

The Behavior of Deep Statistical Comparison Approach for Different Criteria of Comparing Distributions

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Abstract: Deep Statistical Comparison (DSC) is a recently proposed approach for the statistical comparison of meta-heuristic stochastic algorithms for single-objective optimization. The main contribution of the DSC is a ranking scheme, which is based on the whole distribution, instead of using only one statistic, such as average or median, which are commonly used. Contrary to common approach, the DSC gives more robust statistical results, which are not affected by outliers or misleading ranking scheme. The DSC ranking scheme uses a statistical test for comparing distributions in order to rank the algorithms. DSC was tested using the two-sample Kolmogorov-Smirnov (KS) test. However, distributions can be compared using different criteria, statistical tests. In this paper, we analyze the behavior of the DSC using two different criteria, the two-sample Kolmogorov-Smirnov (KS) test and the Anderson-Darling (AD) test. Experimental results from benchmark tests consisting of single-objective problems, show that both criteria behave similarly. However, when algorithms are compared on a single problem, it is better to use the AD test because it is more powerful and can better detect differences than the KS test when the distributions vary in shift only, in scale only, in symmetry only, or have the same mean and standard deviation but differ on the tail ends only. This influence is not emphasized when the approach is used for multiple-problem analysis.

1 INTRODUCTION

Over recent years, many meta-heuristic stochastic optimization algorithms have been developed. Performance analysis of a new algorithm compared with the state-of-the-art is a crucial task and one of the most common ways to compare their performance is to use statistical tests based on hypothesis testing (Lehmann et al., 1986). Making, such statistical comparisons, however, requires sufficient knowledge from the user, which includes knowing which conditions must be fulfilled so that the relevant and proper statistical test (e.g., parametric or nonparametric) can be applied (García et al., 2009).

The nature of stochastic optimization algorithms means that a set of independent runs must be executed on a single instance of a problem in order to get a relevant data set over which either average or median are typically calculated. Further, because the required conditions (normality and homoscedasticity) for the safe use of the parametric statistical tests are usually not satisfied when working with stochastic optimiza-

tion algorithms an appropriate nonparametric statistical test is required. Nonparametric statistical tests are based on a ranking scheme that is used to transform the data prior to analysis. The standard ranking scheme used by many of the nonparametric statistical tests ranks the algorithms for each problem (function) separately with the best performing being ranked number 1, the second best ranked number 2, and so on. In case of ties, average rankings are assigned. Unfortunately, using either the average or the median can negatively affect the outcome of the results of a statistical test (Eftimov et al., 2016). For example, averaging is sensitive to outliers, which should be handled appropriately since stochastic optimization algorithms can produce poor runs. For instance, let us suppose that two algorithms, A_1 and A_2 , are used to optimize the parabola problem, $y = x^2$, and the results after 10 runs are 0, 0, 0, 0, 0, 0, 0, 0, 10, and 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, for each algorithm, respectively. We see that the average of A_1 is 1, while the average of A_2 is 0.5. The standard ranking scheme will rank, A_1 and A_2 as 2 and 1, respectively. From this,

it follows that A_2 is better, but we can see that the algorithm A_1 has only one poor run (outlier), which affects the average. In the case when poor runs are not present, the average can be in some ε -neighborhood, which is defined as the set of all numbers whose distance from a number is less than some specified number ε , and the algorithms will obtain different rankings. In order to overcome this problem, medians are sometimes used because they are more robust to outliers. However, medians can be in some ε -neighborhood, and based on these the algorithms will obtain different rankings. The question is, therefore, how to define the ε -neighborhood for different test problems that have different ranges of obtained data e.g. 10^{-9} , 10^{-2} , 10^1 , etc. Let us suppose that two algorithms, A_1 and A_2 , are new algorithms used to optimize a given problem, and the results from 100 runs of both algorithms are distributed according to $N(0; 1)$. Figure 1(a), shows the probability density function of the two algorithms. In this case, the distributions are the same, the median values are in some ε -neighborhood, and because of this the algorithms should obtain the same ranking. Now let us suppose that two new algorithms, A_1 and A_2 , are used for the optimization of a given problem, and the results from 100 runs are distributed according to $N(0; 1)$ and $N(0; 2.5)$, respectively. Figure 1(b), shows the probability density function of the two algorithms. In this case, the distributions are not the same and the median values are in some ε -neighborhood, and because of this the algorithms should obtain different rankings. If this is a case, then the algorithms rankings are obtained either by the averages or the medians, so the algorithm which has smaller value for average or median is the better one. For these reasons, a novel approach was proposed, called *Deep Statistical Comparison (DSC)* (Eftimov et al., 2017), which removes the sensitivity of the simple statistics to the data and enables calculation of more robust statistics without fear of outliers influence or some errors inside ε -neighborhood.

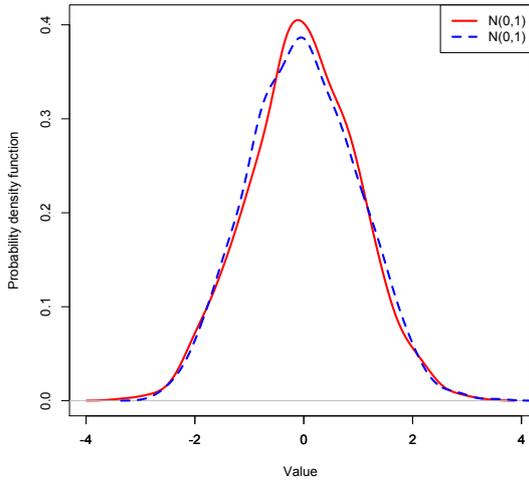
The remainder of the paper is organized as follows. Section II gives an overview of the related work, while Section III reintroduces the *DSC* ranking scheme used to compose a sample of results for each algorithm for multiple-problem analysis. Section IV presents statistical comparisons of stochastic optimization algorithms over multiple problems. This is then followed by a discussion of the results. In Section V, power analysis of *DSC* is presented when different statistical tests for comparing distributions are used in the ranking scheme. The conclusions of the paper are presented in Section VI.

2 RELATED WORK

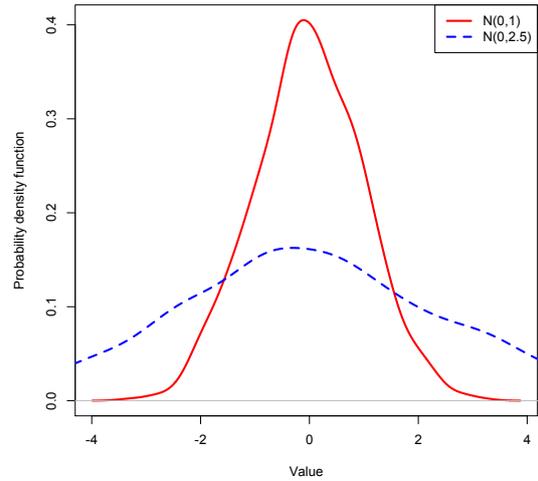
Statisticians (Gill, 1999) have shown that researchers have difficulties performing empirical studies and this could lead them into misinterpreting their results. Leven et al. (Levine et al., 2008) also state that the hypothesis testing is also frequently misunderstood and abused. To select an appropriate statistical test and to choose between a parametric and a nonparametric test (García et al., 2009), the first step is to check the assumptions of the parametric tests, in other words, the required conditions for the safe use of parametric tests.

Demšar (Demšar, 2006) theoretically and empirically examined several suitable statistical tests that can be used for comparing machine learning algorithms. Following a statistical tutorial on machine learning algorithms, Garcia et al. (García et al., 2009) presented a study on the use of nonparametric tests for analyzing the behaviour of evolutionary algorithms for optimization problems. They conducted their study in two ways: single-problem analysis and multiple-problem analysis. Single-problem analysis is a scenario when the data derives from multiple runs of the stochastic optimization algorithms on one problem. This scenario is common in stochastic optimization algorithms, since they are stochastic in nature, meaning we do not have any guaranty that the result will be the same for every run. Even the path leading to the final solution is often different. To test the quality of an algorithm, it is not sufficient to performed just a single run, but many runs are needed to draw a conclusion. The second scenario, the multiple-problem analysis, is the scenario when several stochastic optimization algorithms are compared over multiple problems. In this case, as in most published papers on this topic, the authors use the average of the results for each problem in order to produce results for each algorithm. We call this approach the “common approach” because it is the one most often used for making a statistical comparison of meta-heuristic stochastic optimization algorithms.

Recently, a novel approach for statistical comparison of stochastic optimization algorithms for multiple single-objective problems was proposed. The approach is known as *Deep Statistical Comparison (DSC)* (Eftimov et al., 2017). The term *Deep statistics* comes from the ranking scheme that is based on the whole distribution of multiple runs obtained on a single problem instead of using some simple statistics such as averages or medians. For this propose, the *DSC* was tested using one criteria for comparing distributions, the two-sample KS test. The approach consists of two steps, the first step uses a newly pro-



(a) Same distributions



(b) Different distributions

Figure 1: Probability density functions.

posed ranking scheme to obtain the data that will be used to make a statistical comparison. The second step is a standard omnibus statistical test that can be a parametric or a nonparametric one. *DSC* approach also allows users to calculate more robust statistics, while avoiding wrong conclusions due to either the presence of outliers or a ranking scheme that is used in some standard statistical tests.

3 DEEP STATISTICAL COMPARISON

In this paper, we analyze the behavior of *DSC* taking into account different criteria for comparing distributions used in the *DSC* ranking scheme. We did this to see if different criteria for comparing distributions influence the results from the *DSC*. The two most commonly used statistical tests for comparing distributions were selected, the two-sample *Kolmogorov-Smirnov (KS) test* and the two-sample *Anderson-Darling (AD) test* (Engmann and Cousineau, 2011), as different criteria for comparing distributions. To explain this, we start by reintroducing the *DSC*.

3.1 DSC Ranking Scheme

Let m and k be the number of algorithms and the number of problems that are used for statistical comparison, respectively, and n be the number of runs performed by each algorithm on the same problem.

Let X_i be a $n \times m$ matrix, where $i = 1, \dots, k$. The rows of this matrix correspond to the results obtained by multiple runs on the i -th problem, and the columns

correspond to the different algorithms that are used. The matrix element $X_i[j, l]$, where $j = 1, \dots, n$, and $l = 1, \dots, m$, corresponds to the result obtained by the j -th run on the i -th problem of the l -th algorithm.

The ranking scheme is based on the whole distribution, instead of ranking according to the averages or medians. The first step is to compare the probability distributions of multiple runs of each algorithm on each problem. For this purpose, a statistical test for comparing distributions should be used. Two most commonly used tests are the two-sample *KS test* and the two-sample *AD test*.

The two-sample *KS test* is a nonparametric test of the equality of continuous, one-dimensional probability distributions that can be used to compare two samples. The two-sample *KS test* is one of the most useful and general non-parametric methods for comparing two samples, as it is sensitive to differences in both location and shape of the empirical cumulative distribution functions of the two samples (e.g., a difference with respect to location/central tendency, dispersion/variability, skewness, and kurtosis) (Senger and Çelik, 2013).

The two-sample *AD test* has the same advantages mentioned for the *KS test*, sensibility to shape and scale of a distribution and its applicability to small samples. In addition, it has two extra advantages over the *KS test*. First, it is especially sensitive towards differences at the tails of distributions, and second it is better capable of detecting very small differences, even between large sample sizes. More about comparison of the two tests when shift, scale, and symmetry of distributions are varied independently for different sample sizes can be found in (Engmann and Cousineau, 2011).

Let α_T be the significance level used by the statistical test that is selected for comparing distributions. By using a statistical test for comparing distributions, $m \cdot (m - 1)/2$ pairwise comparisons between the algorithms are performed, and the results are organized into a $m \times m$ matrix, M_i as follows:

$$M_i[p, q] = \begin{cases} p_{\text{value}}, & p \neq q \\ 1, & p = q \end{cases}, \quad (1)$$

where p and q are different algorithms and $p, q = 1, \dots, m$.

Because multiple pairwise comparisons are made, this could lead to the *family-wise error rate (FWER)* (van der Laan et al., 2004), which is the probability of making one or more false discoveries, or type I errors, among all hypotheses when performing multiple hypotheses tests. In the cases when a few algorithms are compared the influence of multiple comparison in the FWER may not be very large, but as the number of compared algorithms increases, the FWER can increase dramatically. In order to counteract the problem of multiple comparisons, the *Bonferroni correction* (García et al., 2010) is used to correct the obtained p-values. The *Bonferroni correction* is based on the idea of testing u different hypotheses. One way of reducing the FWER is to test each individual hypothesis at a statistical significance level of $\frac{1}{u}$ times the desired maximum significance level. In our case the number of multiple pairwise comparisons, or the number of different hypotheses, is given as $C_m^2 = \binom{m}{2} = \frac{m \cdot (m-1)}{2}$.

The matrix M_i is reflexive. Also, it is symmetric because $M_i = M_i^T$, but the key point for the ranking scheme is to check the transitivity, since the ranking is made according to it. For this purpose, the matrix M'_i is introduced using the following equation:

$$M'_i[p, q] = \begin{cases} 1, & M_i[p, q] \geq \alpha_T / C_m^2 \\ 0, & M_i[p, q] < \alpha_T / C_m^2 \end{cases}. \quad (2)$$

The elements of the matrix, M'_i , are defined according to the p-values obtained by the statistical test used for comparing distributions corrected by the *Bonferroni correction*. For example if the element $M'_i[p, q]$ is 1 that means that the null hypothesis used in the statistical test for comparing distributions, which is the hypothesis that the two data samples obtained by the p -th and the q -th algorithm come from the same distribution, is not rejected. If the element $M'_i[p, q]$ is 0 that means that the null hypothesis is rejected, so the two data samples come from different distributions.

Before the ranking is performed, the matrix $M_i'^2$ is calculated to check the transitivity. If the M'_i has a 1

in each position for which $M_i'^2$ has non-zero element, the transitivity is satisfied, otherwise it is not.

If the transitivity is satisfied, the first step is to split the set of algorithms into w disjoint sets of algorithms Φ_f , $f = 1, \dots, w$, such as each algorithm belongs only in one of these sets. Each of these sets contains the indices of the algorithms that are used in the comparison for which the transitivity is satisfied. The cardinality of the union of these sets needs to be m , $\sum_{f=1}^w |\Phi_f| = m$. The next step is to define a $w \times 2$ matrix, W_i . The elements of this matrix are defined with the following equation

$$W_i[f, x] = \begin{cases} \text{mean}(\Phi_f \{h\}), & x = 1 \\ |\Phi_f|, & x = 2 \end{cases}, \quad (3)$$

where h is the number that is ceiled to the nearest integer of a number obtained by the uniform distribution of a random variable $\mathbf{Y} \sim U(1, |\Phi_f|)$. The algorithm from each set, whose average value will be used, can be chosen randomly because the data samples for all the algorithms that belong to the same set come from the same distribution. Then the rows of the matrix are reordered according to the first column sorted in ascending order. Let $Mean_i$ and C be a $w \times 1$ vectors that correspond to the first and the second column of the matrix W_i , respectively. Finally, the rankings to the sets, Φ_f , need to be assigned and organized into a $w \times 1$ vector $Rank_s$. For the set with lowest average value, $Mean_i[1]$, the ranking is defined as

$$Rank_s[1] = \sum_{r=1}^{C[1]} r / C[1]. \quad (4)$$

For remaining sets, the ranking is defined as

$$Rank_s[f] = \sum_{r=C[f-1]+1}^{C[f-1]+C[f]} r / C[f]. \quad (5)$$

After obtaining the rankings of the sets, each algorithm obtains its ranking according to the set to which it belongs by using the following equation:

$$Rank[i, l] = Rank_s[f], \quad l \in \Phi_f. \quad (6)$$

If the transitivity is not satisfied, the first step is to define two $1 \times m$ vectors, $Index_i$ and $Mean_i$, whose elements are the indices of the algorithms and the average values of the multiple runs for each algorithm. The both vectors are sorted in ascending order according to the average values. Then, the rankings of the sorted algorithms are organized into a $1 \times m$ vector, $Rank_s$, whose elements are defined with the following equation.

$$Rank_s[l] = \begin{cases} l, & \exists! Mean_i[l] \in Mean_i \\ \sum_{r=l-c+1}^l r / c, & otherwise \end{cases}, \quad (7)$$

where c is the number of elements from $Mean_i$ that have value $Mean_i[l]$. Finally, the algorithms obtain their rankings according to the rankings assigned to their average values by using the following equation:

$$Rank[i, Index_i[l]] = Rank_s[l]. \quad (8)$$

By using the ranking scheme for the algorithms on each problem, a $k \times m$ matrix, $Rank$, is defined. The i -th row of this matrix corresponds to the rankings of the algorithms obtained by the ranking scheme using the data samples from the i -th problem. Further, this matrix is used as input data for statistical comparison for multiple-problem analysis.

3.2 Selection of a Standard Omnibus Statistical Test

After ranking the algorithms, the next step is to choose an appropriate statistical test. The guidelines on which test to choose are given in (García et al., 2009). Using the new ranking scheme we transformed only the data that is available for further analysis, while everything else remain the same.

4 RESULTS AND DISCUSSION

4.1 Black-Box Benchmarking 2015 Test Functions

To evaluate the behavior of the DSC taking into account different statistical tests for comparing distributions, the results from the Black-Box Benchmarking 2015 (BBOB 2015) (Black Box Optimization Competition,) are used. BBOB 2015 is a competition that provides single-objective functions for benchmarking. From the competition 15 algorithms are used. The algorithms used are: BSif (Pošík and Baudiš, 2015), BSifeg (Pošík and Baudiš, 2015), BSqi (Pošík and Baudiš, 2015), BSrr (Pošík and Baudiš, 2015), CMA-CSA (Atamna, 2015), CMA-MSR (Atamna, 2015), CMA-TPA (Atamna, 2015), GP1-CMAES (Bajer et al., 2015), GP5-CMAES (Bajer et al., 2015), RAND-2xDefault (Brockhoff et al., 2015), RF1-CMAES (Bajer et al., 2015), RF5-CMAES (Bajer et al., 2015), Sif (Pošík and Baudiš, 2015), Sifeg (Pošík and Baudiš, 2015), and Srr (Pošík and Baudiš, 2015). For each of them results for 22 different noiseless test problems in 5 dimensionality (2, 3, 5, 10, and 20) are selected. More details about test problems can be found in (Hansen et al., 2010). Each algorithm provided data for 15 runs.

4.2 Experiments

Let Ψ be the set of the selected 15 stochastic optimization algorithms. For the experiments, the dimension is fixed to 10. A 100 random combinations of 3 distinct algorithms were generated (combinations without repetition) and used for statistical comparisons.

For each combination, the DSC ranking scheme is used to rank the algorithms for each problem separately. To compare the results using different criteria for comparing distributions, the DSC ranking scheme is used with the two-sample *KS test* and the two-sample *AD test*. Then, the results are also compared with the common approach, for which a sample of results for each algorithm is composed by averaging the data from multiple runs for each problem.

After obtaining the data for statistical comparison over multiple problems, the next step is to select an appropriate statistical test. In the case of the DSC the significance level for the ranking scheme is set to $\alpha_T = 0.05$, and the significance level for the statistical test is set to $\alpha = 0.05$.

Table 1 presents the p-values obtained for 6 combinations out of 100 generated, using the two versions of DSC approach, by using the *KS test* and the *AD test*, and the common approach with averages. The p_{value_F} corresponds to the p-values obtained by the *Friedman test*, which in our case is one of the appropriate omnibus statistical tests. From this table, we can see that the results for the first 2 combinations (1-2) differ. By using the common approach with averages, the null hypothesis is rejected, so there is significant statistical difference between the performance of the algorithms, while with both versions of DSC approach, the null hypothesis is not rejected, so we can assume that there is no significant statistical difference between the performance of the algorithms. The number of such combinations in our experiment is 5 out of 100. For the next 2 combinations (3-4), the common approach and both DSC versions give the same results, the p-values obtained are greater than 0.05, so the null hypothesis is not rejected, and we can assume that there is no significant statistical difference between the performance of the algorithms. The number of such combinations in our experiment is 9 out of 100, from which 2 randomly selected are presented in the table. For the last 2 combinations from this table (5-6), the results obtained are the same, the p-values are smaller than the significance level used, so the null hypothesis is rejected, and we can assume that there is a significant statistical difference between the performance of the algorithms. The number of such combinations in our experiment is 86 out of 100,

from which 2 are randomly selected and presented in the table.

In order to explain the difference that appears using the common approach and the two versions of the DSC, one example where the result from the common approach and both versions of the DSC differs is randomly selected and presented in detail. The first combination from the Table 1 is selected, which is a comparison between the algorithms GP5-CMAES, Sifeg, and BSif. For the analysis, the *Friedman test* was selected. In the case of the common approach, the null hypothesis is rejected, while when the DSC approach is used, the null hypothesis is not rejected. The rankings obtained by the *Friedman test* using the common approach with averages and both versions of the DSC ranking scheme are presented in Table 2. Comparing the rankings obtained by the common approach and the DSC ranking scheme, the difference between the rankings that appears by using them can be clearly observed. To explain the difference, separate problems are discussed in detail.

In Figure 2 the cumulative distributions (the step functions) and the average values (the horizontal lines) obtained from the multiple runs for different functions of the three algorithms are presented. In this figure details about the function, f_7 , are presented. The rankings obtained using the common approach with averages are 1.00, 2.00, and 3.00, and they are different because all of them have different averages. The rankings obtained using the two versions of DSC ranking scheme, by using *KS test* and *AD test*, are 1.00, 2.50, and 2.50. The DSC ranking scheme used the cumulative distributions in order to assign the rankings of the algorithms. From the figure, one may assume that there is no significant difference between the cumulative distributions of Sifeg and BSif, but they differ from the cumulative distribution of GP5-CMAES. This result is also obtained by using the two-sample *KS* and *AD test*. The p-values obtained for the pairs of algorithms are 0.00 (GP5-CMAES, Sifeg), 0.00 (GP5-CMAES, BSif), and 0.07 (Sifeg, BSif), by using the *KS test*, while the p-values obtained for the same pairs of algorithms by using the *AD test* are 0.00 (GP5-CMAES, Sifeg), 0.00 (GP5-CMAES, BSif), and 0.02 (Sifeg, BSif). Because multiple pairwise comparisons are made, these p-values are further corrected by using the *Bonferroni correction*. In this case, the transitivity of the matrix M_7' is satisfied, so the set of all algorithms is split into two disjoint sets {GP5-CMAES}, and {Sifeg, BSif}, and the rankings are defined using Equations 4 and 5.

In Figure 2(c) the results for the function, f_{21} , are presented. The rankings obtained using the common approach with averages are 1.00, 2.00, and 3.00,

and they are different because all of them have different average. The rankings obtained using the DSC ranking scheme by using the *KS test* and *AD test* are 2.00, 2.00, and 2.00. From the figure, it is not clear if there is a significant difference between the cumulative distributions of GP5-CMAES, Sifeg, and BSif. To check this, the two-sample *KS test* and *AD test* are used. The p-values obtained for the pairs of algorithms are 0.38 (GP5-CMAES, Sifeg), 0.07 (GP5-CMAES, BSif), and 0.38 (Sifeg, BSif), and 0.41 (GP5-CMAES, Sifeg), 0.02 (GP5-CMAES, BSif), and 0.29 (Sifeg, BSif), respectively. Because multiple pairwise comparisons are made, these p-values are further corrected using the *Bonferroni correction*. In this case, the transitivity of the matrix M_{21}' is satisfied, but the set of all algorithms is not split into disjoint sets because all algorithms belong to one set, {GP5-CMAES, Sifeg, BSif}.

In Figure 2(b), the results for the function, f_{18} , are presented. This example is interesting because both versions of the DSC ranking scheme that use different criteria for comparing distributions, the *KS test* and *AD test*, give different results. For the function f_{18} , the rankings obtained by the common approach are 1.00, 2.00, and 3.00. The rankings obtained by the DSC ranking scheme with *KS test* are 1.00, 2.50, and 2.50, while the *AD test* are 1.00, 2.00, and 3.00. So the two different criteria used by the DSC ranking scheme give different results. The p-values obtained by using the *KS test* for the pairs of algorithms are 0.00 (GP5-CMAES, Sifeg), 0.00 (GP5-CMAES, BSif), and 0.03 (Sifeg, BSif). Because multiple pairwise comparisons are made, these p-values are further corrected by using the *Bonferroni correction*. In this case, the transitivity of the matrix M_{18}' is satisfied, so the set of all algorithms is split into two disjoint sets {GP5-CMAES}, and {Sifeg, BSif}. The p-values obtained by using the *AD test* for the pairs of algorithms are 0.00 (GP5-CMAES, Sifeg), 0.00 (GP5-CMAES, BSif), and 0.01 (Sifeg, BSif). Because multiple pairwise comparisons are made, these p-values are further corrected using *Bonferroni correction*. In this case, the transitivity of the matrix M_{18}' is not satisfied, so the algorithms obtain their rankings according to their averages. So, the two different criteria give different results. This result is important when we compare algorithms on one problem (function), while it does not influence the result when we are performing multiple-problem analysis. Even more, when we are comparing algorithms on one problem, it is better to use *AD test* because it is more powerful and it can better detect differences than the *KS test* when the distributions vary in shift only, in scale only, in symmetry only, or that have the same mean and standard deviation but

Table 1: Statistical comparisons of 3 algorithms.

Algorithms		Common approach	DSC approach (KS)	DSC approach (AD)
		P_{valueF}	P_{valueF}	P_{valueF}
1	<i>GP5-CMAES, Sifeg, BSif</i>	*(.02)	(.42)	(.44)
2	<i>BSif, RFI-CMAES, Sifeg</i>	*(.00)	(.28)	(.33)
3	<i>BSifeg, RFI-CMAES, BSrr</i>	(.16)	(.28)	(.48)
4	<i>Sif, Bsrr, GPI-CMAES</i>	(.35)	(.77)	(.83)
5	<i>BSifeg, GPI-CMAES, CMA-CSA</i>	*(.00)	*(.00)	*(.00)
6	<i>BSrr, RAND-2xDefault, Srr</i>	*(.00)	*(.00)	*(.00)

* indicates that the null hypothesis is rejected, using $\alpha = 0.05$
 P_{valueF} corresponds to the p-value obtained by the *Friedman test*

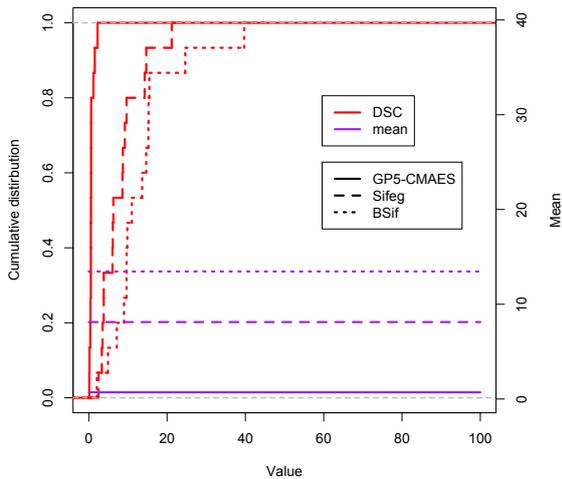
Table 2: Rankings for the algorithms $A_1=GP5-CMAES$, $A_2=Sifeg$, and $A_3=BSif$.

F	Common approach (<i>Friedman test</i>)			DSC ranking scheme (<i>KS test</i>)			DSC ranking scheme (<i>AD test</i>)		
	A_1	A_2	A_3	A_1	A_2	A_3	A_1	A_2	A_3
f_1	3.00	2.00	1.00	3.00	2.00	1.00	3.00	2.00	1.00
f_2	3.00	2.00	1.00	3.00	2.00	1.00	3.00	2.00	1.00
f_3	13.00	2.00	1.00	3.00	2.00	1.00	3.00	2.00	1.00
f_4	3.00	1.00	2.00	3.00	1.00	2.00	3.00	1.00	2.00
f_5	3.00	1.50	1.50	2.00	2.00	2.00	2.00	2.00	2.00
f_6	3.00	1.00	2.00	3.00	1.00	2.00	2.50	1.00	2.50
f_7	1.00	2.00	3.00	1.00	2.50	2.50	1.00	2.50	2.50
f_8	3.00	1.00	2.00	3.00	1.50	1.50	3.00	1.00	2.00
f_9	3.00	1.00	2.00	3.00	1.50	1.50	3.00	1.50	1.50
f_{10}	1.00	2.00	3.00	1.00	2.50	2.50	1.00	2.50	2.50
f_{11}	1.00	2.00	3.00	1.00	2.50	2.50	1.00	2.50	2.50
f_{12}	3.00	2.00	1.00	3.00	1.50	1.50	3.00	1.50	1.50
f_{13}	2.00	1.00	3.00	1.50	1.50	3.00	1.50	1.50	3.00
f_{14}	3.00	1.00	2.00	2.50	2.50	1.00	2.50	2.50	1.00
f_{15}	2.00	1.00	3.00	2.00	2.00	2.00	2.00	2.00	2.00
f_{16}	2.00	1.00	3.00	2.00	2.00	2.00	2.00	1.00	3.00
f_{17}	1.00	2.00	3.00	1.00	2.50	2.50	1.00	2.50	2.50
f_{18}	1.00	2.00	3.00	1.00	2.50	2.50	1.00	2.00	3.00
f_{19}	3.00	1.00	2.00	3.00	1.50	1.50	3.00	1.50	1.50
f_{20}	3.00	1.00	2.00	3.00	1.50	1.50	3.00	1.50	1.50
f_{21}	1.00	2.00	3.00	2.00	2.00	2.00	2.00	2.00	2.00
f_{22}	1.00	2.00	3.00	2.00	2.00	2.00	2.00	2.00	2.00

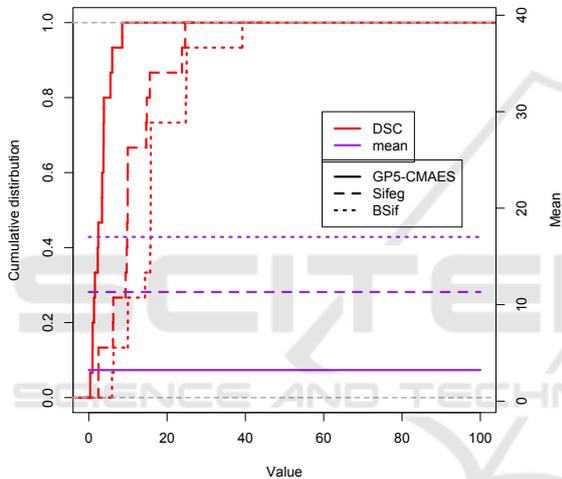
differ on the tail ends only (Engmann and Cousineau, 2011). Also, it requires less data than the *KS test* to reach sufficient statistical power. To see this difference, in Figure 3 the probability density functions for the data of each algorithm obtained on the function f_{18} are presented. From it, we can see that the *KS test* can not detect the small differences that exist between the probability distributions of the algorithms *Sifeg* and *BSif*, but the *AD test* can detect them.

If the null hypothesis is rejected by an omnibus

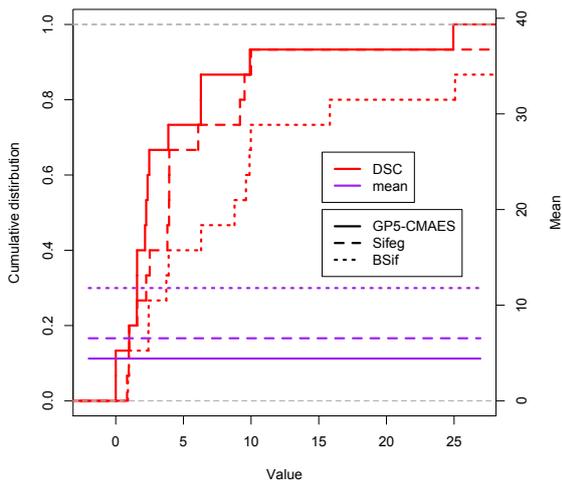
statistical test, an appropriate post-hoc test should be used (Demšar, 2006; García et al., 2009). Post-hoc tests are not a subject in this paper because we analyze the behavior of a ranking scheme that can be used for omnibus statistical tests. However, one of them is used to show differences that exist in the case of the common approach and the DSC approach. Since we are interested to compare all algorithms to each other, we decided to use the *Nemenyi test* (Nemenyi, 1963), which is an appropriate post-hoc test for the *Friedman*



(a) f_7



(b) f_{18}



(c) f_{21}

Figure 2: Cumulative distributions (the step functions) and mean values (the horizontal lines) for different functions of GP5-CMAES, Sifeg, and BSif.

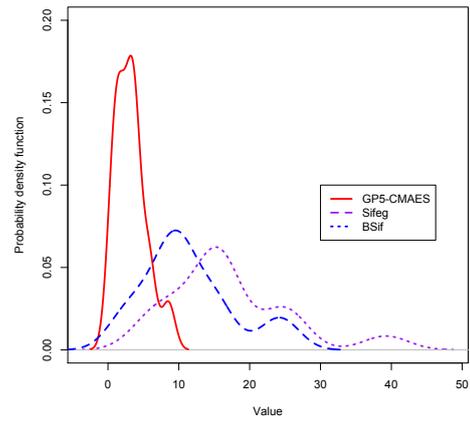
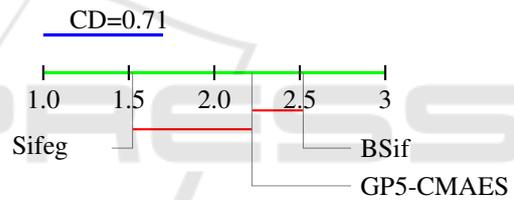
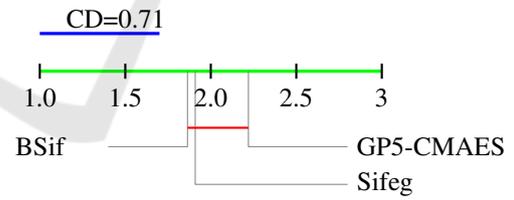


Figure 3: Probability density functions for f_{18} for the algorithms GP5-CMAES, Sifeg, and BSif.

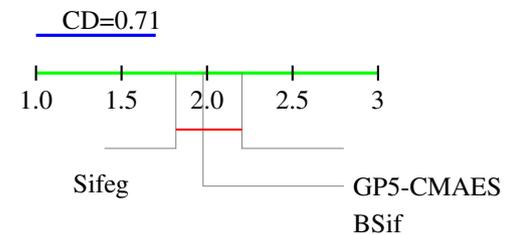
test. In Figure 4, the results obtained by the *Nemenyi test* are presented for the common approach with averages and both versions of DSC approach. When comparing all the algorithms against each other, the groups of algorithms that are not significantly different are connected together.



(a) Comparison of all algorithms against each other with the *Nemenyi test* (common approach using averages).



(b) Comparison of all algorithms against each other with the *Nemenyi test* (DSC approach with *KS test*).



(c) Comparison of all algorithms against each other with the *Nemenyi test* (DSC approach with *AD test*).

Figure 4: Visualization of post-hoc tests used for the algorithms GP5-CMAES, Sifeg, and BSif.

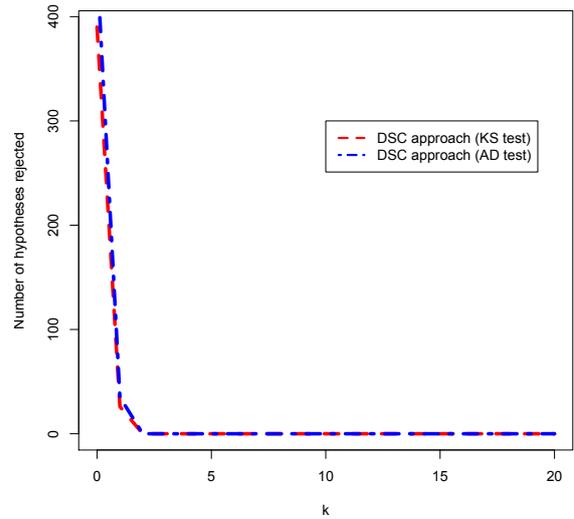
5 POWER ANALYSIS

The power of a statistical test is defined as the probability that the test will (correctly) reject the false null hypothesis. The comparison of the statistical power of the two-sample *KS test* and the two-sample *AD test* is presented in (Engmann and Cousineau, 2011). The comparison of the statistical power between the DSC and the common approach is presented in (Eftimov et al., 2017). Here we focus on power analysis between the two versions of the DSC approach that use different criteria for comparing distributions. For this purpose the power analysis is presented through an experimental analysis introduced in (Demšar, 2006). The experimental analysis of the power is made by a Monte-Carlo simulation. When we are comparing algorithms, samples of ten problems (functions) were randomly selected so that the probability for the problem i being chosen was proportional to $\frac{1}{1+e^{-kd_i}}$, where d_i is the difference between the rankings of the algorithms that are randomly chosen on that problem and k is the bias through which we can regulate the differences between the algorithms. Figure 5(a) represents the number of hypotheses rejected between the three algorithms considered with a significance level of 0.05 and Figure 5(b) represents their associated average p-values. From them, we can conclude that both versions of DSC approach behave similarly. The two-sample *AD test* has better power than the two-sample *KS test* (Engmann and Cousineau, 2011), but this influence is not emphasized in the case of the DSC approach.

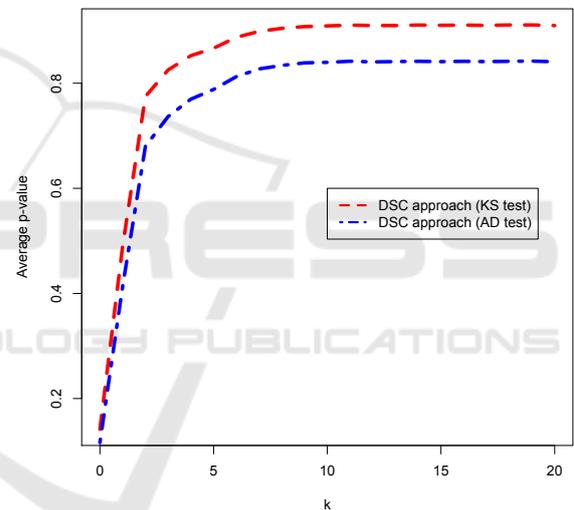
6 CONCLUSION

We analyze the behavior of the deep statistical comparison (DSC) approach taking into account different criteria for comparing distributions. The approach consists of two steps. In the first step, a new ranking scheme is used to obtain data for multiple-problem analysis. The ranking scheme is based on comparing distributions, instead of using simple statistics such as averages or medians. The second step is a standard omnibus statistical test, which uses the data obtained by the DSC ranking scheme as input data. By using the DSC, the wrong conclusions caused by the presence of outliers or ranking scheme used by some standard statistical tests can be avoided.

In this paper, different criteria for comparing distributions were used in the DSC ranking scheme, to see if there is a difference between the obtained results. We used two criteria for comparing distributions, the two-sample *Kolmogorov-Smirnov (KS)*



(a) Number of hypotheses rejected



(b) Average p-value

Figure 5: Power analysis for the combination CMA-MSR, Sifeg, and BSifeg.

test and the two-sample *Anderson-Darling (AD) test*. From the experimental results obtained over the algorithms presented in the BBOB 2015, we can conclude that both versions of DSC approach behave similarly. However, when we are comparing algorithms on one problem, it is better to use *AD test* because it is more powerful and it can better detect differences than the *KS test* when the distributions vary in shift only, in scale only, in symmetry only, or have the same mean and standard deviation but differ on the tail ends only (Engmann and Cousineau, 2011). Also, it requires less data than the *KS test* to reach sufficient statistical power. The two-sample *AD test* has better power than the two-sample *KS test*, but this influence

is not emphasized when the DSC approach is used for multiple-problem analysis.

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