the GECCO2017 niching competition, RS-CMSA. HillVallEA-AMu achieves an average peak ratio of 0.847 over all 20 problems, compared to 0.856 of RS-CMSA. The choice of core search algorithm does not heavily affect the overall performance of HillVallEA.

Using an elitist archive and including it in the population upon restarting further improves the performance of HillVallEA on problems with large niches, preventing these niches to be explored multiple times. A drawback is that budget is still spent on re-clustering these niches in the initialization phase. A way to overcome this would likely improve HillVallEA further.

By automatically tuning the restart parameters, we showed that the manually chosen parameters perform well, and performance is robust with respect to many different parameter choices.

In comparison to NEA2+, HillVallEA considers multiple neighbours in the cluster process. The cutting rules are not tuned based on the EA's performance, but each edge is individually considered using the Hill-Valley test. This improves the robustness and quality of clustering, justifying the additionally spent function evaluations. Furthermore, we showed that HillVallEA does not significantly improve by replacing the initial sampling algorithm. Standard uniform sampling is therefore preferred, as it is computationally cheaper.

Benchmark problems with a budget-limit do demonstrate the full potential of the tested algorithms. By comparing the performance of HillVallEA-AMu with RS-CMSA over time, we showed that Hill-VallEA outperforms RS-CMSA in the long run. However, even for a budget of 1 billion evaluations, neither of the methods are able to detect all global optima for all problems, which suggests limitations in the core search algorithms, in the two-phase approach, or an extremely increased problem complexity for some of the global optima as the dimensionality increases. Re-clustering after a number of generations might help to overcome this. On the other hand, re-clustering might deteriorate performance on problems where it is not necessary, so an adaptive approach may be warranted.

Other improvements to HillVallEA might consist of terminating core search algorithms early when it is clear that they are not approaching global optima, or include task-scheduling in order to run each of the core search algorithms based on their probability of finding a new global optimum.

7 CONCLUSION

In this work, we introduced the Hill-Valley Evolutionary Algorithm (HillVallEA), a two-phase niching method, driven by the principled, yet remarkably simple Hill-Valley Clustering. The resulting model-based evolutionary algorithm was found to be surprisingly competitive to state-of-the-art niching methods, and is even superior in the long run on a variety of benchmark problems. We conclude that

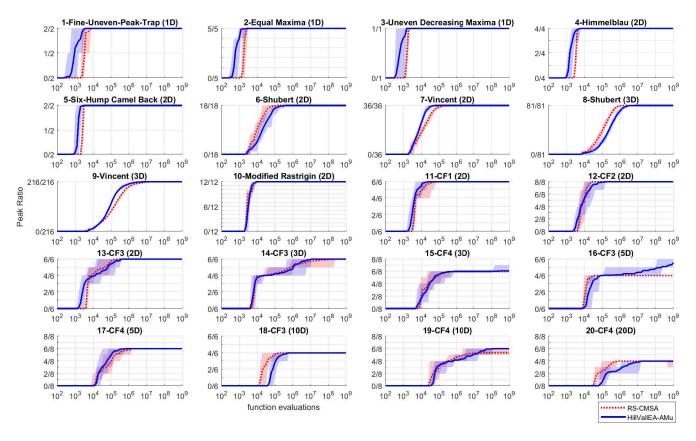


Figure 2: For each benchmark problem, the peak ratio is shown as a function of the number of function evaluations, for RS-CMSA and HillVallEA-AMu, with an increased budget of 50 times the original benchmark budget. Shaded areas show the min-max range over 20 repetitions of the experiment.

HillVallEA is an interesting new addition to the existing spectrum of evolutionary algorithms for multi-modal optimization, with high potential that may still be further exploited.

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