MinCenter: using clustering in global optimization

Abstract

A common problem arises in many scientific fields is that of locating the global minimum of a multimodal function. A novel clustering technique that tackles this problem is introduced here. The proposed method creates clusters from uniform samples of the objective function with the usage of the Kmeans clustering technique. For every cluster a center is created. Finally, a simple rejection procedure is applied to the created clusters in order to remove clusters that are close to others. The proposed method is tested on a series of well - known optimization problems from the relevant literature and the results are reported and compared against the simple Multistart global optimization method.

Keywords: Global optimization, clustering, hubrid methods, numerical methods.

1 Introduction

A novel method that estimates the global minimum of a continuous and differentiable function $f : S \to R, S \subset R^n$ is proposed in the current article. The global optimum location problem is usually defined as:

$$x^* = \arg\min_{x \in S} f(x) \tag{1}$$

where S is

$$S = [a_1, b_1] \otimes [a_2, b_2] \otimes \ldots [a_n, b_n]$$

A review of the recent advantages in the area of the Global Optimization can be found in [1]. Methods that discover the global minimum can be used in many areas such as: economics [2, 3], physics [4, 5], chemistry [6, 7], medicine [8, 9] etc. Global optimization methods usually are divided into two main categories: deterministic and random search methods. Common methods of the first category are the so called Interval methods [10, 11], where the set S is divided iteratively in subregions using some criteria. On the other hand, random search methods are used in the majority of cases, because they can be implement easy

and they do not depend on a some a priori information about the objective function. A small set of random search methods may include Controlled Random Search methods [12, 13, 14], Simulated Annealing methods [15, 16], Differential Evolution methods [17, 18], Particle Swarm Optimization methods [19, 20], Ant Colony Optimization [21, 22], Genetic algorithms [23, 24, 25] etc.

A subclass of random search methods are the clustering techniques as proposed by Rinnooy Kan [26], Ali [27], Tsoulos [28], etc. These methods are try to estimate the clusters of function in order to minimize the effort required to compute the global minimum or all the local minima of the function. The term cluster refers to a set of points that are believed, under some asymptotic considerations, to belong to the same region of attraction of the function. The region of attraction for a local minimum x^* is defined as:

$$A(x^*) = \{x : \ x \in S \subset R^n, \ L(x) = x^*\} \tag{2}$$

where $L(x)$ is a local search procedure that starts from a given point x and terminates when a local minimum is discovered. Common local search procedures are BFGS[29, 30], Steepest Descent[31], L-Bfgs [32] for large scaled functions etc.

The proposed method creates clusters iteratively using the well - known technique of the K-Means clustering introduced by MacQueen[33]. For every cluster a representative is constructed using the K-means method and afterwards a rejection procedure is utilized in order to reduce the number of representatives. Finally, for every remain point a local search procedure is started to locate the global minimum of the function.

The rest of this article is organized as follows: in section 2 the proposed method is described in detail, in section 3 some experimental test functions from the relevant literature are described and a series of test are performed on those functions and finally in section 4 some conclusions are discussed as well as some guidelines to improve the proposed method.

2 Method description

The proposed method is initially based on the commonly used global optimization method named Multistart. The proposed method creates clusters from the objective function. The multistart method is one of the simplest global optimization technique which start a local search optimizer such as BFGS from different random points and yields the lowest discovered minimum as the global one. As it was demonstrated by various researchers [34, 35], if the number of local minimum is finite then Multistart method is capable to locate the global minimum.

Due to its simplicity, the Multistart method is the base method for a series of stochastic methods in the relevant literature such as hybrid methods[36, 37], GRASP methods[38] etc. Also, it has been tested on many practical problems such as the TSP problem [39], the vehicle routing problem [40], the maximum clique problem [41], flowshop rescheduling [42], energy consumption etc. The

Algorithm 1 The main steps of the Multistart method.

1. **Initialization** step.
   (a) **Set** M as the total number of samples.
   (b) **Set** (x^*, y^*) as the global minimum. Initialize y^* to a very large value.
2. **Sampling** step.
   (a) **For** $i = 1 \ldots M$ **Do**
      i. **Sample** a point $x_i \in S$
      ii. $y_i = \mathrm{LS}(x_i)$. Where LS(x) is a local search procedure.
      iii. **If** $y_i \leq y^*$ then $x^* = x_i, y^* = y_i$
   (b) **EndFor**

base method has been extended in different works such as enhanced stopping rules for the multistart method [34, 35, 44, 45], parallel techniques [46, 47], multistart hybrid methods executed on modern GPU architectures [48, 49] etc. Also, a variety of methods have been introduced that enhance the sampling in the procedure such as the repulsion sampling [50], methods where the sampling is guided by a neural network [51] etc. Furthermore, the multistart method has been extended to solve also constrained optimization problems [52].

The main steps of a typical Multistart procedure are shown in Algorithm 1.

The proposed method replaces the sampling step of the Multistart method with the usage of centroids constructed by Kmeans clustering. The main steps of the Kmeans method are given in Algorithm 2. The estimated centroids are iteratively enhanced with Kmeans and new samples that added each time for a predefined number of times. Having created the centroids a rejection procedure is applied to reduce the number of centroids. The rejection procedure removes from the set of centers, points that have many neighbors in a predefined radius. The rejection procedure is necessary to remove from the set samples, that possible will yeld the same local optimum after the application of the local search procedure. The proposed method is described in Algorithm 3.

3 Experiments

3.1 Test functions

In order to measure the effectiveness of the proposed approach we utilize several benchmark functions from the relevant literature [53, 54]. The names and the complete forms of these equations can be obtained from [51].

Algorithm 2 The algorithm Kmeans.

1. **Repeat**
   (a) $S_j = \{\}\,,\ j = 1..K$
   (b) **For** every sample x_i **Do**
      i. **Set** $j^* = \min_{i=1}^{K} \{D(x_i, c_j)\}$, where j^* is the nearest center for sample x_i.
      ii. **Set** $S_{j^*} = S_{j^*} \cup \{x_i\}$.
   (c) **EndFor**
   (d) **For** every center c_j **Do**
      i. **Set** M_j=number of elements in S_j
      ii. **Update** c_j

$$c_j = \frac{1}{M_j} \sum_{i=1}^{M_j} x_i$$

   (e) **EndFor**
2. **Terminate** when c_j no longer change.

3.2 Experimental Results

The proposed method was tested against the traditional multistart global optimization method on the series of benchmark problems. The parameters used in the conducted experiments are listed in Table 1. The method was coded using the OpenMP library[55] in order to take advantage of multi - core modern computing systems and the experiments were conducted on a cluster of systems running the Linux operating system. The results from the experiments are listed in Tables 2 and 3. The methods were executed 30 times for each test function and averages were measured. In every run different seed for the random generator (drand48() function of C) was used. The numbers appeared in the parentheses stand for the percentage of runs where the global minimum was located sucessfully and they are not shown if the success was 100%. The last row represents the total number of function calls. The local search procedure used (denotes as LS(x)) was a BFGS variant due to Powell[30].

In Table 2 the results for the Multistart global optimization procedure are shown. The column $M = 100$ denotes the application of the algorithm given in Algorithm 1 with $M = 100$ samples. The column $M = 200$ stands for the Multistart algorithm using 200 samples. The last column stands for the results of the Multistart method with 100 samples and the application of the proposed rejection procedure of algorithm in Algorithm 4 in the samples before the application of the local search procedure. It is evident that the application of the rejection procedure does not reduce significantly the number of function

Algorithm 3 The proposed method.

1. **Initialization** step.
   (a) **Set** M as the number of samples.
   (b) **Set** (x^*, y^*) as the global minimum. Initialize y^* to a very large value.
   (c) **Set** K the number of teams, where $K < M$.
   (d) **Set** K_{MAX} the number of construction iterations for the KMeans algorithm.
   (e) **Set** $C = \{\}$, as the set of constructed centers.
2. **Construction** step.
   (a) **For** $i = 1..K_{\text{MAX}}$ **Do**
      i. **Sample** M points from the objective function $S = \{x_1, x_2, \ldots, x_M\}$
      ii. **Update** the centers C with the set S, using Kmeans.
   (b) **EndFor**
3. **Create** the set R from C using the rejection algorithm of Algorithm 4.
4. **Evaluation** step.
   (a) **For** $i = 1 \ldots |R|$ **Do**
      i. **Set** $x_i = R_i$
      ii. $y_i = \text{LS}(x_i)$. Where LS(x) is a local search procedure.
      iii. **If** $y_i \leq y^*$ then $x^* = x_i, y^* = y_i$
   (b) **EndFor**

Algorithm 4 The rejection algorithm.

1. **Set** C the set of centers.
2. **Set** $R = \emptyset$ the outcome of the rejection algorithm.
3. **Set** $D_{\min} = \min_{i \neq j} \|c_i - c_j\|$
4. **Set** $F > 1$, a double value.
5. **Set** $N_{\min} > 1$, an integer value.
6. **For** every center c_i **Do**
   (a) **Set** $N = 0$
   (b) **For** every center c_j, $i \neq j$ **Do**
      i. **If** $\|c_i - c_j\| \leq FD_{\min}$ **then** $N = N + 1$
   (c) **EndFor**
   (d) **If** $N < N_{\min}$ **then** $R = R \cup c_i$
7. **EndFor**
8. **Return** R

calls for the multistart case.

In Table 3 the experimental results for the proposed method are listed. The column $M = 100$ stands for the usage of 100 samples in the proposed method (parameter M) and the column $M = 200$ for 200 samples. The proposed method has significantly lower number of function calls than the Multistart method and as the number of samples increases (parameter M) the method requires lower amount of function calls to estimate the global minimum. This means that the method tends to create more accurate clusters (clusters that emulate the true regions of attraction) of the objective function as the number of samples increases. To fully demonstrate it an additional run has been performed using different values for the samples and constant number of centers: in Figure 1 the number of required function calls for Function EXP4 are plotted against the number of samples. The number of samples varies between 100 and 900 and the number of centers remains $K = 100$ for all the runs. As is is evident, the number of function calls remains constant, regarding the increase of samples. Also, to compare the proposed method with M=100 with the M=200 for different functions, the Wilcoxon signed-rank test was used. The results obtained with this statistical test are shown in Figure 2.

Table 1: The values for the parameters used in the conducted experiments.

PARAMETER	VALUE
K	100
$K_{\max}$	100
F	1.5
$N_{\min}$	3

Figure 1: Plot for the EXP4 functions as the number of samples increases.

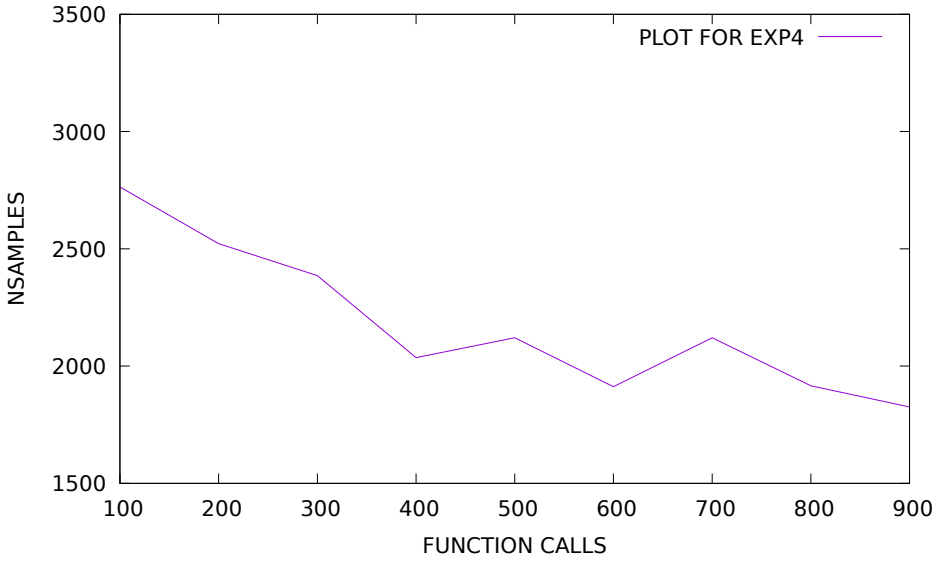


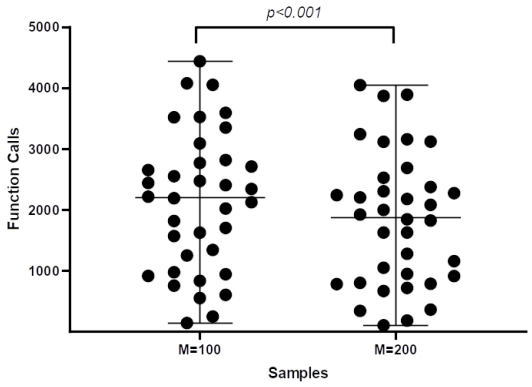
Table 2: Multistart results.

Function	$M = 100$	$M = 200$	$M = 100$,Rejection
B2	4518	8849	4472
Easom	943	1949	933
Bf1	4508	9300	4469
Bf2	3750	7621	3666
Branin	1948	3855	1938
Camel	2669	4983	2502
CM4	5714	11783	5644
CM8	7341(0.33)	14813(0.60)	7289(0.33)
DIFFPOWER10	123729	248924	121012
ELP4	1203	2474	1158
ELP8	1721	3395	1652
ELP16	2789	5485	2252
EXP4	3646	7063	3609
EXP8	3723	7447	3651
EXP16	3835	7486	3310
GKLS250	1486	2928	1426
GKLS350	1030(0.97)	2007	913(0.87)
GKLS3100	1020(0.77)	2005	1018(0.77)
GRIEWANK2	3131(0.70)	6197(0.97)	3048(0.70)
GRIEWANK10	10449	20763	10226
HANSEN	2482	4997	2422
HARTMAN3	2911	5753	2868
HARTMAN6	3825	7875	3787
POTENTIAL3	5237	10784	5178
POTENTIAL5	11594	22331	10127
POTENTIAL10	20361	40592	5089(0.70)
RASTRIGIN	2345	4731	2242(0.93)
SHEKEL5	3852	7841	3730
SHEKEL7	3951	7149	3885
SHEKEL10	3982	6987	3890
SINU4	3317	6624	3246
SINU8	4883	10015	4791
SINU16	8731	17005	8692
TEST2n4	3258	6608	3216
TEST2n5	3565	7128	3534
TEST2n6	3804(0.90)	7790	3850(0.90)
TEST2n7	4203(0.83)	8501(0.97)	4155(0.77)
TOTAL	**281454(0.93)**	**562038(0.98)**	**259160(0.92)**

Table 3: The proposed method with $K = 100$ centers.

Function	$M = 100$	$M = 200$
B2	4073	3886
Easom	830	782
Bf1	4046	3864
Bf2	3346	3153
Branin	1699	1623
Camel	2338	2237
CM4	4434	4043
CM8	3084(0.63)	1819(0.50)
DIFFPOWER10	26726	17980
ELP4	971	908
ELP8	601	338
ELP16	139	100
EXP4	2764	2522
EXP8	1564	943
EXP16	245	179
GKLS250	1337	1275
GKLS350	911(0.93)	777(0.83)
GKLS3100	939(0.97)	796(0.97)
GRIEWANK2	2812(0.77)	2684(0.70)
GRIEWANK10	1812	1152(0.80)
HANSEN	2210	2077
HARTMAN3	2400	1993
HARTMAN6	2707	2369
POTENTIAL3	1246	714
POTENTIAL5	752	664
POTENTIAL10	1621(0.23)	1045(0.10)
RASTRIGIN	2016	1917
SHEKEL5	3520	3116
SHEKEL7	3515	3113
SHEKEL10	3586	3237
SINU4	2548	2268
SINU8	2121	1624
SINU16	546	358
TEST2n4	2436	2198
TEST2n5	2186(0.97)	1840(0.97)
TEST2n6	2648(0.80)	2300(0.83)
TEST2n7	2469(0.77)	2173(0.73)
TOTAL	**103198(0.95)**	**84067(0.93)**

Figure 2: Scatter plot representation and Wilcoxon rank-sum test results of the comparison between the samples M=100 with the M=200 for different functions. A p-value of less than 0.05 (2-tailed) was used for statistical significance.



4 Conclusions

A new clustering method was introduced in this article to tackle to global optimization problem. For every cluster gradually a representative is created using the well - known Kmeans method. Afterwards, the clusters are reduced in number using a simple rejection procedure. The proposed method was tested on a series of benchmark problems from the relevant literature and it is compared against the Multistart method and the results are reported. The proposed method outperforms traditional multistart due to creation of the centers, that seem to be more accurate representatives of the regions of attractions for the underlying function. To achieve better representation of these regions of attractions, the centers are gradually improved through repetitive sampling of the objective function.

Despite the promising results, the method has some limitations that can be addressed in future research. First of all the method depends heavily on the Kmeans algorithm. The method creates K clusters that are enhanced through sampling, but this processes can take long especially for problems of higher dimension. A possible solution could be to use parallel techniques for the estimation of the clusters. Also, the method requires an excessive memory storage just to hold the member points for the K clusters and some efficient memory allocation mechanism should be incorporated here.

Judging from the reported results, the proposed method seems to be very promising and a series of enhancements could be applied on the method such as:

1. Dynamic selection of K in Kmeans algorithm.

2. Better estimation of the critical distance between clusters in the rejection procedure.

3. Usage of more efficient stopping rules to prevent the method from unnecessary local searches, that could lead to the same global optimum many times.

Compliance with Ethical Standards

All authors declare that they have no has no conflict of interest.

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