

Opleiding Informatica

Applying a Mixed-integer evolutionary strategy

for the configuration and parameterization of a CMA-ES

Jelle Lubben

Supervisors: Prof. dr. Thomas Bäck & Sander van Rijn

BACHELOR THESIS

Leiden Institute of Advanced Computer Science (LIACS) www.liacs.leidenuniv.nl

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Abstract

This paper will focus on finding improvements in a Covariance Matrix Adaption Evolution Strategy (CMA-ES). We will make use of a framework that allows us to combine the many variants of the CMA-ES to find the best combination. The configuration of the framework consist of a mix of discrete options and parameter values. To find a optimization for both the framework and the parameters we will make use of a Mixed Integer Evolution Strategy(MIES). We made an attempt to find improvements over previous results. However while the combination of MIES and the CMA-ES were better then common variants, previous experiments that did not involve parameter optimizations were mostly better with the same amount of resources.

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Introduction

Genetic Algorithms (GA) and Evolutionary Strategies (ES) have been proven useful in solving a wide variety of optimization problems. This paper will focus on the Covariance Matrix Adaption Evolutionary Strategy(CMA-ES). Recent study [1] has shown that optimizing the configuration of a CMA-ES using a GA is a useful way of improving it. The configurations found by the GA have been proven better than any literature described configuration for the CMA-ES. Our goal is to explore if even more improvement can be found. In the previous study only eleven configuration options were altered during the process. We add the ability to alter the remaining continuous and integer parameters, in order to explore the performance of tuned configurations.

1.1 Applications

We will make use of the original framework from [1]. To find the best configuration and parameterization for the CMA-ES, we will use a Mixed Integer Evolution Strategy(MIES) [2]. The MIES is an ES that can be used to find an approximately optimal solution for a problem whose solution consists of discrete, integer and continuous values. The configuration for the CMA-ES consists of eleven discrete values, one integer value and fourteen continuous values.

1.2 Thesis Overview

The thesis consists of the problem description in Chapter 2. Next we will explain the techniques used in solving the problem in the Chapter 3. The experiments and results will be shown in the Chapter 4 and the results will be discussed in Chapter 5. Finally, in Chapter 6 we present some ideas for future work.

Problem

Our goal is to find the best algorithm for any given optimization problem. Specifically to explore if improvements can be found over previous results by van Rijn et al [1]. We do this by comparing our MIES-based results to those of the original research. For testing the optimization, we use the common Black Box Optimization Benchmark (BBOB) suite [3].

2.1 BBOB

The BBOB suite is a standardized collection of 24 test functions that are representative for real-world optimization problems. This allows us to compare optimization algorithms by comparing their performance results on the test problems in the BBOB suite.

The 24 functions are split into four subgroups:3 unimodal and separable,7 unimodal and non-separable, 2 multimodal and separable and 12 multimodal and non-separable. Examples of these subgroups can be found in Figure 2.1.

Unimodal means that only a single global optimal point in the function exists. Multimodal means that there are multiple local optima.

A function is called separable if the global optimal point composed of all optimal points from each dimension. In other words, the function can be optimized independently for each dimension. Non-separable means that the global optimal point cannot be optimized individually for each dimension.







Figure 2.1: Examples of the four function subgroups in the BBOB suite.

Evolution Strategies

There are many forms of Evolution strategies. In this paper we will be using two different ones. The CMA-ES for solving the benchmark functions and the MIES for determining the configuration of the CMA-ES.

3.1 Standard Evolution strategy

An evolution strategy is based on biological evolution. Initially, individual are created as possible solutions for an optimization problem. These initial individuals will be the first generation. Each individual is evaluated using a fitness function.By recombining the best individual and applying mutation a new generation is created. The idea is that each generation will produce better results then its predecessor. This process will repeat itself until a fitness objective is met or the resources run out.

3.2 CMA-ES

What makes a CMA-ES different from other ES's is that it uses a covariance matrix to adapt the distribution of mutations during the search process. This covariance matrix is updated each generation. For a more detailed description of the CMA-ES, please refer to [4]. A 2-dimensional example is shown in figure 3.1 The framework we use for the CMA-ES [4] has eleven independent modules that each have two or three options resulting in 4608 different module combinations. Some of these modules have associated continuous parameters. Some parameters are not dependent on a module but are simply parameters of the basic CMA-ES. These parameters are the integer parameter population size μ and the continuous parameters α_{mu} , c_{σ} , damps, c_c , c_1 and c_{mu} .

The eleven modules and their respective parameters are as follows:



Figure 3.1: The black dots represent the individuals generated by the ES. The white rings mean equal fitness in the search space. The white circle is the global optimum, and the orange dotted line shows the sampling distribution described by the covariance matrix.

1. Active update

Active update uses not only the best individuals in the population, but also the least favored individuals in order to adapt the covariance matrix. Without active update, only the best individuals will be used. The Active Update is introduced by Jastrebski el al. [5].

2. Elitism

Elitism allows individuals from previous generations to survive and reproduce if no better solutions are found. If turned off the algorithm is forced to choose an new set of parents from the current generation. If this option is enabled $(\mu + \lambda)$ is used. If this option is disabled (μ, λ) is used.

3. Mirrored Sampling

Mirrored sampling is used to evenly spread the sampling of the search space. First sampling half of the intended number of mutation vectors, and then using those and their mirror images to create the new candidates. [6].

4. Orthogonal Sampling

Another sampling options is orthogonal sampling [7]. The desired number of samples is first drawn from the usual distribution. Then the Gram-Schmidt process is used to orthonormalize the set of vectors. Although orthogonal sampling was introduced in combination with Mirrored Sampling, both can be used independently in this framework.

5. Sequential Selection

Normally the entire new generation is evaluated before selecting the new best individual. However Sequential Selection stops the evaluation of further offspring once an improvement has been found. [6].

6. Threshold Convergence

Threshold Convergence can be used to prevent the the ES to get stuck on a local optimum. [8]. Threshold convergence forces the ES to stay in the exploratory phase for a longer time. It does this by requiring the mutation vectors to be of greater or equal length than a certain threshold. The decrease over time of this threshold is governed by associated continuous parameters 'initial-threshold' and 'decay-factor'.

7. Two-Point Step-Size Adaptation (TPA)

TPA determines if the step-size of the CMA-ES should increase or decrease [9]. It does this by creating two additional candidates after the selection step. They are sampled along the direction of the weighted average mutation vector, one shorter and one longer. The step size is increased if the candidate with longer mutation step is more successful, otherwise the step size is decreased. It makes use of the following continuous parameters:'tpa-factor', ' β_{tpa} , c_{α} and α .

8. Pairwise Selection

In order to prevent mirrored-sampling pairs from (partially) cancelling each other out when updating the covariance matrix or step size, Pairwise Selection as introduced by Auger, et al. [10] can be used. It selects the best offspring from each pair, and then regular selection is applied to that set. Although Pairwise Selection was introduced together with Mirrored Sampling and Orthogonal Sampling, all can be used individually.

9. Recombination Weights

In the CMA-ES, recombination is performed with the following weight vector: $w_i = log(\mu + \frac{1}{2}) - \frac{log(i)}{\Sigma_j w_j}$ for i, \dots, μ . Alternative weights are the arithmetic mean $w_i = \frac{1}{\mu}$.

10. Quasi-Gaussian Sampling

Instead of drawing samples from a Gaussian distribution sampling they can also be drawn from a Quasi-random uniform sequence, which are then transformed to a Gaussian distribution. This was proposed in [11]. Available Quasi-random sequences are the 'Sobol' and 'Halton' sequences.

11. Increasing Population Size (IPOP/BIPOP)

With no improvement in recent generations one can restart an ES with increasing generation size(IPOP) [12]. Alternatively bi-population(BIPOP) [13] can be used. BIPOP varies between increasing and decreasing population size. Both IPOP and BIPOP make use of the continuous parameter 'population-increase-factor'.

3.3 MIES

Mixed Integer Evolution Strategies(MIES) are designed for optimizing problems where the objective function uses a mix of continuous, integer and discrete values. As a regular ES assumes that the candidate consists of only of continuous values, typical mutation would not work for discrete and integer values. MIES has a separate mutation process for each set of values.

TABLE I: A table showing all continuous and integer variables and the associated module. λ is the only integer value, all the other values are continuous. μ is shown as a percentage of λ

Integer/Continuous variable	default values	Associated module
λ	$4 + (3 \cdot \log(n))$	
μ	0.5	
α_{mu}	2	
$\mathcal{C}_{\mathcal{O}}$	$\frac{\mu_{eff}+2}{\mu_{eff}+n+5}$	
damps	$1 + 2 \cdot \sqrt{\frac{\mu_{eff} - 1}{n+1}} - 1 + c_{\sigma}$	
C _C	$\frac{(4+\mu_{eff}/n)}{(n+4+2\cdot\mu_{eff}/n)}$	
<i>c</i> ₁	$\frac{2}{(n+1.3)^2 + \mu_{eff}}$	
C _{mu}	$min(1-c_{1})$,	
	$\alpha_{mu} \cdot \frac{\mu_{eff} - 2 + 1/\mu_{eff}}{(n+2)^2 + \alpha_{mu} \cdot \mu_{eff}/2} \big)$	
initial-threshold	0.2	Threshold Convergence
decay-factor	0.995	Threshold Convergence
tpa-factor	0.5	TPA
β_{tpa}	0	TPA
C_{α}	0.3	TPA
α	0.5	TPA
'population-increase-factor'	2	IPOP/BIPOP.

The candidate solution representation is split into three subsets. Let *S* denote the entire representation of a candidate solution. Then D^n , I^m and R^k are the sets of discrete, integer and continuous values from S respectively. Thus an individual in the MIES can be represented by $S = D^n I^m R^k$. In the remainder of this section we will elaborate on the mutations of each of these three sets of values.

3.3.1 Nomenclature

In the next sections the following definitions are used:

N(a, b) denotes a normally distributed random number with expectation *a* and standard deviation *b*.

U(a, b) is a uniformly distributed number between *a* and *b*.

T[a, b](x) is the transformation function, as proposed in [2] that keeps the value x between upper and lower bound a and b. The algorithm is shown in Algorithm 1.

Algorithm 1 Transformation function

1: y = (x - a)/(b - a)2: if $\lfloor y \rfloor \mod 2 = 0$ then 3: $y = |y - \lfloor y \rfloor|$ 4: else 5: $y = 1 - |y - \lfloor y \rfloor|$ 6: end ifreturn y

3.3.2 Discrete values

Each discrete parameter in D^n has its own step-size value. The step-size is represented as $P_1 \cdots P_k$ with P_i the step-size of D_i . As shown in algorithm 2 in step 3-5 all step-sizes are updated. Then if U(0,1) is lower then the step-size, the discrete value is randomly chanced to a different possible value in step 6-7.

Algorithm 2 Discrete values

1: $N_c = N(0, 1)$ 2: Initialize global and local learning curve $\tau = \frac{1}{\sqrt{2k}}$; $\tau' = \frac{1}{\sqrt{2\sqrt{k}}}$ 3: for all $i \in \{1, \dots, n\}$ do 4: $P_i = \frac{1}{1 + \frac{1-P_i}{P_i} \cdot exp(-\tau \cdot N_c - \tau' \cdot N(0,1))}$ 5: $P_i = T[\frac{1}{3n}, 0.5]P_i$ 6: if $U(0, 1) < P_i$ then 7: choose random from set of possible discrete values without the current value of the discrete 8: end if 9: end for

3.3.3 Integer Values

A individual step-size is kept for each integer parameter. The step-size is represented as $\varsigma_1 \cdots \varsigma_m$ with ς_i the step-size of I_i .

Algorithm 3 shows the mutation procedure for integer values. First the step-size is updated in step 4. Then variables are created in order to create G_1 and G_2 . Which are then added and subtracted from the value of the initial integer respectively.

Algorithm 3 Integer values

1: $N_c = N(0, 1)$ 2: Initialize global and local learning curve $\tau = \frac{1}{\sqrt{2k}}$; $\tau' = \frac{1}{\sqrt{2\sqrt{k}}}$ 3: for all $i \in \{0, \dots, m\}$ do 4: $\varsigma_i = max(1, \varsigma_i exp(\tau N_c + \tau' N(0, 1)))$ 5: $u_1 = U(0, 1); u_2 = U(0, 1)$ 6: $\psi = 1 - (\varsigma_i/m)(1 + \sqrt{1 + (\frac{\varsigma_i}{m})})^{-1}$ 7: $G_1 = \lfloor \frac{ln(1-u_1)}{ln(1-\psi)} \rfloor$ 8: $G_2 = \lfloor \frac{ln(1-u_2)}{ln(1-\psi)} \rfloor$ 9: $I_i = I_i + G_1 - G_2$ 10: end for

3.3.4 Continuous Values

The mutation of the continuous values is similar to integer mutation. A individual step-size is kept for each continuous parameter. As shown in algorithm 4, the step-size is updated with the learning curve and added to the previous continuous value. The value of the parameter is then kept in-bounds by the transformation function T. The step-size is represented as $\sigma_1 \cdots \sigma_k$ with σ_i the step-size of R_i .

Algorithm 4 Continuous values

1: $N_c = N(0, 1)$ 2: Initialize global and local learning curve $\tau = \frac{1}{\sqrt{2k}}$; $\tau' = \frac{1}{\sqrt{2\sqrt{k}}}$

- 3: for all $i \in \{0, \dots, k\}$ do 4: $\sigma_i = \sigma_i \cdot exp(\tau N_c + \tau' N(0, 1))$ 5: $R_i = Ri + \sigma i$ 6: $R_i = T(max_i, min_i)R_i$ 7: end for

Experiments

The experiments reported in this thesis are based on the ideas discussed in [1]. The key difference between our study and [1] is that the MIES changes all the parameters. As it is well known that tuned algorithms perform better. These experiments test the possibility of tuning the CMA-ES while optimizing its structure at the same time.

The experiments are run on all 24 Black Box Optimization Problems. The evolutionary strategy was executed om 2,3 and 5 dimensional problems. All test are run 10 times to get a solid estimate of how good this approach works.

4.1 Settings

The previous experiments of [1] use a generation size of 12 and a budget of 20 generations for the Genetic Algorithm (GA) that optimizes the configuration. Although the search-space is larger due to the inclusion of the CMA-ES parameters, we choose to give the MIES the same budget of 20 generations. This is done both to facilitate a more direct comparison with previous results, and due to time-constraints.

MIES sets the configuration of the CMA-ES. In order for the MIES to properly work, all integer and continuous values of the CMA-ES need to have an upper and lower bound. For the single integer value λ (population size), we use a minimum of 2 and a maximum of 200. It is argued that with a budget of $1000 \cdot D$ (where D denotes the dimensionality of the problem), a larger population size would probably lead to too few generation. For continuous values we define the range as $[0.7\sigma_i, 1.3\sigma_i]$ where σ_i are the default values from literature as mentioned in Chapter 3. A standard range of [0,1] was used if a default value was o.

4.2 Conditional mutation

Roughly half of the continues variables are uniquely used by a discrete option. If they are mutated while the discrete option is not enabled, we will get no feedback on this continuous value and potentially lose information. Therefore if a discrete option is disabled all continuous values that depend on the option will remain unaltered. All discrete options with the corresponding continuous values can be found in Table I.

4.3 FCE and ERT

In order to make the results measurable we make use of the same criteria as in [1]. Fixed Cost Error (FCE) and Expected Run Time (ERT) will be used. FCE is the lowest distance to the global optimum after using fixed computation cost. The target fitness is set at 10^{-8} . If a run reached the target within the given resources it has been successful, then the FCE will be 10^{-8} . The ERT gives more information but can only be calculated if at least one of the runs reached the target fitness. If none of the runs reach the target fitness, the ERT is considered invalid and only FCE can be used.

ERT is calculated as follows:

$$ERT(f_{target}) = RT_S \frac{1 - p_s}{p_s} RT_{US}$$

With RT_S the mean of r_s , and r_s rated with the number of function evaluations needed to reach f_{target} . Furthermore, let RT_{US} be the mean of r_{us} with r_{us} runs that did not reach f_{target} and are considered unsuccessful. These are rated with the full run time. The ratio of successful runs to all runs yields the value p_s . In words, the ERT is the time you expect the CMA-ES to take in order to reach the taget value f_{target} .

As the goal of an optimizer is to find the best possible (i.e. minimal) fitness value in the least amount of time, we want to minimize both FCE and ERT measures. Comparisons between two configurations are done on FCE only if neither have valid ERT values, otherwise the only valid/better ERT wins.

4.4 Results

As you can see in the graphs of F1,F3 and F6 for example, the FCE line is horizontal with a value of 1e-8. This is because that is the target value of the fitness function and therefore the minimum(best) result. If the target value has been found in the first generation, the FCE will most likely stay at that value.

The FCE can still rise during the process(for example, F₃, F₅ and F₁₆). This is because in that generation only higher values were found.







Figure 4.2: plots of F3 - F4

However, the ERT (Expected Run Time) will not increase because, only when improvements are found the ERT can be updated.

While most improvements of ERT are found in later generation, it is very possible for the best solution to be found somewhere in the first few generation. For example F10 3-dim and 5-dim show no improvement after generation 2, and F10 2-dim has reached its best result in generation 1.

However it can be discussed that better results will be found after the initial 20 generations.

If we look at F6 2-dim, we can see that there are multiple moments where improvements were found. And we could argue that given more generations, even better results will be found.

It is often that a higher dim results in a higher ERT. In almost all cases ERT of 5-dim is higher then 3-dim which in turn is higher then 2-dim. However this is not always the case. F10 and F18 3-dim has a higher ERT then 5-dim. It would make sense that a higher dimensionality would cost more run time.

All the data has been processed into graphs and tables. The graph shows the average improvements over 10 runs. And the tables show the best results found. In 4.1 - 4.12 the results of 20 generations with F1- F24 in 2, 3 and 5 dimensions are shown.

All best results are shown in Table I and Table II. The results are compared to previous results in Table III and Table IV to the results from [1].









105

10³

ò

2-dim 3-dim 5-dim

10 Generation







Figure 4.6: plots of F11 - F12

















Figure 4.10: plots of F19 - F20









Conclusion

Looking at table III and table IV in the CV/MIES column you can clearly see that the MIES mostly performs better then the Common Variant. If an improvement has been found the value is higher then 1. If the value is lower, the MIES failed to find better results. When comparing the MIES against the GA, results aren't as clear. While some improvements are found (for example F18 dim 5 in table IV), most results of the GA are still better or somewhat the same. Of the total 72 results, 9 improvements were found, 9 results are somewhat the same (between 0.95 and 1.05) and 54 are worse.

The reason that most results are worse is most likely due to the fact that there are too many parameters and discrete values to calculate within the given calculation time. While the GA only needed to optimize 11 discrete values, the MIES had to optimize 15 additional parameters. This causes the MIES to create results that are mostly worse then the GA.

Future research

There are some possible improvements in this research. Because there are so many parameters that influence the result, what could be done is running 20 generations to find the best discrete value combination for the CMA-ES configuration and using another 20 generations for parameter optimization.

Right now all continuous values have been set at a predetermined setting and given a limit 30% up and down of that value. It would be possible to find better results in the long run if the limits are allowed to optimize in a wider range

The discrete options IPOP/BIPOP both use the same parameter 'increase-population-size'. This means that if IPOP is being used, the parameter for BIBOP will change. This could give undesirable results. a suggestion would be to split the parameter 'increase population size' into two different parameters. There are still experiments from 10 and 20 dimensionality that can be tested and compared.

TABLE I: All best found by the MIES F1 -F12. discrete values represent the same option as in the previous experminents from [1]. The first number in the next colom is the population size λ . The discrete values from left to right are μ , $\alpha_{mu} c_{\sigma}$, damps, c_c , c_1 , c_{mu} , 'initial-threshold', 'decay-factor', 'tpa-factor, 'beta-tpa','c-alpha', 'alpha' and population-increase-factor.

F-ID	N	GA-discrete							1	paramet	ers							FCE	ERT
F1	2	00011010000	18	0.068	2.43	0.1	1.285	0.13	0.003	0.654	0.2	0.955	0.383	0.462	0.214	0.546	2.133	1e-08	157.72
F1	3	00110010112	11	0.062	1.854	0.077	0.838	0.153	0.002	0.664	0.227	0.906	0.408	0.053	0.254	0.585	2.564	1e-08	251.91
F1	5	01011010112	11	0.059	1.748	0.097	1.181	0.129	0.002	0.255	0.2	0.955	0.415	0.192	0.221	0.441	2.254	1e-08	453.84
F2	2	00010011002	8	0.101	2.22	0.101	1.373	0.153	0.002	0.597	0.2	0.955	0.392	0.621	0.375	0.383	2.209	1e-08	426.44
F2	3	10010011001	9	0.092	1.96	0.098	1.113	0.096	0.003	0.454	0.151	0.885	0.426	0.298	0.366	0.422	1.515	1e-08	878.13
F2	5	01000010012	21	0.072	1.797	0.068	1.299	0.152	0.002	0.131	0.2	0.955	0.527	0.473	0.371	0.379	1.81	1e-08	168.00
F3	2	01011010112	52	0.076	2.211	0.104	0.782	0.142	0.002	0.958	0.2	0.955	0.559	0.239	0.215	0.434	2.381	1e-08	3213.44
F3	3	10111011021	111	0.098	1.731	0.091	1.396	0.121	0.002	0.857	0.2	0.955	0.386	0.388	0.314	0.463	1.817	0.0679	12877.64
F3	5	00000010001	146	0.091	2.588	0.072	1.402	0.137	0.003	0.324	0.2	0.955	0.401	0.126	0.278	0.462	1.935	0.308	47132.73
F4	2	11110000011	18	0.108	1.56	0.087	1.154	0.174	0.002	0.141	0.2	0.955	0.374	0.545	0.28	0.469	1.713	0.0509	7054.41
F4	3	00101000022	13	0.061	2.176	0.066	1.226	0.124	0.002	0.127	0.253	0.936	0.601	0.246	0.328	0.394	2.211	0.887	62078.69
F4	5	11100000102	45	0.101	1.936	0.08	0.823	0.108	0.003	0.283	0.2	0.955	0.441	0.687	0.35	0.571	2.587	3.48	1553072.00
F5	2	10101100022	25	0.085	1.523	0.103	1.228	0.126	0.003	0.893	0.223	0.818	0.5	0	0.3	0.5	2.196	1e-08	505.25
F5	3	10111000020	24	0.078	2.284	0.121	1.35	0.155	0.002	0.329	0.141	0.755	0.605	0.024	0.339	0.566	1.756	1e-08	809.13
F5	5	01101000001	14	0.083	1.705	0.116	1.416	0.108	0.002	0.106	0.254	1.052	0.5	0	0.3	0.5	2.31	1e-08	1772.66
F6	2	11010001112	7	0.093	2.59	0.093	1.184	0.169	0.003	0.965	0.2	0.955	0.471	0.32	0.281	0.36	2.154	1e-08	334.72
F6	3	11110010010	13	0.102	2.486	0.114	1.066	0.145	0.002	0.218	0.2	0.955	0.405	0.298	0.373	0.446	1.888	1e-08	566.69
F6	5	00110001120	6	0.087	2.037	0.072	0.812	0.119	0.002	0.06	0.2	0.955	0.466	0.018	0.302	0.571	2	1e-08	1099.38
F7	2	11001010012	14	0.086	2.392	0.118	1.009	0.096	0.002	0.184	0.172	0.787	0.62	0.809	0.221	0.448	2.318	1e-08	268.63
F7	3	00000011121	12	0.08	2.061	0.077	1.393	0.125	0.003	0.695	0.2	0.955	0.471	0.361	0.217	0.472	2.367	1e-08	564.66
F7	5	10000010021	94	0.072	1.455	0.091	1.301	0.173	0.002	0.675	0.195	0.754	0.384	0.947	0.279	0.431	2.25	1e-08	1270.25
F8	2	01001000100	5	0.101	1.983	0.093	0.939	0.101	0.002	0.44	0.229	0.727	0.423	0.347	0.211	0.52	1.933	1e-08	506.09
F8	3	00000111002	2	0.096	1.538	0.092	1.197	0.137	0.002	0.059	0.153	0.902	0.603	0.746	0.332	0.615	1.67	1e-08	1041.81
F8	5	11000111111	12	0.075	2.067	0.078	1.323	0.118	0.002	0.01	0.234	0.737	0.482	0.591	0.383	0.456	2.579	1e-08	2406.09
F9	2	01001011011	9	0.099	2.049	0.113	1.072	0.118	0.002	0.21	0.219	0.899	0.648	0.023	0.222	0.496	1.903	1e-08	398.00
F9	3	11010010011	16	0.081	1.632	0.098	0.998	0.132	0.003	0.514	0.2	0.955	0.575	0.058	0.289	0.465	2.096	1e-08	959.75
F9	5	00110010011	20	0.074	1.592	0.113	1.416	0.162	0.002	0.442	0.2	0.955	0.501	0.02	0.31	0.552	1.492	1e-08	2307.69
F10	2	11000000110	4	0.101	2.491	0.088	0.844	0.117	0.003	0.07	0.183	0.779	0.471	0.222	0.365	0.529	2.207	1e-08	502.38
F10	3	10110011101	16	0.082	2.006	0.113	1.2	0.162	0.002	0.523	0.2	0.955	0.488	0.562	0.297	0.399	2.59	1e-08	956.50
F10	5	00110010002	20	0.093	1.452	0.119	0.937	0.154	0.002	0.165	0.228	0.789	0.507	0.571	0.251	0.432	1.902	1e-08	1915.88
F11	2	10011010020	24	0.108	1.734	0.093	0.927	0.149	0.002	0.81	0.218	0.979	0.645	0.561	0.287	0.457	1.958	1e-08	448.65
F11	3	00111010011	13	0.058	2.557	0.103	1.132	0.163	0.003	0.478	0.2	0.955	0.428	0.081	0.365	0.51	1.442	1e-08	864.66
F11	5	00001010012	22	0.077	2.312	0.112	1.033	0.161	0.003	0.308	0.164	0.798	0.649	0.199	0.317	0.429	1.532	1e-08	1272.06
F12	2	11001011122	23	0.099	1.525	0.096	0.908	0.125	0.003	0.336	0.247	0.867	0.522	0.729	0.219	0.476	1.68	1e-08	1126.00
F12	3	01111010102	22	0.099	2.52	0.116	1.119	0.122	0.002	0.341	0.143	1.164	0.63	0.069	0.222	0.37	2.042	1e-08	2672.03
F12	5	01010010022	12	0.085	1.462	0.076	0.765	0.161	0.002	0.272	0.142	0.795	0.471	0.114	0.356	0.419	2.049	1e-08	6082.81

TABLE II: All best found by the MIES F13 -F24. discrete values represent the same option as in the previous experminents from [1]. The first number in the next colom is the population size λ . The discrete values from left to right are μ , $\alpha_{mu} c_{\sigma}$, damps, c_c , c_1 , c_{mu} , 'initial-threshold', 'decay-factor', 'tpa-factor', 'calpha', 'calpha', 'alpha' and population-increase-factor.

E-ID	N	CA-discrete							r	aramot	ore							ECE	FRT
Eto		10010010112	01	0.062	2 2 2 7	0.082	1.208	0.1.11	1	0 520	0.2	0.055	0 ==6	0.00	0.256	0.055	2.286	10.08	600.25
F13 Eco	2	10010010112	21	0.002	2.237	0.062	1.290	0.141	0.002	0.539	0.2	0.955	0.550	0.09	0.250	0.355	2.200	10-00	099.25
Г13 Бар	3	01000010101	0	0.104	2.359	0.07	0.921	0.130	0.001	0.239	0.2	0.955	0.501	0.25	0.216	0.447	2.437	10-00	1400.91
Г13 Бал	5	10110010021	30	0.091	1.925	0.110	1.279	0.110	0.001	0.295	0.2	0.955	0.500	0.200	0.339	0.426	1.994	10-00	33/1.44
114 Eau	2	00110011000	9	0.009	2.302	0.090	1.321	0.112	0.002	0.522	0.199	1.034	0.467	0.713	0.339	0.030	1.075	10-00	571.100
Г14 Бал	3	00111000000	12	0.064	2.013	0.101	0.014	0.129	0.003	0.257	0.2	0.955	0.5	0	0.3	0.5	2.201	10-00	093.75
F14 E15	5	10010010020	14	0.1	2.000	0.113	1.335	0.107	0.002	0.013	0.229	1.117	0.300	0.143	0.211	0.353	1 8 2 1	10.08	1745.01
F15	2	10010010022	17	0.071	1.050	0.103	1.105	0.140	0.003	0.002	0.100	0.920	0.034	0.53	0.32	0.401	1.021	10-00	2013.91
F15	3	11101110012	71	0.087	2.301	0.098	1.154	0.162	0.003	0.043	0.202	0.871	0.366	0.828	0.357	0.495	1.982	16-08	8473.56
F15	5	00100010101	154	0.093	2.125	0.077	0.818	0.097	0.002	0.188	0.2	0.955	0.411	0.998	0.224	0.535	2.05	0.0938	21929.10
F10	2	11110010001	10	0.087	1.491	0.111	1.111	0.169	0.002	0.473	0.141	1.217	0.499	0.118	0.326	0.635	1.477	16-08	1171.25
F10	3	00100010001	53	0.078	1.56	0.11	1.263	0.118	0.001	0.288	0.2	0.955	0.551	0.937	0.369	0.536	1.475	16-08	2091.88
F16	5	00111010002	111	0.096	1.781	0.079	1.231	0.161	0.003	0.603	0.2	0.955	0.553	0.854	0.335	0.424	1.499	10-08	6479.91
F17	2	00110011122	28	0.083	1.63	0.09	1.303	0.144	0.002	0.821	0.247	1.192	0.484	0.38	0.215	0.635	2.387	1e-08	1397.16
F17	3	00110110122	44	0.07	2.479	0.075	0.966	0.094	0.002	0.019	0.167	0.825	0.565	0.473	0.363	0.502	2.177	1e-08	2346.06
F17	5	00100011002	87	0.069	1.646	0.091	0.849	0.153	0.002	0.354	0.21	0.785	0.4	0.446	0.336	0.504	1.822	1e-08	5595.91
F18	2	11000011001	7	0.105	2.207	0.102	0.793	0.153	0.001	0.388	0.2	0.955	0.534	0.125	0.35	0.571	1.572	1e-08	1723.22
F18	3	11010011112	13	0.101	2.007	0.067	0.964	0.136	0.003	0.314	0.2	0.955	0.43	0.144	0.383	0.403	2.433	1.02	3904.69
F18	5	10010010022	13	0.091	2.487	0.08	1.233	0.156	0.002	0.246	0.2	0.955	0.541	0.028	0.23	0.42	2.292	0.000000449	9190.69
F19	2	11010010112	15	0.082	2.127	0.081	1.074	0.106	0.003	0.491	0.2	0.955	0.387	0.093	0.252	0.438	2.197	1e-08	3460.33
F19	3	00101010022	119	0.066	2.222	0.084	1.314	0.167	0.002	0.808	0.2	0.955	0.577	0.656	0.317	0.411	2.207	0.0191	15316.14
F19	5	10000011021	126	0.08	1.887	0.102	0.999	0.169	0.002	0.724	0.2	0.955	0.526	0.237	0.342	0.501	1.808	0.144	71535.75
F20	2	00001010011	9	0.059	2.446	0.107	0.901	0.137	0.003	0.308	0.2	0.955	0.481	0.824	0.252	0.527	1.762	1e-08	2330.81
F20	3	00101010022	9	0.068	1.626	0.103	1.416	0.1	0.003	0.32	0.2	0.955	0.644	0.262	0.33	0.407	2.415	0.000172	7668.806
F20	5	11011110011	103	0.097	2.319	0.113	1.396	0.103	0.002	0.09	0.252	1.042	0.515	0.453	0.297	0.446	2.436	0.258	37833.27
F21	2	11011110022	11	0.064	1.9	0.087	1.141	0.094	0.002	0.489	0.256	1.129	0.57	0.908	0.24	0.505	2.515	1e-08	590.19
F21	3	11100110102	9	0.087	2.506	0.085	0.789	0.155	0.003	0.554	0.237	0.774	0.45	0.324	0.28	0.477	2.544	1e-08	1842.44
F21	5	00100010122	12	0.086	2.115	0.103	1.214	0.104	0.002	0.367	0.2	0.955	0.599	0.337	0.321	0.479	1.887	0.102	7702.25
F22	2	11000110112	18	0.103	2.572	0.101	0.886	0.166	0.002	0.551	0.193	0.95	0.506	0.443	0.259	0.431	1.948	1e-08	434-44
F22	3	01010010111	26	0.065	2.5	0.096	1.298	0.095	0.002	0.614	0.165	1.04	0.447	0.541	0.211	0.573	1.762	3.22	1565.60
F22	5	01001001022	12	0.069	1.567	0.086	0.909	0.155	0.002	0.176	0.2	0.955	0.5	0	0.3	0.5	2.352	0.0804	7979-37
F23	2	00100010001	14	0.095	2.185	0.081	1.229	0.149	0.002	0.294	0.2	0.955	0.359	0.223	0.334	0.554	1.984	1e-08	1664.84
F23	3	11011110022	19	0.084	1.414	0.086	0.924	0.096	0.003	0.022	0.219	1.03	0.639	0.136	0.389	0.514	2.47	0.00443	8568.60
F23	5	10011000111	75	0.074	2.565	0.115	0.767	0.136	0.001	0.088	0.2	0.955	0.481	0.126	0.219	0.412	2.136	0.0622	22807.11
F24	2	00010100111	17	0.085	1.895	0.111	1.173	0.151	0.002	0.54	0.224	0.679	0.382	0.185	0.223	0.399	2.214	0.0113	4016.00
F24	3	00010100021	10	0.097	1.827	0.069	0.908	0.142	0.002	0.142	0.228	0.934	0.359	0.879	0.298	0.518	1.648	2.63	50193.29
F24	5	11011011111	135	0.061	2.573	0.084	0.845	0.156	0.002	0.726	0.2	0.955	0.612	0.037	0.279	0.624	2.392	20	1557063.00

TABLE III: F1 - F12 from the MIES Compared to the best from GA and best from Commen Variant, both from [1]. In the last two coloms, if the value is greater then 1 then there is an improvement. Note: All experiments from the MIES only had 10 runs while both other options had 30 tries.

F	dim	MIES ERT	MIES FCE	GA ERT	GA FCE	CV ERT	CV FCE	GA/MIES	CV/MIES
F1	2	157.72	1e-08	165.25	1e-08	294.75	1e-08	1.048	1.869
F1	3	251.91	1e-08	235.38	1e-08	473.59	1e-08	0.934	1.880
F1	5	453.84	1e-08	440.00	1e-08	797.50	1e-08	0.969	1.757
F2	2	426.44	1e-08	434.06	1e-08	660.375	1e-08	1.018	1.549
F2	3	878.13	1e-08	678.59	1e-08	1160.03	1e-08	0.773	1.321
F2	5	1685.00	1e-08	1364.50	1e-08	2314.75	1e-08	0.810	1.374
F3	2	3213.44	1e-08	1795.73	0.2912	2742.35	0.3582	0.559	0.853
F3	3	12877.64	0.0679	5380.40	0.8374	12346.00	1.062	0.418	0.959
F3	5	47132.73	0.308	29318.80	2.067	76816.00	5.074	0.622	1.630
F4	2	7054.41	0.0509	4089.00	0.8781	11877.60	1.464	0.580	1.684
F4	3	62078.69	0.887	17803.20	3.561	93793.00	4.469	0.287	1.511
F4	5	1553072.00	3.48	155980.00	8.998	N/A	5.905	0.100	1.697
F5	2	505.25	1e-08	466.59	1e-08	605.81	1e-08	0.923	1.199
F5	3	809.13	1e-08	736.66	1e-08	940.69	1e-08	0.910	1.163
F5	5	1772.66	1e-08	1324.28	1e-08	1731.00	1e-08	0.747	0.977
F6	2	334.72	1e-08	315.06	1e-08	428.44	1e-08	0.941	1.280
F6	3	566.69	1e-08	573.38	1e-08	805.00	1e-08	1.012	1.421
F6	5	1099.38	1e-08	1141.78	1e-08	1574.00	1e-08	1.039	1.432
F_7	2	268.63	1e-08	266.00	1e-08	438.75	1e-08	0.990	1.633
F_7	3	564.66	1e-08	540.72	1e-08	886.51	0.002689	0.958	1.570
F_7	5	1270.25	1e-08	1613.56	1e-08	2587.50	1e-08	1.270	2.037
F8	2	506.09	1e-08	422.31	1e-08	574.13	1e-08	0.834	1.134
F8	3	1041.82	1e-08	827.44	1e-08	1034.69	1e-08	0.794	0.993
F8	5	2406.09	1e-08	1705.62	1e-08	2705.43	0.4914	0.709	1.124
F9	2	398.00	1e-08	423.06	1e-08	546.94	1e-08	1.063	1.374
F9	3	959.75	1e-08	764.25	1e-08	1183.66	1e-08	0.796	1.233
F9	5	2307.69	1e-08	1710.22	1e-08	2607.23	0.000213	0.741	1.130
F10	2	502.38	1e-08	441.69	1e-08	662.06	1e-08	0.879	1.318
F10	3	956.50	1e-08	693.50	1e-08	1146.25	1e-08	0.725	1.198
F10	5	1915.88	1e-08	1458.25	1e-08	2228.00	1e-08	0.761	1.163
F11	2	448.66	1e-08	453.19	1e-08	629.63	1e-08	1.010	1.403
F11	3	864.66	1e-08	734.97	1e-08	1334.38	1e-08	0.850	1.543
F11	5	1272.06	1e-08	1518.75	1e-08	2493.00	1e-08	1.194	1.960
F12	2	1126	1e-08	776.07	1e-08	1360.85	0.1986	0.689	1.209
F12	3	2672.03	1e-08	2084.69	0.02808	3426.50	390.5	0.780	1.282
F12	5	6082.81	1e-08	4017.19	1.03e-05	6489.26	0.1195	0.660	1.067

TABLE IV: F13 - F24 from the MIES Compared to the best from GA and best from Commen Variant, both from [1]. In the last two coloms, if the value is greater then 1 then there is an improvement. Note: All experiments from the MIES only had 10 runs while both other options had 30 tries.

EID	N	EDT MIES	ECE MIES	EPTCA	ECE C A	EPT CV	ECE CV		CV/MIES
г-1D F12		EK1-MIES		EKI-GA	гСЕ-GA 10-08	EKI-CV	FCE-CV	GA/ MIES	
F13		1400.02	10.08	882.24	10.08	9/1.01	10.08	0.742	1.390
Г13 Б13	3	1400.92	10-08	003.34	10-08	1704.20	2 8622 08	0.031	1.217
Г13 Гал	5	3371.44	10-00	2967.23	10-08	3030.93	3.0020-00	0.000	1.130
Г14 Бал	2	5/1.19	10-00	410.70	10-08	610.94	10-00	0.719	1.084
Г14 Бал	3	093.75	10-00	630.75	16-08	1196.12	10-00	0.706	1.338
F14	5	1745.81	10-08	1327.00	16-08	2506.75	1e-08	0.760	1.436
F15	2	2813.91	10-08	1881.09	0.2853	2864.47	0.5562	0.668	1.018
F15	3	8473.56	16-08	3882.94	0.6265	7889.00	1.29	0.458	0.931
F15	5	21929.10	0.0622	17811.00	1.492	51880.00	2.363	0.812	2.366
F16	2	1171.25	1e-08	740.37	0.001413	897.80	0.003917	0.632	0.767
F16	3	2091.88	1e-08	1490.07	0.03927	3739.84	0.06716	0.712	1.788
F16	5	6479.91	1e-08	4970.18	0.0417	16217.80	0.07463	0.767	2.503
F17	2	1397.16	1e-08	952.00	9.01e-07	1643.77	0.0001774	0.681	1.177
F17	3	2346.06	1e-08	1610.94	2.25e-07	5332.69	0.01492	0.687	2.273
F17	5	5595.91	1e-08	4393.66	5.07e-07	79964.00	0.01337	0.785	14.290
F18	2	1723.22	1e-08	2210.00	1.14e-01	7734.00	0.3329	1.282	4.488
F18	3	3904.69	1e-08	4807.50	1.48e-01	22480.50	0.1571	1.231	5.757
F18	5	9190.69	1e-08	22228.60	1.41e-02	N/A	0.1211	2.419	12110000
F19	2	3460.33	0.000982	1833.39	4.91e-03	3498.86	0.01915	0.530	1.011
F19	3	15316.14	0.00144	12702.00	2.47e-01	22671.20	0.2385	0.829	1.480
F19	5	71535.75	0.0579	77184.00	1.95e-01	N/A	0.405	1.079	6.995
F20	2	2330.81	1e-08	1571.65	2.31e-01	3022.40	0.2513	0.674	1.297
F20	3	7668.81	0.01	7379.73	5.70e-01	29992.70	0.8192	0.962	3.911
F20	5	37833.27	0.0449	50956.70	9.24e-01	156448.00	1.313	1.347	4.135
F21	2	590.19	1e-08	352.97	1e-08	459.75	1e-08	0.598	0.779
F21	3	1842.44	1e-08	1265.62	9.91e-02	2074.62	0.1948	0.687	1.126
F21	5	7702.25	1e-08	4443.60	6.98e-01	6536.50	0.9468	0.577	0.849
F22	2	434.44	1e-08	360.00	1e-08	431.63	1e-08	0.829	0.994
F22	3	1565.59	1e-08	1136.71	5.94e-02	1609.25	0.08648	0.726	1.028
F22	5	7979.37	0.0123	3573.12	2.99e-01	5198.74	1.097	0.448	0.652
F23	2	1664.84	1e-08	1528.80	1.04e-01	2652.00	0.2817	0.918	1.593
F23	3	8568.59	1e-08	3794.21	2.22e-01	7371.58	0.4927	0.443	0.860
F23	5	22807.11	0.0155	21265.10	2.67e-01	52277.30	0.5859	0.932	2.292
F24	2	4016.00	0.0348	4411.54	1.501	63342.00	2.09	1.098	15.772
F24	3	50193.29	2.44	30994.00	6.622	N/A	4.007	0.617	1.642
F24	5	1557063.00	19.7	N/A	5.931	N/A	7.855	0.301	0.399

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