

# A Hitchhiker's Guide to Statistical Tests for Assessing Randomized Algorithms in Software Engineering<sup>1</sup>

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## Abstract

Randomized algorithms are widely used to address many types of software engineering problems, especially in the area of software verification and validation with a strong emphasis on test automation. However, randomized algorithms are affected by chance, and so require the use of appropriate statistical tests to be properly analyzed in a sound manner. This paper features a systematic review regarding recent publications in 2009 and 2010 showing that, overall, empirical analyses involving randomized algorithms in software engineering tend to not properly account for the random nature of these algorithms. Many of the novel techniques presented clearly appear promising, but the lack of soundness in their empirical evaluations casts unfortunate doubts on their actual usefulness. In software engineering, though there are guidelines on how to carry out empirical analyses involving human subjects, those guidelines are not directly and fully applicable to randomized algorithms. Furthermore, many of the text books on statistical analysis are written from the viewpoints of social and natural sciences, which present different challenges from randomized algorithms. To address the questionable overall quality of the empirical analyses reported in the systematic review, this paper provides guidelines on how to carry out and properly analyze randomized algorithms applied to solve software engineering tasks, with a particular focus on software testing which is by far the most frequent application area of randomized algorithms within software engineering.

**Keyword:** Statistical difference, effect size, parametric test, non-parametric test, confidence interval, Bonferroni adjustment, systematic review, survey.

## 1 Introduction

Many problems in software engineering can be alleviated through automated support. For example, automated techniques exist to generate test cases that satisfy some desired coverage criteria on the system under test, such as for example branch [58] and path coverage [51]. Because often these problems are undecidable, deterministic algorithms that are able to provide optimal solutions in reasonable time do not exist. The use of heuristics, implemented as randomized algorithms [86], is hence necessary to address this type of problems.

At a high level, a randomized algorithm is an algorithm that has one or more of its components based on randomness. Therefore, running twice the same randomized algorithm on the same problem instance may yield different results. The most well-known example of randomized algorithm in software engineering is perhaps *random testing* [31, 13]. Techniques that use random testing are of course randomized, as for example DART [51] (which combines random testing with symbolic execution). Furthermore, there is a large body of work on the application of *search algorithms* in software engineering [57], as for example Genetic Algorithms. Since search algorithms are typically randomized and numerous software engineering problems can be

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<sup>1</sup>This paper is an extension of a conference paper [10] published in the International Conference on Software Engineering (ICSE), 2011.

34 addressed with search algorithms, randomized algorithms therefore play an increasingly important role. Appli-  
35 cations of search algorithms include software testing [81], requirement engineering [18], project planning and  
36 cost estimation [2], bug fixing [14], automated maintenance [84], service-oriented software engineering [22],  
37 compiler optimisation [26] and quality assessment [67].

38 A randomized algorithm may be strongly affected by chance. It may find an optimal solution in a very  
39 short time or may never converge towards an acceptable solution. Running a randomized algorithm twice on  
40 the same instance of a software engineering problem usually produces different results. Hence, researchers in  
41 software engineering that develop novel techniques based on randomized algorithms face the problem of how  
42 to properly evaluate the effectiveness of these techniques.

43 To analyze the cost and effectiveness of a randomized algorithm, it is important to study the *probability*  
44 *distribution* of its output and various performance metrics [86]. Though a practitioner might want to know  
45 what is the execution time of those algorithms *on average*, this might be misleading as randomized algorithms  
46 can yield very complex and high variance probability distributions.

47 The probability distribution of a randomized algorithm can be analyzed by running such an algorithm  
48 several times in an independent way, and then collecting appropriate data about its results and performance.  
49 For example, consider the case in which one wants to trigger failures by applying random testing (assuming  
50 that an automated oracle is provided) on a specific software system. As a way to assess its cost and effectiveness,  
51 test cases can be sampled at random until the first failure is detected. For example, in the first experiment, a  
52 failure might be detected after sampling 24 test cases. Assume the second run of the experiment (if a pseudo-  
53 random generator is employed, there would be the need to use a different seed for it) triggers the first failure  
54 when executing the second random test case. If in a third experiment the first failure is obtained after generating  
55 274 test cases, the *mean* value of these three experiments would be 100. Using such a mean to characterize  
56 the performance of random testing on a set of programs would clearly be misleading given the extent of its  
57 variation.

58 Since randomness might affect the reliability of conclusions when performing the empirical analysis of  
59 randomized algorithms, researchers hence face two problems: (1) how many experiments should be run to  
60 obtain reliable results, and (2) how to assess in a rigorous way whether such results are indeed reliable. The  
61 answer to these questions lies in the use of *statistical tests*, and there are many books on their various aspects  
62 (e.g., [99, 25, 71, 55, 119]). Notice that though statistical testing is used in most if not all scientific domains  
63 (e.g., medicine and behavioral sciences), each field has its own set of constraints to work with. Even within  
64 a field like software engineering the application context of statistical testing can vary significantly. When  
65 human resources and factors introduce randomness (e.g., [33, 63]) in the phenomena under study, the use of  
66 statistical tests is also required. But the constraints a researcher would work with are quite different from those  
67 of randomized algorithms, such as for example the size of data samples and the types of distributions.

68 Because of the widely varying situations across domains and the overwhelming number of statistical tests,  
69 each one with its own characteristics and assumptions, many practical guidelines have been provided targeting  
70 different scientific domains, such as biology [89] and medicine [64]. There are also guidelines for running  
71 experiment with human subjects in software engineering [120]. In this paper, the intent is to do the same for  
72 randomized algorithms in software engineering, with a particular focus on verification and validation, as they  
73 entail specific issues regarding the application of statistical testing.

74 To assess whether the results obtained with randomized algorithms are properly analyzed in software en-  
75 gineering research, and therefore whether precise guidelines are required, a systematic review was carried out.  
76 The analyses were limited to the years 2009 and 2010, as the goal was not to perform an exhaustive review  
77 of all research that was ever published but rather to obtain a recent, representative sample on which to draw  
78 conclusions about current practices. The focus was on research venues that deal with all aspects of software en-  
79 gineering, such as IEEE Transactions of Software Engineering (TSE), IEEE/ACM International Conference on  
80 Software Engineering (ICSE) and International Symposium on Search Based Software Engineering (SSBSE).  
81 The former two are meant to get an estimate of the extent to which randomized algorithms are used in software  
82 engineering. The latter, more specialized venue provides additional insight into the way randomized algorithms  
83 are assessed in software engineering. Furthermore, because randomized algorithms are more commonly used in  
84 software testing, the journal Software Testing, Verification and Reliability (STVR) was also taken into account.  
85 The review shows that, in many cases, statistical analyses are either missing, inadequate, or incomplete. For  
86 example, though journal guidelines in medicine require a mandatory use of standardized *effect size* measure-

87 ments [55] to quantify the effect of treatments, only one case was found in which a standardized effect size was  
88 used to measure the relative effectiveness of a randomized algorithm [96]. Even more surprising, in many of  
89 the surveyed empirical analyses, randomized algorithms were evaluated based on the results of only one run.  
90 Only few empirical studies reported the use of statistical analysis.

91 Given the results of this survey, it was necessary to devise *practical* guidelines for the use of statistical  
92 testing in assessing randomized algorithms in software engineering applications. Note that, though guidelines  
93 have been provided for other scientific domains [89, 64] and for other types of empirical analyses in software  
94 engineering [33, 63], they are not directly applicable and complete in the context of randomized algorithms. The  
95 objective of this paper is therefore to account for the specific properties of randomized algorithms in software  
96 engineering applications.

97 Notice that Ali *et al.* [3] have recently carried out a systematic review of search-based software testing  
98 which includes some limited guidelines on the use of statistical testing. This paper builds upon that work by: (1)  
99 analyzing software engineering as whole and not just software testing, (2) considering all types of randomized  
100 algorithms and not just search algorithms, and (3) giving precise, practical, and complete suggestions on many  
101 aspects related to statistical testing that were either not discussed or just briefly mentioned in the work of Ali *et*  
102 *al.* [3].

103 The main contributions of this paper can be summarized as follows:

- 104 • A systematic review is performed on the current state of practice of the use of statistical testing to analyze  
105 randomized algorithms in software engineering. The review shows that randomness is not properly taken  
106 into account in the research literature.
- 107 • A set of practical guidelines is provided on the use of statistical testing that are tailored to randomized  
108 algorithms in software engineering applications, with a particular focus on verification and validation  
109 (including testing), and the specific properties and constraints they entail.

110 The paper is organized as follows. Section 2 discusses a motivating example. The systematic review  
111 follows in Section 3. Section 4 presents the concept of statistical difference in the context of randomized  
112 algorithms. Section 5 compares two kinds of statistical tests and discusses their implications on randomized  
113 algorithms. The problem of censored data and how it applies to randomized algorithms is discussed in Section  
114 6. How to measure effect sizes and therefore the practical impact of randomized algorithms is presented in  
115 Section 7. Section 8 investigates the question of how many times randomized algorithms should be run. The  
116 problems associated with multiple tests are discussed in Section 9, whereas Section 10 deals with the choice  
117 of artifacts, which has usually a significant impact on results. Practical guidelines on how to use statistical  
118 tests are summarized in Section 11. The threats to validity associated with the work presented in this paper are  
119 discussed in Section 12. Finally, Section 13 concludes the paper.

## 120 2 Motivating Example

121 In this section, a motivating example is provided to show why the use of statistical tests is a necessity in the  
122 analyses of randomized algorithms in software engineering. Assume that two techniques  $\mathcal{A}$  and  $\mathcal{B}$  are used  
123 in a type of experiment in which the output is binary: either *pass* or *fail*. For example, in the context of  
124 software testing,  $\mathcal{A}$  and  $\mathcal{B}$  could be testing techniques (e.g., random testing [31, 13]), and the experiment would  
125 determine whether they trigger or not any failure given a limited testing budget. The technique with highest  
126 *success rate*, that is failure rate in the testing example, would be considered to be superior. Further assume  
127 that both techniques are run  $n$  times, and  $a$  represents the times  $\mathcal{A}$  was successful, whereas  $b$  is the number of  
128 successes for  $\mathcal{B}$ . The *estimated* success rates of these two techniques are defined as  $a/n$  and  $b/n$ , respectively.  
129 A related example in software testing (in which success rates are compared) that currently seems very common  
130 in industry (especially for online companies such as Google and Amazon) is “A/B testing”<sup>2</sup>.

131 Now, consider that such experiment is repeated  $n = 10$  times, and the results show that  $\mathcal{A}$  has a 70%  
132 estimated success rate, whereas  $\mathcal{B}$  has a 50% estimated success rate. Would it be safe to conclude that  $\mathcal{A}$  is  
133 better than  $\mathcal{B}$ ? Even if  $n = 10$  and the difference in estimated success rates is quite large (i.e., 20%), it would

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<sup>2</sup>[en.wikipedia.org/wiki/A/B\\_testing](http://en.wikipedia.org/wiki/A/B_testing), accessed October 2012.

134 actually be unsound to draw any conclusion about the respective performance of the two techniques. Because  
 135 this might not be intuitive, the exact mathematical reasoning is provided below to explain the above statement.

136 A series of repeated  $n$  experiments with binary outcome can be described as a *binomial distribution* [36],  
 137 where each experiment has probability  $p$  of success, and the mean value of the distribution (i.e., number of  
 138 successes) is  $pn$ . In the case of  $\mathcal{A}$ , one would have an estimated success rate  $p = a/n$  and an estimated number  
 139 of successes  $pn = a$ . The probability mass function of a binomial distribution  $B(n,p)$  with parameters  $n$  and  $p$   
 140 is:

$$P(B(n,p) = k) = \binom{n}{k} p^k (1-p)^{n-k} .$$

141  $P(B(n,p) = k)$  represents the probability that a binomial distribution  $B(n,p)$  would result in  $k$  successes.  
 142 Exactly  $k$  runs would be successful (probability  $p^k$ ) while the others  $n - k$  would fail (probability  $(1-p)^{n-k}$ ).  
 143 Since the order of successful experiments is not important, there are  $\binom{n}{k}$  possible orders. Using this probability  
 144 function, what is the probability that  $a$  equals the expected number of successes? Considering the example  
 145 provided in this section, having a technique with an *actual* 70% success rate, what is the probability of having  
 146 exactly 7 successes out of 10 experiments? This can be calculated with:

$$P(B(10,0.7) = 7) = \binom{10}{7} 0.7^7 (0.3)^3 = 0.26 .$$

147 This example shows that there is only a 26% chance to have exactly  $a = 7$  successes if the actual success  
 148 rate is 70%! This shows a potential misconception: expected values (e.g., successes) often have a relatively low  
 149 probability of occurrence. Similarly, the probability that both techniques have a number of successes equal to  
 150 their expected value would be even lower:

$$P(B(10,0.7) = 7) \times P(B(10,0.5) = 5) = 0.06 .$$

151 Reversely, even if one obtains  $a = 7$  and  $b = 5$ , what would be the probability that both techniques have an  
 152 equal actual success rate of 60%? We would have:

$$P(B(10,0.6) = 7) \times P(B(10,0.6) = 5) = 0.04 .$$

153 Though 0.04 seems a rather “low” probability, it is not much lower than 0.06, the probability of the observed  
 154 number of successes to be actually equal to their expected values. Therefore, one cannot really say that the  
 155 hypothesis of equal actual success rates (60%) is much more implausible than the one with 70% and 50%  
 156 actual success rates. But what about the case where the two techniques have exactly the same actual success  
 157 rate equal to 0.2? Or what about the cases in which  $\mathcal{B}$  would actually have a better actual success rate than  
 158  $\mathcal{A}$ ? What would be the probability for these situations to be true? Figure 1 shows all these probabilities, when  
 159  $a = 0.7n$  and  $b = 0.5n$ , for two different numbers of runs:  $n = 10$  and  $n = 100$ . For  $n = 10$ , there is a great  
 160 deal of variance in the probability distribution of success rates. In particular, the cases in which  $\mathcal{B}$  has a higher  
 161 actual success rate do not have a negligible probability. On the other hand, in the case of  $n = 100$ , the variance  
 162 has decreased significantly. This clearly shows the importance of using sufficiently large samples, an issue that  
 163 will be covered in more detail later in the paper.

164 In this example, with  $n = 100$ , the use of statistical tests (e.g., Fisher Exact test) would yield strong  
 165 evidence to conclude that  $\mathcal{A}$  is better than  $\mathcal{B}$ . At an intuitive level, a statistical test would estimate the probability  
 166 of mistakenly drawing the conclusion that  $\mathcal{A}$  is better than  $\mathcal{B}$ , under the form of a so-called  $p$ -value, as further  
 167 discussed later in the paper. The resulting  $p$ -value would be quite small for  $n = 100$  (i.e., 0.005), whereas for  
 168  $n = 10$  it would far much larger (i.e. 0.649), thus confirming and quantifying what is graphically visible in  
 169 Figure 1. So even for what might appear to be large values of  $n$ , the capability to draw reliable conclusions  
 170 could still be weak. Though some readers might find the above example rather basic, the fact of the matter is  
 171 that many papers reporting on randomized algorithms overlook the principles and issues illustrated above.

### 172 3 Systematic Review

173 Systematic reviews are used to gather, in an unbiased and comprehensive way, published research on a specific  
 174 subject and analyze it [65]. Systematic reviews are a useful tool to assess general trends in published research,

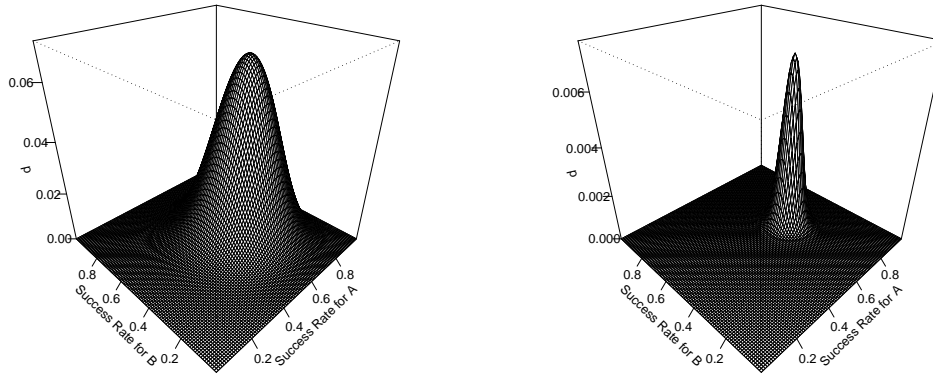


Figure 1: Probabilities to obtain  $a = 0.7n$  and  $b = 0.5n$  when  $n = 10$  (left) and  $n = 100$  (right) for different success rates of the algorithms  $\mathcal{A}$  and  $\mathcal{B}$ .

175 and they are becoming increasingly common in software engineering [70, 33, 63, 3].

176 The systematic review reported in this paper aims at analyzing: (RQ1) how often randomized algorithms  
 177 are used in software engineering, (RQ2) how many runs were used to collect data, and (RQ3) which types of  
 178 statistical analyses were used for data analysis.

179 To answer RQ1, two of the main venues that deal with all aspects of software engineering were selected:  
 180 IEEE Transactions of Software Engineering (TSE) and IEEE/ACM International Conference on Software En-  
 181 gineering (ICSE). The International Symposium on Search-Based Software Engineering (SSBSE) was also  
 182 considered, which is a specialized venue devoted to the application of search algorithms in software engi-  
 183 neering. Furthermore, because many of the applications of randomized algorithms are in software testing, the  
 184 journal Software Testing, Verification and Reliability (STVR) was included as well. Because the goal of this  
 185 paper is not to perform an exhaustive survey of published works, but rather to get an up-to-date snapshot of  
 186 current practices regarding the application of randomized algorithms in software engineering research, only  
 187 2009 and 2010 publications were included.

188 Only full length research papers were retained and, as a result, 77 papers at ICSE and 11 at SSBSE were  
 189 excluded. A total of 246 papers were considered: 96 in TSE, 104 in ICSE, 23 in SSBSE and 23 in STVR. These  
 190 papers were manually checked to verify whether they made use of randomized algorithms, thus leading to a  
 191 total of 54 papers. The number of analyzed papers is in line with other systematic reviews (e.g., in the work of  
 192 Ali *et al.* [3] a total of 68 papers were analyzed). For example, in their systematic review on systematic reviews  
 193 in software engineering, Kitchenham *et al.* [70] show that 11 out 20 systematic reviews involved less than 54  
 194 publications. Table 1 summarizes the details of the systematic review divided by venue and year.

195 Notice that papers were excluded if it was not clear whether randomized algorithms were used. For exam-  
 196 ple, the techniques described in the work of Hsu and Orso [60] and the work of Thum *et al.* [112] use external  
 197 SAT solvers, and those might be based on randomized algorithms, though it was not possible to tell with cer-  
 198 tainty. Furthermore, papers that involve *machine learning* algorithms that are randomized were not considered  
 199 since they require different types of analysis [85]. On the other hand, if a paper focused on presenting a deter-  
 200 ministic, novel technique, then it was included when randomized algorithms were used for comparison purposes  
 201 (e.g., fuzz testing [43]). Table 2 (for the year 2009) and Table 3 (for the year 2010) summarize the results of  
 202 this systematic review for the final selection of 54 papers. The first clearly visible result is that randomized  
 203 algorithms are widely used in software engineering (RQ1): they were found in 15% of the regular articles in  
 204 TSE and ICSE, which are general-purpose and representative software engineering venues. More specifically,  
 205 72% of all the papers (i.e., 39 out of 54) are on verification and validation (V&V).

206 To answer RQ2, the data in Table 2 and Table 3 shows the number of times a technique was run to collect  
 207 data regarding its performance on each artifact in the case study. Only 27 cases out of 54 show at least 10 runs.

Table 1: Number of publications grouped by venue, year and type.

Venue	Year	All	Regular	Randomized Algorithms
TSE	2009	48	48	3
	2010	48	48	12
ICSE	2009	70	50	4
	2010	111	54	10
SSBSE	2009	17	9	9
	2010	17	14	11
STVR	2009	12	12	4
	2010	11	11	1
<b>Total</b>		334	246	54

208 In many cases, data are collected from only one run of the randomized algorithms. Furthermore, notice that  
 209 the case in which randomized algorithms are evaluated based on *only one run per case study artifact* is quite  
 210 common in the literature. Even very influential papers, such as DART [51], feature this problem which poses  
 211 serious threats to the validity of their reported empirical analyses.

212 In the literature, there are empirical analyses in which randomized algorithms are run only once per case  
 213 study artifact, but a large number of artifacts were generated at random (e.g., [90, 118]). The validity of such  
 214 empirical analyses depends on the representativeness of instances created with the random generator. At any  
 215 rate, the choice of a case study that is statistically appropriate, and its relations to the required number of runs  
 216 for evaluating a randomized algorithm, needs careful consideration and will be discussed in more detail in  
 217 Section 10.

218 Regarding RQ3, only 19 out of 54 articles include empirical analyses supported by some kind of statistical  
 219 testing. More specifically, those are *t*-tests, Welch and U-tests when algorithms are compared in a pairwise  
 220 fashion, whereas ANOVA and Kruskal-Wallis are used for multiple comparisons. Furthermore, in some cases  
 221 linear regression is employed to build prediction models from a set of algorithm runs. However, in only one  
 222 article [96] standardized *effect size* measures (see Section 7) are reported to quantify the relative effectiveness  
 223 of algorithms.

224 Results in Table 2 and 3 clearly show that, when randomized algorithms are employed, empirical analyses in  
 225 software engineering do not properly account for their random nature. Many of the novel proposed techniques  
 226 may indeed be useful, but the results in Table 2 and 3 cast serious doubts on the validity of most existing results.

227 Notice that some of empirical analyses in Table 2 and 3 do not use statistical tests since they do not perform  
 228 any comparison of the technique they propose with alternatives. For example, in the award winning paper at  
 229 ICSE 2009, a search algorithm (i.e., Genetic Programming) was used and was run 100 times on each artifact  
 230 in the case study [117]. However this algorithm was not compared against simpler alternatives or even random  
 231 search (e.g., successful applications of automated bug fixing on real-world software can be traced back at least  
 232 down to the work of Griesmayer *et al.* [54]). When looking more closely at the reported results in order to assess  
 233 the implications of such lack of comparison, one would see that the total number of fitness evaluations was 400  
 234 (a population size of 40 individuals that is evolved for 10 generations). This sounds like a very low number (for  
 235 example, for test data generation in branch coverage, it is common to see 100,000 fitness evaluations for *each*  
 236 branch [58]) and one can therefore conclude that there is very limited search taking place. This implies that a  
 237 random search might have yielded similar results, and this would have warranted a comparison with random  
 238 search. This is directly confirmed in the reported results in the work of Weimer *et al.* [117], in which in half  
 239 of the subject artifacts in the case study, the average number of fitness evaluations per run is at most 41, thus  
 240 implying that, on average, appropriate patches are found in the random initialization of the first population  
 241 before the actual evolutionary search even starts.

242 As the search operators were tailored to specific types of bugs, then the choice of the case study and its  
 243 representativeness play a major role in assessing the validity of an empirical study (more details in Section 10).  
 244 Therefore, as discussed by Ali *et al.* [3], a search algorithm should always be compared against at least random  
 245 search in order to check that success is not due to the search problem (or case study) being easy. Notice,  
 246 however, that the previous work on automated bug fixing does not seem to feature comparisons neither (e.g.,

Table 2: Results of systematic review for the year 2009.

Reference	Venue	V&V	Repetitions	Statistical Tests
[1]	TSE	yes	1/5	U-test
[80]	TSE	yes	1	None
[90]	TSE	no	1	None
[83]	ICSE	no	100	$t$ -test, U-test
[117]	ICSE	yes	100	None
[43]	ICSE	yes	1	None
[68]	ICSE	yes	1	None
[7]	SSBSE	yes	1000	Linear regression
[48]	SSBSE	yes	30/500	None
[32]	SSBSE	no	100	U-test
[46]	SSBSE	yes	50	None
[72]	SSBSE	yes	10	Linear regression
[66]	SSBSE	yes	10	None
[79]	SSBSE	yes	1	None
[69]	SSBSE	no	1	None
[106]	SSBSE	no	1	None
[21]	STVR	yes	1/100	None
[95]	STVR	yes	1	None
[104]	STVR	yes	1	None
[61]	STVR	yes	Undefined	None

see [111, 110, 54, 14]). The work of Weimer *et al.* [117] was discussed only because it was among the sampled papers in the systematic review, and it is a good example to point out the importance of comparisons.

Since comparisons with simpler alternatives (at a very minimum random search) is a necessity when one proposes a novel randomized algorithm or addresses a new software engineering problem [3], statistical testing should be part of all publications reporting such empirical studies. In this paper, specific guidelines are provided on how to use statistical tests to support comparisons among randomized algorithms. One might argue that, depending on the addressed problem and the aimed contribution, there might be cases when comparisons with alternatives are either not possible or unnecessary, thus removing the need for statistical testing. However, such cases should be rare and in any case not nearly as common as what can be observed in the systematic review.

## 4 Statistical Difference

When a novel randomized algorithm  $\mathcal{A}$  is developed to address a software engineering problem, it is common practice to compare it against existing techniques, in particular simpler alternatives. For simplicity, consider the case in which just one alternative randomized algorithm (called  $\mathcal{B}$ ) is used in the comparisons. For example,  $\mathcal{B}$  can be random testing, and  $\mathcal{A}$  can be a search algorithm such as Genetic Algorithms or an hybrid technique that combines symbolic execution with random testing (e.g., DART [51]).

To compare  $\mathcal{A}$  versus  $\mathcal{B}$ , one first needs to decide which criteria are used in the comparisons. Many different measures ( $M$ ), either attempting to capture the effectiveness or the cost of algorithms, can be selected depending on the problem at hand and contextual assumptions, e.g., source code coverage, execution time. Depending on the selected choice, one may want to either minimize or maximize  $M$ , for example maximize coverage and minimize execution time.

To enable statistical analysis, one should run both  $\mathcal{A}$  and  $\mathcal{B}$  a large enough number ( $n$ ) of times, in an independent way, to collect information on the probability distribution of  $M$  for each algorithm. A *statistical test* should then be run to assess whether there is enough empirical evidence to claim, with a high level of confidence, that there is a difference between the two algorithms (e.g.,  $\mathcal{A}$  is better than  $\mathcal{B}$ ). A *null hypothesis*  $H_0$  is typically defined to state that there is no difference between  $\mathcal{A}$  and  $\mathcal{B}$ . In such a case, a statistical test aims

Table 3: Results of systematic review for the year 2010.

Reference	Venue	V&V	Repetitions	Statistical Tests
[45]	TSE	yes	1000	None
[125]	TSE	yes	100	<i>t</i> -test
[58]	TSE	yes	60	U-test
[96]	TSE	yes	32	U-test, $\hat{A}_{12}$
[30]	TSE	yes	30	Kruskal-Wallis, undefined pairwise
[109]	TSE	no	20	None
[20]	TSE	no	10	U-test, <i>t</i> -test, ANOVA
[34]	TSE	no	3	U-test
[6]	TSE	yes	1	None
[16]	TSE	yes	1	None
[19]	TSE	yes	1	None
[118]	TSE	no	1	None
[74]	ICSE	yes	100	None
[126]	ICSE	yes	50	None
[50]	ICSE	yes	5	None
[87]	ICSE	yes	5	None
[42]	ICSE	yes	1	None
[56]	ICSE	yes	1	None
[62]	ICSE	no	1	None
[123]	ICSE	yes	1	None
[92]	ICSE	yes	1	None
[103]	ICSE	no	1	None
[28]	SSBSE	yes	100	<i>t</i> -test
[29]	SSBSE	no	100	None
[78]	SSBSE	no	50	<i>t</i> -test
[82]	SSBSE	yes	50	U-test
[122]	SSBSE	yes	30	U-test
[124]	SSBSE	yes	30	<i>t</i> -test
[75]	SSBSE	yes	30	Welch
[115]	SSBSE	no	30	ANOVA
[17]	SSBSE	yes	3/5	None
[77]	SSBSE	yes	3	None
[127]	SSBSE	no	1	None
[128]	STVR	yes	24/480	Linear regression



272 to verify whether one should reject the null hypothesis  $H_0$ . However, what aspect of the probability distribution  
273 of  $M$  is being compared depends on the used statistical test. For example, a  $t$ -test compares the mean values of  
274 two distributions whereas others tests focus on the median or proportions, as discussed in Section 5.

275 There are two possible types of error when performing statistical testing: (I) one rejects the null hypothesis  
276 when it is true (i.e., claiming that there is a difference between two algorithms when actually there is none),  
277 and (II)  $H_0$  is accepted when it is false (there is a difference but the researcher claims the two algorithms to be  
278 equivalent). The  $p$ -value of a statistical test denotes the probability of a Type I error. The *significant level*  $\alpha$   
279 of a test is the highest  $p$ -value one accepts for rejecting  $H_0$ . A typical value, inherited from widespread practice  
280 in natural and social sciences, is  $\alpha = 0.05$ .

281 Notice that the two types of error are conflicting; minimizing the probability of one of them necessarily  
282 tends to increase the probability of the other. But traditionally there is more emphasis on not committing a  
283 Type I error, a practice inherited from natural sciences where the goal is often to establish the existence of a  
284 natural phenomenon in a conservative manner. In this context, one would only conclude that an algorithm  $\mathcal{A}$   
285 is better than  $\mathcal{B}$  when the probability of a Type I error is below  $\alpha$ . The price to pay for a small  $\alpha$  value is  
286 that, when the data sample is small, the probability of a Type II error can be high. The concept of statistical  
287 *power* [25] refers to the probability of rejecting  $H_0$  when it is false (i.e., the probability of claiming statistical  
288 difference when there is actually a difference).

289 Getting back to the comparison of techniques  $\mathcal{A}$  and  $\mathcal{B}$ , assume one obtains a  $p$ -value equal to 0.06. Even  
290 if one technique seems significantly better than the other in terms of effect size (Section 7), the researcher  
291 would then conclude that there is no difference when using the traditional  $\alpha = 0.05$  threshold. In software  
292 engineering, or in the context of *decision-making* in general, this type of reasoning can be counter-productive.  
293 The tradition of using  $\alpha = 0.05$ , discussed by Cowles [27], has been established in the early part of the last  
294 century, in the context of natural sciences, and is still applied by many across scientific fields. It has, however,  
295 an increasing number of detractors [52, 53] who believe that such thresholds are arbitrary, and that researchers  
296 should simply report  $p$ -values and let the readers decide in context what risks they are willing to take in their  
297 decision-making process.

298 When there is the need to make a choice between techniques  $\mathcal{A}$  and  $\mathcal{B}$ , an engineer would like to use the  
299 technique that is more likely to outperform the other. If one is currently using  $\mathcal{B}$ , and a new technique  $\mathcal{A}$   
300 seems to show better results, then a high level of confidence (i.e., a low  $p$ -value) might be required before  
301 opting for the “cost” (e.g., buying licenses and training) of switching from  $\mathcal{B}$  to  $\mathcal{A}$ . On the other hand, if  
302 the “cost” of applying the two techniques is similar, then whether one gets a  $p$ -value lower than  $\alpha$  bears little  
303 consequence from a practical standpoint, as in the end an alternative *must* be selected, for example to test a  
304 system. However, as it will be shown in Section 8, obtaining  $p$ -values lower than  $\alpha = 0.05$  should not be a  
305 problem when experimenting with randomized algorithms. The focus of such experiments should rather be  
306 on whether a given technique brings any practically significant advantage, usually measured in terms of an  
307 estimated effect size and its confidence interval, an important concept addressed in Section 7.

308 In practice, the selection of an algorithm would depend on the  $p$ -value of effectiveness comparisons, the  
309 effectiveness effect size, and the cost difference among algorithms (e.g., in terms of user-provided inputs or  
310 execution time). Given a context-specific decision model, the reader, using such information, could then decide  
311 which technique is more likely to maximize benefits and minimize risk. In the simplest case where compared  
312 techniques would have comparable costs, one would simply select the technique with the highest effectiveness  
313 regardless of the  $p$ -values of comparisons, even if as a result there is a non-negligible probability that it will  
314 bring no particular advantage.

315 When one has to carry out a statistical test, one must choose between *one-tailed* and a *two-tailed* test.  
316 Briefly, in a two-tailed test, the researcher would reject  $H_0$  if the performance of  $\mathcal{A}$  and  $\mathcal{B}$  are different regardless  
317 of which one is the best. On the other hand, in a one-tailed test, the researcher is making assumptions about  
318 the relative performance of the algorithms. For example, one could expect that a new sophisticated algorithm  
319  $\mathcal{A}$  is better than a naive algorithm  $\mathcal{B}$  used in the literature. In such a case, one would detect a statistically  
320 significant difference when  $\mathcal{A}$  is indeed better than  $\mathcal{B}$ , but ignoring the “unlikely” case of  $\mathcal{B}$  being better than  
321  $\mathcal{A}$ . An historical example in the literature of statistics is the test to check whether there is the right percent of  
322 gold (carats) in coins. One could expect that a dishonest coiner might produce coins with lower percent of gold  
323 than declared, and so a one-tailed test would be used rather than a two-tailed. Such a test could be used if one  
324 wants to verify whether the coiner is actually dishonest, whereas giving more gold than declared would be very

325 unlikely. Using a one-tailed test has the advantage, compared to a two-tailed test, that the resulting  $p$ -value is  
326 lower (so it is easier to detect statistically significant differences).

327 Are there cases in which a one-tailed test could be advisable in the analysis of randomized algorithms in  
328 software engineering? As a rule of thumb, the authors of this paper believe this is not the case: two-tailed tests  
329 should be used. One should use a one-tailed test only if he has strong arguments to support such a decision. In  
330 contrast to empirical analyses in software engineering involving human subjects, most of the time one cannot  
331 make any assumption on the relative performance of randomized algorithms. Even naive testing techniques  
332 such as random testing can fare better than more sophisticated techniques on some classes of problems (e.g.,  
333 [105, 9]). The reason is that sophisticated novel techniques might incur extra computational overhead compared  
334 to simpler alternatives, and the magnitude of this overhead might not only be very high but also difficult to  
335 determine before running the experiments. Furthermore, search algorithms do exhibit complex behavior, which  
336 is dependent on the properties of the search landscape of the addressed problem. It is not uncommon for a  
337 novel testing technique to be better on certain types of software and worse on others. For example, an empirical  
338 analysis in software testing in which this phenomenon is visible with statistical confidence can be found in  
339 the work of Fraser and Arcuri [37]. In that paper, a novel technique for test data generation of object-oriented  
340 software was compared against the state of the art. Out of a total of 727 Java classes, the novel technique  
341 gave better results in 357 cases, but worse on 81 (on the remaining 289 classes there was no difference). In  
342 summary, if one wants to lower the  $p$ -values, it is recommended to have a large number of runs (see Section 8)  
343 when possible rather than using an arguable one-tailed test.

344 Assume that a researcher runs  $n$  experiments and does not obtain significant results. It might be then  
345 tempting to run an additional  $k$  experiments, and base the statistical analyses on those  $n + k$  runs, in the hope  
346 of getting significant results as a result of increased statistical power. However, in this case, the  $k$  runs are not  
347 independent, as the choice of running them depended on the outcome of the first  $n$  runs. As a result, the real  
348  $p$ -value ends up being higher than what is estimated by statistical testing. This problem and related solutions  
349 are referred to in the literature as “sequence statistical testing” or “sequential analysis”, and have been applied  
350 in numerous fields such as repeated clinical trials [108]. In any case, if one wants to run  $k$  more experiments  
351 after analyzing the first  $n$ , it is important to always state it explicitly, as otherwise the reader would be misled  
352 when interpreting the obtained results.

## 353 5 Parametric vs Non-Parametric Tests

354 In the research context of this paper, the two most used statistical tests are the  $t$ -test and the Mann-Whitney  
355 U-test. These tests are in general used to compare two independent data samples (e.g., the results of running  $n$   
356 times algorithm  $\mathcal{A}$  compared to  $\mathcal{B}$ ). The  $t$ -test is *parametric*, whereas the U-test is *non-parametric*.

357 A parametric test makes assumptions on the underlying distribution of the data. For example, the  $t$ -test as-  
358 sumes normality and equal variance of the two data samples. A non-parametric test makes no assumption about  
359 the distribution of the data. *Why* is there the need for two different types of statistical tests? A simple answer is  
360 that, in general, non-parametric tests are less powerful than parametric ones when the latter’s assumptions are  
361 fulfilled. When, due to cost or time constraints, only small data samples can be collected, one would like to use  
362 the most powerful test available if its assumptions are satisfied.

363 There is a large body of work regarding which of the two types of tests should be used [35]. The assumptions  
364 of the  $t$ -test are in general not met. Considering that the variance of the two data samples is most of the time  
365 different, a Welch test should be used instead of a  $t$ -test. But the problem of the normality assumption remains.

366 An approach would be to use a statistical test to assess whether the data is normal, and, if the test is  
367 successful, then use a Welch test. This approach increases the probability of Type I error and is often not  
368 necessary. In fact, the Central Limit theorem tells that, for large samples, the  $t$ -test and Welch test are robust  
369 even when there is strong departure from a normal distribution [99, 102]. But in general one cannot know how  
370 many data points ( $n$ ) he needs to reach reliable results. A rule of thumb is to have at least  $n = 30$  for each data  
371 sample [99].

372 There are three main problems with such an approach: (1) if one needs to have a large  $n$  for handling  
373 departures from normality, then it might be advisable to use a non-parametric test since, for a large  $n$ , it is  
374 likely to be powerful enough; (2) the rule of thumb  $n = 30$  stems from analyses in behavioral science and there  
375 is no supporting evidence of its efficacy for randomized algorithms in software engineering; (3) the Central

376 Limit theorem has its own set of assumptions, which are too often ignored. Points (2) and (3) will be now  
377 discussed in more details by accounting for the specific properties of the application of randomized algorithms  
378 in software engineering, with an emphasis on software testing.

## 379 5.1 Violation of Assumptions

380 Parametric tests make assumptions on the probability distributions of the analyzed data sets, but “The assump-  
381 tions of most mathematical models are always false to a greater or lesser extent” [49]. Consider the following  
382 software testing example. A technique is used to find a test case for a specific testing target (e.g., a test case  
383 that triggers a failure or covers a particular branch/path), and then a researcher evaluates how many test cases  
384  $X_i$  the technique requires to sample and evaluate before covering that target. This experiment can be repeated  
385  $n$  times, yielding  $n$  observations  $\{X_1, \dots, X_n\}$  to study the probability distribution of the random variable  $X$ .  
386 Ideally, one would like a testing technique that minimizes  $X$ .

387 Since using the  $t$ -test assumes normality in the distribution  $X$ , are there cases for which it can be used to  
388 compare distributions of  $X$  resulting from different test techniques? The answer to this question is *never*. First,  
389 a normal distribution is continuous, whereas the number of sampled test cases  $X$  would be discrete. Second,  
390 the density function of the normal distribution is always positive for any value, whereas  $X$  would have zero  
391 probability for negative values. At any rate, asking whether a data set follows a normal distribution is not the  
392 right question [49]. A more significant question is what are the effects of departures from the assumptions on  
393 the validity of the tests. For example, a  $t$ -test returns a  $p$ -value that quantifies the probability of Type I error.  
394 The more the data departs from normality and equal variance, the more the resulting  $p$ -value will deviate from  
395 the true probability of Type I error.

396 Glass *et al.* [49] showed that in many cases the departures from the assumptions do not have serious con-  
397 sequences, particularly for data sets with not too high kurtosis (roughly, the kurtosis is a measure of infrequent  
398 extreme deviations). However, such empirical analyses reported and surveyed by Glass *et al.* [49] are based on  
399 social and natural sciences. For example, Glass *et al.* [49] wrote:

400 “Empirical estimates of skewness and kurtosis are scattered across the statistical literature. Kendall and  
401 Stuart (1963, p. 57) reported the frequency distribution of age at marriage for over 300,000 Australians;  
402 the skewness and kurtosis were 1.96 and 8.33, respectively. The distribution of heights of 8,585 English  
403 males (see Glass & Stanley, 1970, p. 103) had skewness and kurtosis of -0.08 and 3.15, respectively”.

404 Data sets for age at marriage and heights have known bounds (e.g., according to Wikipedia, the tallest  
405 man in world was 2.72 meters, whereas the oldest was 122 years old). As a result, extreme deviations are not  
406 possible. This is not true for software testing, where testing effort can drastically vary across software systems.  
407 For example, one can safely state that testing an industrial system is vastly more complex than testing a method  
408 implementing the triangle classification problem. None of the papers surveyed in Section 3 report skewness or  
409 kurtosis values. Although meta-analyses of the literature are hence not possible, the following arguments cast  
410 even further doubts about the applicability of parametric tests to analyze randomized algorithms in software  
411 testing.

412 Random testing is perhaps the easiest and most known automated software testing technique. It is often  
413 recommended as a comparison baseline to assess whether novel testing techniques are indeed useful [57]. When  
414 random testing is used to find a test case for a specific testing target (e.g., a test case that triggers a failure or  
415 covers a particular branch/path), it follows a geometric distribution. When there is more than one testing target,  
416 e.g., full structural coverage, it follows a coupon’s collector problem distribution [13]. Given  $\theta$  the probability  
417 of sampling a test case that covers the desired testing target, then the expectation (i.e., the average number of  
418 required test cases to sample) of random testing is  $\mu = 1/\theta$  and its variance is  $\delta^2 = (1 - \theta)/\theta^2$  [36].

419 Figure 2 plots the mass function of a geometric distribution with  $\theta = 0.01$  and a normal distribution  
420 with same  $\mu$  and  $\delta^2$ . In this context, the mass function represents the probability that, for a given number of  
421 sampled test cases  $l$ , the target is covered after sampling exactly  $l$  test cases. For random testing, the most  
422 likely outcome is  $l = 1$ , whereas for a normal distribution it is  $l = \mu$ . As it is easily visible from Figure  
423 2, the geometric distribution has a very strong departure from normality! Comparisons of novel techniques  
424 versus random testing (as this is common practice when search algorithms are evaluated [57]) using  $t$ -tests can  
425 be questionable if the number of repeated experiments is “low”. Furthermore, the probability distributions for

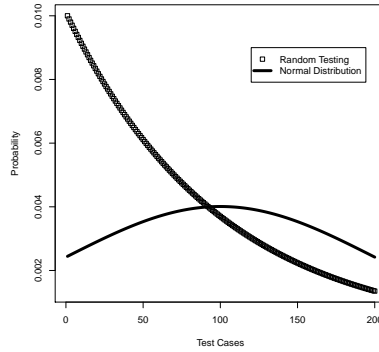


Figure 2: Mass and density functions of random testing and normal distribution given same mean  $\mu = 1/\theta$  and variance  $\sigma^2 = (1 - \theta)/\theta^2$ , where  $\theta = 0.01$ .

426 performance  $M$  (recall Section 4) for search algorithms may also strongly depart from normality. A common  
 427 example is when the search landscape of the addressed problem has trap-like regions [91].

428 Violations of the assumptions of a statistical test such as  $t$ -test can be tolerated as long as they are not too  
 429 “large” (where “large” can be somehow quantified with the kurtosis value [49]). Empirical evidence suggests  
 430 that to be the case for natural and social sciences, and therefore probably so for empirical studies in software  
 431 engineering involving human subjects. On the other end, there is no evidence at all in the literature that con-  
 432 firms it should be the case for randomized algorithms, used for example in the context of software testing. The  
 433 arguments presented in this section actually cast doubts on such possibility. As long as no evidence is provided  
 434 in the randomized algorithm literature to disprove the above concerns, in software testing or other fields of ap-  
 435 plications, one should not blindly follow guidelines provided for experiments with human subjects in software  
 436 engineering or other experimental fields.

## 437 5.2 Central Limit Theorem

438 The Central Limit theorem states that the *sum* of  $n$  random variables converges to a normal distribution [36]  
 439 as  $n$  increases. For example, consider the result of throwing a die. There are only six possible outcomes,  
 440 each one with probability  $1/6$  (assuming a fair die). If one considers the *sum* of two dice (i.e.,  $n = 2$ ), there  
 441 would be 11 possible outcomes, from value 2 to 12. Figure 3 shows that with  $n = 2$ , in the case of dice,  
 442 a distribution that resembles the normal one is already obtained, even though with  $n = 1$  it is very far from  
 443 normality. In the research context of this paper, these random variables are the results of the  $n$  runs of the  
 444 analyzed algorithm. This theorem makes four assumptions: the  $n$  variables should be independent, coming  
 445 from the same distribution and their mean  $\mu$  and variance  $\delta^2$  should exist (i.e., they should be different from  
 446 infinity). When using randomized algorithms, having  $n$  independent runs coming from the same distribution  
 447 (e.g., the same algorithm) is usually trivial to achieve (one just needs to use different seeds for the pseudo-  
 448 random generators). But the existence of the mean and variance requires more scrutiny. As shown before, those  
 449 values  $\mu$  and  $\delta^2$  exist for random testing. A well known “paradox” in statistics in which mean and variance do  
 450 not exist is the Petersburg Game [36]. Similarly, the existence of mean and variance in search algorithms is not  
 451 always guaranteed, as discussed next.

452 To put this discussion on a more solid ground, the Petersburg Game is here briefly described. Assume  
 453 a player tosses an unbiased coin until a head is obtained. The player first gives an amount of money to the  
 454 opponent which needs to be negotiated, and then she receives from the opponent an amount of money (Kroner)  
 455 equal to  $k = 2^t$ , where  $t$  is the number of times the coin was tossed. For example, if the player obtains two  
 456 tails and then a head, then she would receive from the opponent  $k = 2^3 = 8$  Kroner. *On average*, how many  
 457 Kroner  $k$  will she receive from the opponent in a single match? The probability of having  $k = 2^x$  is equivalent  
 458 to get first  $x - 1$  tails and then one head, so  $p(2^x) = 2^{-(x-1)} \times 2^{-1} = 2^{-x}$ . Therefore, the average reward is  
 459  $\mu = E[k] = \sum_k kp(k) = \sum_t 2^t p(2^t) = \sum_t 2^t \times 2^{-t} = \sum_t 1 = \infty$ . Unless the player gives an *infinite* amount  
 460 of money to the opponent before starting tossing the coin, then the game would not be fair *on average* for the

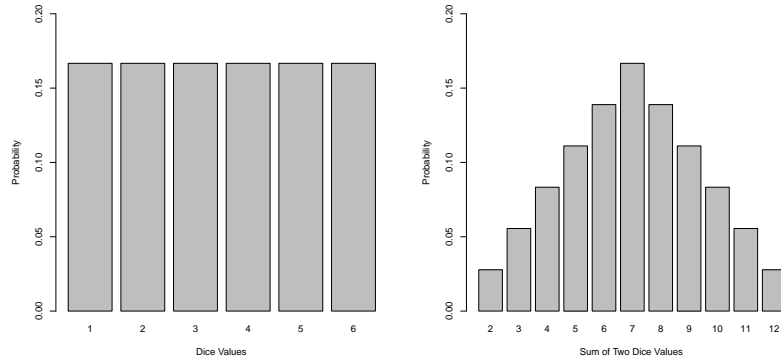


Figure 3: Density functions of the outputs of one dice and the sum of two dice.

461 opponent! This is a classical example of a random variable where it is not intuitive to see that it has no finite mean  
 462 value. For example, obtaining  $t > 10$  is very unlikely, and if one tries to repeat the game  $n$  times, the average  
 463 value for  $k$  would be quite low and would be a very wrong estimate of the actual, theoretical average (infinity).

464 Putting the issue illustrated by the Petersburg Game principle in the research context of this paper, if the  
 465 performance of a randomized algorithm is bounded within a predefined range, then the mean and variance  
 466 always exist. For example, if an algorithm is run for a predefined amount of time to achieve structural test  
 467 coverage, and if there are  $z$  structural targets, then the performance of the algorithm would be measured with a  
 468 value between 0 and  $z$ . Therefore, one would have  $\mu \leq z$  and  $\delta^2 \leq z^2$ , thus making the use of a  $t$ -test valid.

469 The problems arise if no bound is given on how the performance is measured. A randomized algorithm  
 470 could be run until it finds an optimal solution to the addressed problem. For example, random testing could be  
 471 run until the first failure is triggered (assuming an automated oracle is provided). In this case, the performance  
 472 of the algorithm would be measured in the number of test cases that are sampled before triggering the failure  
 473 and there would be no upper limit for a run. If a researcher runs a search algorithm on the same problem  $n$   
 474 times, and he has  $n$  variables  $X_i$  representing the number of test cases sampled in each run before triggering  
 475 the first failure, the mean would be estimated as  $\hat{\mu} = \frac{1}{n} \sum_{i=1}^n X_i$ , and one would hence conclude that the mean  
 476 exists. As the Petersburg Game shows, this can be wrong, because  $\hat{\mu}$  is only an *estimation* of  $\mu$ , which might  
 477 not exist.

478 For most search algorithms convergence in finite time is proven under some conditions (e.g., [100]), and  
 479 hence mean and variance exist. But in software engineering, when new problems are addressed, standard search  
 480 algorithms with standard search operators may not be usable. For example, when testing for object-oriented  
 481 software using search algorithms (e.g., [114]), complex non-standard search operators are required. Without  
 482 formal proofs (e.g., as done by Fraser and Arcuri [40]), it is not safe to speak about the existence of the mean  
 483 in those cases.

484 However, the non-existence of the mean is usually not a problem from a practical standpoint. In practice,  
 485 there usually are upper limits to the amount of computational resources a randomized algorithm can use. For  
 486 example, a search algorithm can be prematurely stopped when reaching a time limit. Random testing could  
 487 be stopped after 100,000 sampled test cases if it has found no failure so far. But, in these cases, one is actu-  
 488 ally dealing with *censored* data [71] (in particular, right-censorship) and this requires proper care in terms of  
 489 statistical testing and the interpretation of results, as it will be discussed in Section 6.

### 490 5.3 Differences in the Compared Properties

491 Even under proper conditions for using a parametric test, one aspect that is often ignored is that the  $t$ -test and  
 492 U-test analyze two different properties. Consider a random testing example in which one counts the number of  
 493 test cases run before triggering a failure. Considering a failure rate  $\theta$ , the mean value of test cases sampled by  
 494 random testing is hence  $\mu = 1/\theta$ . Assume that a novel testing technique  $\mathcal{A}$  yields a normal distribution of the  
 495 required number of test cases to trigger a failure. If one further considers the same variance as random testing  
 496 and a mean that is 85% of that of random testing, which one is better? Random testing with mean  $\mu$  or  $\mathcal{A}$  with

497 mean  $0.85\mu$ ? Assuming a large number of runs (e.g.,  $n$  is equal to one million), a  $t$ -test would state that  $\mathcal{A}$  is  
498 better, whereas a Mann-Whitney U-test would state exactly the opposite. How come? This is not an error as the  
499 two tests are measuring different things: The  $t$ -test measures the difference in mean values whereas the Mann-  
500 Whitney U-test deals with their stochastic ranking, i.e., whether observations in one data sample are more likely  
501 to be larger than observations in the other sample. Notice that this latter concept is technically different from  
502 detecting difference in *median* values (which can be stated only if the two distributions have same shape). In  
503 a normal distribution, the median value is equal to the mean, whereas in a geometric distribution the median is  
504 roughly 70% of the mean [36]. On one hand, half of the data points for random testing would be lower than  
505  $0.7\mu$ . On the other hand, with  $\mathcal{A}$  half of the data points would be above  $0.85\mu$ , and a significant proportion  
506 between  $0.7\mu$  and  $0.85\mu$ . This explains the apparent contradiction in results: though the average is higher for  
507 random testing, its median is lower than that of  $\mathcal{A}$ .

508 From a practical point of view, which statistical test should be used? Based on the discussions in this  
509 section, and in line with Leech and Onwuegbuzie [76], it is recommendable to use Mann-Whitney U-test (to  
510 assess difference in stochastic order) rather than the  $t$ -test and Welch test (to assess difference in mean values).  
511 However, the full motivation will become clearer once censored data, effect size, and the choice of  $n$  will be  
512 discussed in the next sections.

## 513 5.4 Rank Transformation

514 There is an important aspect that needs to be considered: data can be “transformed” before being given as input to  
515 a statistical test. As discussed by Ruxton [101], a Welch test can be used instead of a U-test if the raw values in  
516 the data are replaced by their rank. For example, consider the data set  $\{24, 2, 274\}$  discussed in the introduction  
517 regarding random testing. Those values could be substituted with their ranks  $\{2, 1, 3\}$  before being given as  
518 input to a statistical test. What would be the motivation of doing so? The U-test might be negatively affected if  
519 the two compared distributions have “significantly” different variance, and in such case a Welch test on ranked  
520 data might be better (in the sense that it would have lower probability of Type I and II errors). However, the  
521 Welch test would still be negatively affected by violations of the normality assumption (ranked data might not  
522 be normal). Ruxton [101] reports on some cases in which a Welch test on ranked data is better than a U-test, but  
523 the results of those *empirical* analyses might not generalize to the context of randomized algorithms applied to  
524 software engineering problems.

525 For simplicity and because it has widespread applications, the authors of this paper recommend to use a U-  
526 test rather than a Welch test on ranked data. There might be cases in which this latter test could be preferable, but  
527 it might be difficult, for a non-expert in statistics, to clearly identify those cases. Nevertheless, it is important to  
528 clarify that a Welch test on ranked data does not assess any more whether there is a statistical difference among  
529 the mean values of the two compared distributions. Rather, it assesses differences in mean values of the ranks  
530 and therefore determine whether there is any difference in stochastic ordering between the two distributions.  
531 For example, assume the two data sets  $X = \{1, 2, 3, 4, 5, 6, 49\}$  and  $Y = \{7, 8, 9, 10, 11, 12, 13\}$ . If it were  
532 not for the “outlier” 49 in  $X$ , then all the values in  $Y$  would be greater than the values in  $X$ . Both data sets  
533 have a mean value equal to 10. A Welch test on raw values would result in a  $p$ -value equal to 1, which is not  
534 surprising considering that the two data sets have the same mean. However, if one does a rank transformation,  
535 then the outlier 49 would be replaced by the value 14 (all the other values in  $X$  and  $Y$  remain the same). In this  
536 case, the resulting  $p$ -value of the Welch test would be 0.02, which suggests a strong difference in the stochastic  
537 ordering (i.e., ranks) between the two distributions.

## 538 5.5 Test for Randomized vs Deterministic Algorithm

539 In the discussions above, it was assumed that both algorithms  $\mathcal{A}$  and  $\mathcal{B}$  are randomized. If one of them is  
540 deterministic (e.g.,  $\mathcal{B}$ ), it is still important to use statistical testing. Consistent with the above recommendation,  
541 the non-parametric *One-Sample Wilcoxon* test should be used. Given  $m_{\mathcal{B}}$  the performance measure of the  
542 deterministic algorithm, a one-sample Wilcoxon test would verify whether the performance of  $\mathcal{A}$  is symmetric  
543 about  $m_{\mathcal{B}}$ , i.e., whether by using  $\mathcal{A}$  one is as likely to obtain a value lower than  $m_{\mathcal{B}}$  as otherwise.

## 544 6 Censored Data

545 Assume that the result of an experiment is dichotomous: either one finds a solution to solve the software  
546 engineering problem at hand (*success*), or he does not (*failure*). For example, in software testing, if the goal is  
547 to cover a particular target (e.g., a specific branch), one can run a randomized algorithm with a time limit  $L$ ,  
548 chosen based on available computing resources. The algorithm will be stopped as soon as a solution is found,  
549 otherwise the search stops after time  $L$ . Another example is bug fixing [117] where one finds a patch within  
550 time  $L$ , or does not.

551 The above types of experiments are dealing with *right-censored* data, and their properties are equivalent to  
552 survival/failure time analysis [71, 41]. Let  $X$  be the random variable representing the time a randomized algo-  
553 rithm takes to solve a software engineering problem, and consider  $n$  experiments in which a researcher collects  
554  $X_i$  values. This is a case of right-censorship since, assuming a time limit  $L$ , one will not have observations  
555  $X_i$  for the cases  $X > L$ . Although there are several ways to deal with this problem [71], in this paper the  
556 discussions are limited to simple solutions.

557 One interesting special case is when one cannot say for sure whether the chosen target has been achieved,  
558 e.g., generation of test cases that achieve code branch coverage. Putting aside trivial cases, there are usually  
559 infeasible targets (e.g., unreachable code) and their number is unknown. As a result, such experiments are  
560 not dichotomous because one cannot know whether all feasible targets have been covered. Even when using a  
561 time limit  $L$ , these cases would still not be considered as involving censored data. However, if in the experi-  
562 ments the comparisons are made reusing artifacts from published studies in the literature, and if one wants to  
563 know whether or not, within a given time, he can obtain better coverage than these reported studies, then such  
564 experiments can be considered dichotomous despite infeasible targets.

565 Consider the case in which one needs to compare two randomized algorithms  $\mathcal{A}$  and  $\mathcal{B}$  on a software  
566 engineering problem with dichotomous outcome. Let  $X$  be the random variable representing the time  $\mathcal{A}$  takes  
567 to find a valid solution, and let  $Y$  be the same type of variable for  $\mathcal{B}$ . Assume that a researcher runs  $\mathcal{A}$  and  $\mathcal{B}$   
568  $n$  times, collecting observations  $X_i$  and  $Y_i$ , respectively. Using a time limit  $L$ , to evaluate which of the two  
569 algorithms is better, one can consider their *success rate*  $\gamma = k/n$ , i.e., the proportion of number of times  $k$ , out  
570 of the  $n$  runs, for which a valid solution is found. To evaluate whether there is statistical difference between the  
571 success rates of  $\mathcal{A}$  and  $\mathcal{B}$ , a test for differences in proportions is then appropriate, such as the Fisher exact test  
572 [71].

573 The Fisher exact test is a parametric test, which assumes that the analyzed data follows a binomial distribu-  
574 tion. In contrast to other parametric tests (e.g., the  $t$ -test), its assumptions are always valid: if the experiments  
575 are independent, then the success rate of a series of randomized experiments would always follow a binomial  
576 distribution, where  $\gamma$  represents the estimated probability of success. Furthermore, for values of  $n$  until roughly  
577 100, the test is “exact”. This means that the resulting  $p$ -values are precise, and not estimates based on how close  
578 the data are from satisfying the conditions of a test (e.g., normality and equal variance in a  $t$ -test). However, for  
579 larger values of  $n$ , the computational cost of the test would start to be too prohibitive, and approximations are  
580 then used to calculate the  $p$ -values (this is often done automatically in many statistical tools).

581 Assume that out of  $n = 100$  runs the success rate of  $\mathcal{A}$  is  $\gamma_{\mathcal{A}} = 1\%$ , whereas for  $\mathcal{B}$  it is  $\gamma_{\mathcal{B}} = 5\%$ . A  
582 Fisher exact test has a resulting  $p$ -value equal to 0.21, which might be considered high, i.e., there is a 21%  
583 probability that the success rates of the two algorithms are actually equal. In such cases, one can run more  
584 experiments (i.e., increase  $n$ ) to obtain higher statistical power (i.e., decrease the  $p$ -value). Alternatively, if  
585 there is no statistically or practically significant difference between the success rates of  $\mathcal{A}$  and  $\mathcal{B}$ , a practical  
586 question is then to determine which technique uses *less* time. This is particularly relevant if the success rates  
587 of both techniques are high. There can be different ways to analyze such cases, such as considering artificial  
588 censorships at different times before  $L$ . For example, one can consider censorship at  $L/2$ , i.e., the success rate  
589 with half the time, and determine which technique still fares better and whether its success rate is acceptable.  
590 Note that such analysis does not require to run any further experiments as success rates can be computed at  
591  $L/2$  from existing runs. Another alternative to compare execution times is to apply a Mann-Whitney U-test,  
592 recommended above, using only the times of successful runs, which have  $X_i$  and  $Y_i$  values lower or equal to  $L$ .

593 A more complex situation is when one algorithm shows a significantly higher success rate, but takes more  
594 time to produce valid solutions than the other. This is a typical situation, that is not so uncommon, where  
595 a choice needs to be made. For example, on one hand, a *local search* [81] might be very fast in generating  
596 appropriate testing data if it starts from the right area of the search landscape. But, at the same time, it could

597 yield a low success rate if most of the search landscape has gradient toward local optima, and if the number  
598 of such optima is low. (Notice that this is just an example: it is not in the scope of the paper to give lengthy  
599 explanations of why that would be a problem for local search; see the work of Arcuri [8] for further details on  
600 this topic.) On the other hand, a population-based search algorithm, such as Genetic Algorithms, could avoid  
601 the problem of local optima, which in turn would result in higher success rate than a local search. However,  
602 because an entire population is evolved at the same time, depending on the selection pressure of the algorithm  
603 (e.g., the value of the tournament size in tournament selection) and the population size, a Genetic Algorithm  
604 might take much longer than a local search to converge towards a solution in its successful runs.

## 605 7 Effect Size

606 When comparing a randomized algorithm  $\mathcal{A}$  against another  $\mathcal{B}$ , given a large enough number of runs  $n$ , it is  
607 most of the time possible to obtain statistically significant results with a  $t$ -test or U-test. Indeed, two different  
608 algorithms are extremely unlikely to have exactly the same probability distribution. In other words, with a large  
609 enough  $n$  one can obtain statistically significant differences even if they are so small as to be of no practical  
610 value.

611 Though it is important to assess whether an algorithm fares statistically better than another, it is in addition  
612 crucial to assess the magnitude of the improvement. To analyze such a property, *effect size* measures are needed  
613 [55, 63, 89]. Effect sizes can be divided in two groups: standardized and unstandardized. Unstandardized  
614 effect sizes are dependent on the unit of measurement used in the experiments. Consider the difference in  
615 means between two algorithms  $\Delta = \mu^{\mathcal{A}} - \mu^{\mathcal{B}}$ . This value  $\Delta$  has a measurement unit, that of  $\mathcal{A}$  and  $\mathcal{B}$ . For  
616 example, in software testing,  $\mu$  can be the expected number of test executions to find the first failure. On one  
617 testing artifact it could be that  $\Delta_1 = \mu^{\mathcal{A}} - \mu^{\mathcal{B}} = 100 - 1 = 99$ , whereas on another testing artifact it can be  
618  $\Delta_2 = \mu^{\mathcal{A}} - \mu^{\mathcal{B}} = 100,000 - 200,000 = -100,000$ . Deciding based on  $\Delta_1$  and  $\Delta_2$  which algorithm is better  
619 is difficult to determine since the two scales of measurement are different.  $\Delta_1$  is very low compared to  $\Delta_2$ , but  
620 in that case  $\mathcal{A}$  is 100 times worse than  $\mathcal{B}$ , whereas it is only twice as fast in the case  $\Delta_2$ .

621 Empirical analyses of randomized algorithms, if they are to be reliable and generalizable, require the use of  
622 large numbers of artifacts (e.g., programs). The complexity of these artifacts is likely to widely vary, such as  
623 the number of test cases required to fulfill a coverage criterion on various programs. The use of standardized  
624 effect sizes, that are independent from the evaluation criteria measurement unit, is therefore necessary to be  
625 able to compare results across artifacts and experiments. In their systematic review of empirical analyses in  
626 software engineering involving controlled experiments with human subjects, Kampenes *et al.* [63] found that  
627 standardized effect sizes were reported in only 29% of the cases. In the systematic review performed in this  
628 paper, only one paper [96] was found, which uses the Vargha and Delaney’s  $\hat{A}_{12}$  statistics (described later in  
629 this section).

630 In this section, the most known standardized effect size measure is described first followed by an expla-  
631 nation of why it should *not* be used when analyzing randomized algorithms applied in software engineering.  
632 Then, two other standardized effect sizes are described, and instructions are given on how to apply them in  
633 practice.

634 The most known effect size is the so called  $d$  family which, in the general form, is  $d = (\mu^{\mathcal{A}} - \mu^{\mathcal{B}})/\sigma$ .  
635 In other words, the difference in mean is scaled over the standard deviation (several corrections exists to this  
636 formula, but for more details please see the book of Grissom and Kim [55]). Though one obtains a measure that  
637 has no measurement unit, the problem is that it assumes normality of the data, and strong departures can make  
638 it meaningless [55]. For example, in a normal distribution, roughly 64% of the points lie within  $\mu \pm \sigma$  [36],  
639 i.e., they are at most  $\sigma$  away from the mean  $\mu$ . But for distributions with high skewness (as in the geometric  
640 distribution and as it is often the case for search algorithms), the results of scaling the mean difference by the  
641 standard deviation “would not be valid”, because “standard deviations can be very sensitive to a distribution’s  
642 shape” [55]. In this case, a non-parametric effect size should be preferred. Existing guidelines [63, 89] only  
643 briefly discuss the use of non-parametric effect sizes.

644 The Vargha and Delaney’s  $\hat{A}_{12}$  statistic is a non-parametric effect size measure [116, 55]. Its use has  
645 been advocated by Leech and Onwuegbuzie [76], and one example of its use in software engineering in which  
646 randomized algorithms are involved can be found in the work of Poulding and Clark [96]. In the research  
647 context of this paper, given a performance measure  $M$ ,  $\hat{A}_{12}$  measures the probability that running algorithm  $\mathcal{A}$



648 yields higher  $M$  values than running another algorithm  $\mathcal{B}$ . If the two algorithms are equivalent, then  $\hat{A}_{12} = 0.5$ .  
 649 This effect size is easier to interpret compared to the  $d$  family. For example,  $\hat{A}_{12} = 0.7$  entails one would obtain  
 650 better results 70% of the time with  $\mathcal{A}$ . Though this type of non-parametric effect size is not common in statistical  
 651 tools, it can be very easily computed [76, 55]. The following formula is reported in the work of Vargha and  
 652 Delaney [116]:

$$\hat{A}_{12} = (R_1/m - (m + 1)/2)/n \quad (1)$$

653 where  $R_1$  is the rank sum of the first data group under comparison. For example, assume the data  $X =$   
 654  $\{42, 11, 7\}$  and  $Y = \{1, 20, 5\}$ . The data set  $X$  would have ranks  $\{6, 4, 3\}$ , whose sum is 13, whereas  $Y$  would  
 655 have ranks  $\{1, 5, 2\}$ . The rank sum is a basic component in the Mann-Whitney U-test, and most statistical tools  
 656 provide it. In Equation 1,  $m$  is the number of observations in the first data sample, whereas  $n$  is the number of  
 657 observations in the second data sample. In most experiments, one would run two randomized algorithms the  
 658 same number of times:  $m = n$ .

659 When dealing with dichotomous results (as discussed in Section 6), several types of effect size measures  
 660 [55] can be considered. The *odds ratio* is the most used and “is a measure of how many times greater the odds  
 661 are that a member of a certain population will fall into a certain category than the odds are that a member of  
 662 another population will fall into that category” [55]. Given  $a$  the number of times algorithm  $\mathcal{A}$  finds an optimal  
 663 solution, and  $b$  for algorithm  $\mathcal{B}$ , the odds ratio is calculated as

$$\psi = \frac{a + \rho}{n + \rho - a} / \frac{b + \rho}{n + \rho - b}, \quad (2)$$

664 where  $\rho$  is any arbitrary positive constant (e.g.,  $\rho = 0.5$ ) used to avoid problems with zero occurrences [55].  
 665 There is no difference between the two algorithms when  $\psi = 1$ . The cases in which  $\psi > 1$  imply that algorithm  
 666  $\mathcal{A}$  has higher chances of success.

667 Both  $\hat{A}_{12}$  and  $\psi$  are standardized effect size measures. But because their calculation is based on a finite  
 668 number of observations (e.g.,  $n$  for each algorithm, so  $2n$  when two algorithms are compared), they are only  
 669 estimates of the real  $\hat{A}_{12}^*$  and  $\psi^*$ . If  $n$  is low, these estimations might be very inaccurate. One way to deal with  
 670 this problem is to calculate *confidence intervals* (CI) for them [55]. A  $(1 - \alpha)$  CI is a set of values for which  
 671 there is  $(1 - \alpha)$  probability that the value of the effect size lies in that range. For example, if one has  $\hat{A}_{12} = 0.54$   
 672 and a  $(1 - \alpha)$  CI with range  $[0.49, 59]$ , then with probability  $(1 - \alpha)$  the real value  $\hat{A}_{12}^*$  lies in  $[0.49, 59]$  (where  
 673  $\hat{A}_{12} = 0.54$  is its most likely estimation). Such effect size confidence intervals can facilitate decision making  
 674 as they enable the comparison of the costs of alternative algorithms while accounting for uncertainty in their  
 675 estimates. To see how confidence intervals are calculated for  $\hat{A}_{12}$ , please see the book of Grissom and Kim [55]  
 676 or the work of Vargha and Delaney [116].

677 Furthermore, general techniques such as *bootstrapping* [24] can be employed to create confidence intervals  
 678 for  $\hat{A}_{12}$  or any other statistics of interest (e.g., mean and median). At a high level, bootstrapping works as  
 679 follows. Assume  $n$  experiments with results  $x_i$ . The arithmetic average would be calculated as  $\mu = \frac{\sum_{i=1}^n x_i}{n}$ .  
 680 Because  $n$  is finite,  $\mu$  is only an estimate of the real average (e.g., recall the Petersburg Game discussed in  
 681 Section 5.2). By defining  $X$  as the set of  $n$  results  $x_i$ , bootstrapping works by resampling  $n$  values with  
 682 replacement from  $X$  and by calculating the statistics of interest (e.g., the mean) on this new set (e.g.,  $\mu_j$ ).  
 683 This process is repeated  $k$  times (e.g.,  $k = 1,000$ ), which provides  $k$  values for the statistics of interest (e.g.,  
 684  $\mu_1, \mu_2, \dots, \mu_k$ ). Then, several different techniques can be used to create a confidence interval at level  $\alpha$   
 685 from these  $k$  estimates. For more details on the properties of bootstrapping, the interested reader is referred to  
 686 Chernick’s book [24].

687 Notice that a confidence interval can replace a test of statistical difference (e.g.,  $t$ -test and U-test). If the  
 688 null hypothesis  $H_0$  lies within the confidence interval, then there is insufficient evidence to claim there is a  
 689 statistically significant difference. In the previous example, because 0.5 is inside the  $(1 - \alpha)$  CI  $[0.49, 59]$ , then  
 690 there is no statistical difference at the selected significance level  $\alpha$ . For a dichotomous result,  $H_0$  would be  
 691  $\psi = 1$ .

## 692 8 Number of Runs

693 How many runs does a researcher need when analyzing and comparing randomized algorithms? A general  
694 answer is: As many as necessary to show with high confidence that the obtained results are statistically sig-  
695 nificant and to obtain a small enough confidence interval for effect size estimates. In many fields of science  
696 (e.g., medicine and behavioral science), a common rule of thumb is to use at least  $n = 30$  observations. In the  
697 many fields where experiments are very expensive and time consuming, it is in general not feasible to work  
698 with high values for  $n$ . Several new statistical tests have been proposed and discussed to cope with the problem  
699 of lack of power and violation of assumptions (e.g., normality of data) when smaller numbers of observations  
700 are available [119].

701 Empirical studies of randomized algorithms usually do not involve human subjects and the number of *runs*  
702 (i.e.,  $n$ ) is only limited by computational resources. When there is access to clusters of computers as this is  
703 the case for many research institutes and universities, and when there is no need for expensive, specialized  
704 hardware (e.g., hardware-in-the-loop testing), then large numbers of runs can be carried out to properly analyze  
705 the behavior of randomized algorithms. Many software engineering problems are furthermore not highly com-  
706 putationally expensive, as for example code coverage at the unit testing level, and can therefore involve very  
707 large numbers of executions. There are however exceptions, such as the system testing of embedded systems  
708 (e.g., [12]) where each test case can be very expensive to run.

709 Whenever possible, in most cases, it is therefore recommended to use a very high number of runs. For  
710 most problems in software engineering, thousands of randomized algorithm runs should be feasible and would  
711 solve most of the problems related to the power and accuracy of statistical tests. For example, as illustrated  
712 in references [83, 32] in Table 2, even with 100 runs, the U-test might not be powerful enough to confirm a  
713 statistical difference at a 0.05 significance level, even when the data seems to suggest such a difference.

714 Most discussions in the literature about statistical tests focus on situations with small numbers of observa-  
715 tions (e.g., [101]). However, with thousands of runs, one would detect statistically significant differences on  
716 practically any experiment (Section 4). It is hence essential to complement such analyses with a study of the  
717 effect size as discussed in Section 7. Even when having large numbers of runs is not necessary, for a set  $\alpha$  level  
718 (e.g., 0.05), to obtain differences that are large enough to show  $p$ -values less than  $\alpha$ , additional runs would help  
719 tighten the confidence intervals for effect size estimates and would be of practical value to support decision  
720 making.

721 In Section 4, it was suggested to use U-test instead of  $t$ -test. For very large samples, such as  $n = 1,000$ ,  
722 there would be no practical difference between them regarding power and accuracy. However, the choice of a  
723 non-parametric test would be driven by its corresponding effect size measure. In Section 7 it was argued that  
724 effect size measures based on the mean (i.e., the  $d$  family) were not appropriate for randomized algorithms in  
725 software engineering due to violations in distribution assumptions. It would then be inconsistent to investigate  
726 the statistical difference of mean values with a  $t$ -test if one cannot use a reliable measure for its effect size.  
727 In other words, it is advisable to use size measures that are consistent with the differences being tested by the  
728 selected statistical test.

## 729 9 Multiple Tests

730 In most situations, researchers need to compare several alternative algorithms. Furthermore, if one is comparing  
731 different algorithm settings (e.g., population size in a Genetic Algorithm), then each setting technically defines  
732 a different algorithm [11]. This often leads to a large number of statistical comparisons. It is possible to use  
733 statistical tests that deal with multiple techniques (treatments, experiments) at the same time (e.g., Factorial  
734 ANOVA), and effect sizes have been defined for those cases [55]. There are several types of statistical tests  
735 addressing multiple comparisons, and the choice depends on which research question one is addressing. This  
736 paper only deals with the two most common research questions, since several books are dedicated to this topic,  
737 and an exhaustive analysis would not be possible:

- 738 • Does the choice of a particular parameter affect the performance of a randomized algorithm?
- 739 • Among a set of randomized algorithms, which one is the best in solving the addressed problem?

740 Given a parameter that can take several different values  $j \in J$ , assume a researcher has carried out a series  
741 of experiments for a set of parameter values  $\{j_1, j_2, \dots, j_k\} \subseteq J$ . For example, in a Genetic Algorithm,  
742 one might want to study whether applying different cross-over rates has any effect on the effectiveness of the  
743 algorithm. One could consider the values  $\{0, 0.25, 0.5, 0.75, 1\}$ , and have  $n = 1,000$  independent experiments  
744 for each of these five rate values. If the goal is to evaluate whether the choice of this rate has any effect on  
745 the effectiveness of a Genetic Algorithm, then an *omnibus* test such as ANOVA can be employed. The null  
746 hypothesis is that the choice of the parameter value has no effect on the mean effectiveness of the algorithm.  
747 However, ANOVA suffers of the same problems as the *t*-test, i.e., assumption about normality of the data and  
748 equal variance. A non-parametric equivalent is the so called Kruskal-Wallis test [73].

749 Assume that the result of a Kruskal-Wallis test suggests that the choice of that crossover rate has a statisti-  
750 cally significant effect (i.e., the resulting *p*-value is low, so one can reject the null hypothesis). A relevant  
751 question might then be which crossover rate should be used (i.e., which one gives the best performance?). An  
752 omnibus test is not able to answer such a research question. This situation is exactly equivalent to the case of  
753 identifying the best algorithm among  $K = 5$  algorithms/variants. In this case, one would like to individually  
754 compare the performance of each algorithm against all other alternatives. Given a set of algorithms, a researcher  
755 would not be interested in simply determining whether all of them have the same mean values. Rather, given  
756  $K$  algorithms, one wants to perform  $Z = K(K - 1)/2$  pairwise tests and measure effect size in each case.

757 However, using several statistical tests inflates the probability of Type I error. If one has only one com-  
758 parison, the probability of Type I error is equal to the obtained *p*-value. On the other hand, if one has many  
759 comparisons, even when all the *p*-values are low, there is usually a high probability that at least in one of the  
760 comparisons the null hypothesis is true as all these probabilities somehow add up. In other words, if in all the  
761 comparisons the *p*-values are lower than  $\alpha$ , then a researcher would normally reject all the null hypotheses. But  
762 the probability that at least one null hypothesis is true could be as high as  $1 - (1 - \alpha)^Z$  for  $Z$  comparisons,  
763 which converges to 1 as  $Z$  increases.

764 One way to address this problem is to use the so called *Bonferroni adjustment* [94, 88]. Instead of applying  
765 each test assuming a significance level  $\alpha$ , a researcher would use an adjusted level  $\alpha/Z$ . For example, if the  
766 probability of Type I error is selected to be 0.05 and two comparisons are performed, two statistical tests are run  
767 with  $\alpha = 0.025$  to check whether both differences are significant (i.e., if both *p*-values are lower than 0.025).  
768 However, the Bonferroni adjustment has been repeatedly criticized in the literature [94, 88], and the authors of  
769 this paper largely agree with those critiques. For example, assume that for both those tests the researcher obtains  
770 *p*-values equal to 0.04. If a Bonferroni adjustment is used, then both tests will not be statistically significant  
771 with  $\alpha = 0.05$ . It would then be tempting to publish the results of only one of them and claiming statistical  
772 significance because  $0.04 < 0.05$ . Such a practice can therefore hinder scientific progress by reducing the  
773 number of published results [94, 88]. This would be particularly true when many randomized algorithms can  
774 be compared to address the same software engineering problem: it would be very tempting to leave out the  
775 results of some of the poorly performing algorithms. Notice that there are other adjustment techniques that are  
776 equivalent to Bonferroni but that are less conservative [44]. However, the statistical significance of a single  
777 comparison would still depend on the number of performed and reported comparisons. Though in general it  
778 is not recommend to use the Bonferroni adjustment, it is important to always report the obtained *p*-values, not  
779 just whether a difference is significant or not at an arbitrarily chosen  $\alpha$  level. If for some reasons the readers  
780 want to evaluate the results using a Bonferroni adjustment or any of its (less conservative) variants, then it is  
781 possible to do so. For a full list of other problems related to the Bonferroni adjustment, the reader is referred to  
782 the work of Perneger [94] and Nakagawa [88].

783 Instead of pairwise tests using Bonferroni-like corrections, another (less popular) approach is to use the so  
784 called *post-hoc* methods, such as the Tukey's range test. This test is applied on each of the  $Z$  pairs, and it is  
785 very similar to a *t*-test. Similar to the Bonferroni method, it employs a *p*-value correction to handle possible  
786 inflation of probability of Type I error.

787 At any rate, alpha level adjustments can be very important when assessing the validity of behavioral or nat-  
788 ural phenomena with high confidence. For example, the leading international journal *Nature* has the following  
789 *requirement*<sup>3</sup> for published research papers regarding multiple tests:

790 Multiple comparisons: When making multiple statistical comparisons on a single data set, authors should

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<sup>3</sup><http://www.nature.com/nature/authors/gta/index.html#a5.6>, accessed November 2011.

791 explain how they adjusted the alpha level to avoid an inflated Type I error rate, or they should select  
792 statistical tests appropriate for multiple groups (such as ANOVA rather than a series of t-tests).

793 However, in Section 4 it was stated that in software engineering in general, and for randomized algorithms  
794 in particular, one mostly deals with decision-making problems. For example, if one must test software, then  
795 one must choose one alternative among  $K$  different techniques. In this case, even if the  $p$ -values are higher  
796 than  $\alpha$ , the software needs to be tested anyhow and a choice must be made. In this context, Bonferroni-  
797 like adjustments make even less sense. Just keep using the current technique because there is no statistically  
798 significant difference at a prefixed arbitrary  $\alpha$  level is not optimal as it ignores available information.

799 Assume that a researcher has analyzed the performance of  $K$  algorithms using pairwise tests and effect  
800 sizes. How to visualize the results of such analyses to grasp how their performance relate? There can be  
801 different ways (e.g., see the recent work of Carrano *et al.* [23]), and the description of a simple but practical  
802 technique is here provided, which was used for example by Fraser and Arcuri [38].

803 In their work [38], the effects of six parameters of a search algorithm were investigated in the context of  
804 automated unit testing of object-oriented software. Five parameters are binary (**Bo**, **Xo**, **Ra**, **Pa** and **Be**) and  
805 one ternary (**W**), for a total of  $2^5 \times 3 = 96$  configurations. Each configuration was compared against all the  
806 other 95 (i.e., a total of  $96 \times 95$  comparisons, which can be divided by two due to the symmetric property of the  
807 comparisons). Pairwise comparisons were made using a U-test, where the  $\alpha$  level was arbitrarily set to 0.05.  
808 Initially, a score of zero is assigned to each configuration. For each comparison in which a configuration is  
809 statistically better, its score is increased by one, whereas it is reduced by one in case it is statistically worse.  
810 Therefore, in the end each configuration obtains a score between -95 and 95, where the higher the score, the  
811 better the configuration. After this first phase, these scores are ranked such that the highest score has the best  
812 rank, where better ranks have lower values. In case of ties, the ranks are averaged. For example, if one has  
813 five configurations with scores  $\{10, 0, 0, 20, -30\}$ , then their ranks will be  $\{2, 3.5, 3.5, 1, 5\}$ . In the work  
814 of Fraser and Arcuri [38], this procedure was repeated for each artifact in the case study (i.e., for all the 100  
815 branches used in that empirical study), and the average of these ranks over all artifacts were calculated for each  
816 configuration, for a total of  $100 \times 96 \times 95/2 = 456,000$  statistical comparisons. After collecting all of these  
817 data, a table (reported in Table 4) was made in which the configurations were ordered based on their average  
818 rank from top (best) to bottom (worst). From this table, not only it is clear which are the best configurations,  
819 but it also possible to visualize some trends in the data (e.g., configurations with **Ra** are always better and **Xo**  
820 does not seem particularly useful). However, the above ranking mechanism has limitations, as it ignores the  
821 effect sizes and the actual  $p$ -values (e.g., a 0.051 value would be treated in the same way as a 1).

## 822 10 Experimenting With Several Artifacts

### 823 10.1 Choice of the Artifacts

824 When assessing randomized algorithms, the choice of artifacts to which these algorithms are applied (e.g.,  
825 source code or executable programs) is of paramount importance as it usually has a strong bearing on the  
826 evaluation results. When analyzing empirical analyses in the software engineering literature evaluating ran-  
827 domized algorithms, many of the studies are carried out on artificial and small artifacts. Empirical analyses  
828 on real industrial systems are rare, thus raising questions about the credibility of results and the usefulness of  
829 the proposed algorithms. However, achieving realism by using representative industrial systems is particularly  
830 challenging. One usually cannot precisely characterize the population of artifacts he is targeting in his studies.  
831 Even if a researcher could, he usually does not have access to large collections of industrial artifacts that are  
832 readily available to be sampled. And even if that were the case, studies are necessarily limited in terms of  
833 resources and time, and the number of artifacts studied is typically much more restricted than one would like.  
834 As a result, studies about randomized algorithms in software engineering typically present threats to external  
835 validity, making it difficult to generalize the results to other systems than the ones under study. In this paper,  
836 because the focus is on how to apply statistical tests, the details of how one should choose artifacts from a  
837 general standpoint are not emphasized. The following discussions in the paper rather concentrate on how this  
838 choice affects the statistical tests procedures and the number of runs required.

839 The first question one faces is whether the selected artifacts are *representative* of the type of problem that  
840 is being addressed. For example, assume one wants to evaluate a new tool for automatically generating unit

Table 4: Results of empirical analysis performed in the work of Fraser and Arcuri [38]. The table shows the performance of the the 96 configurations, ordered from top (best performance) to bottom (worst performance). Symbols are used to indicate whether a particular boolean parameter is activated.

Bo	Xo	Ra	Pa	Be	20	W 50	80	Av. Rank	Av. Success Rate
△		⊕	▽	⊞		W		31.475	0.464
△		⊕	▽			W		31.840	0.456
△		⊕		⊞		W		32.595	0.482
		⊕	▽	⊞		W		32.670	0.456
		⊕	▽			W		34.725	0.447
△		⊕				W		35.415	0.448
△		⊕		⊞		W		36.070	0.442
△		⊕		⊞	W			37.335	0.423
△	⊗	⊕	▽	⊞		W		37.430	0.430
△		⊕		⊞			W	37.605	0.459
△	⊗	⊕		⊞	W			37.615	0.418
△	⊗	⊕		⊞		W		38.080	0.422
	⊗	⊕	▽	⊞		W		39.325	0.419
	⊗	⊕		⊞		W		39.455	0.423
	⊗	⊕	▽			W		39.580	0.413
△		⊕			W			39.790	0.431
		⊕		⊞	W			39.815	0.431
	⊗	⊕			W			40.050	0.414
△		⊕	▽		W			40.140	0.420
△	⊗	⊕	▽			W		40.330	0.425
△		⊕	▽	⊞	W			40.670	0.413
△		⊕	▽	⊞			W	40.700	0.432
△	⊗	⊕		⊞	W			40.835	0.405
△		⊕		⊞			W	40.940	0.438
△		⊕	▽			W		41.200	0.455
△	⊗	⊕			W			41.350	0.410
		⊕	▽	⊞		W		41.695	0.423
		⊕	▽	⊞	W			41.890	0.405
		⊕	▽	⊞	W			41.925	0.413
	⊗	⊕	▽		W			42.150	0.399
	⊗	⊕	▽	⊞			W	42.195	0.401
	⊗	⊕	▽	⊞	W			42.470	0.388
△	⊗	⊕	▽		W			42.500	0.395
	⊗	⊕		⊞		W		42.800	0.422
		⊕			W			43.075	0.407
	⊗	⊕				W		43.095	0.421
△	⊗	⊕			W			43.255	0.420
△	⊗	⊕	▽	⊞	W			43.635	0.377
		⊕				W		45.160	0.398
	⊗	⊕	▽				W	45.205	0.393
		⊕	▽			W		45.285	0.412
△	⊗	⊕	▽			W		45.450	0.392
△		⊕				W		45.850	0.418
△		⊕				W		46.460	0.401
△	⊗	⊕				W		46.625	0.388
△	⊗	⊕		⊞		W		46.700	0.409
△	⊗	⊕	▽	⊞		W		47.760	0.379
△	⊗	⊕				W		47.850	0.384
△			▽	⊞		W		48.985	0.342
			▽			W		49.585	0.329
			▽	⊞		W		49.705	0.334
△			▽	⊞	W			49.995	0.369
△	⊗		▽	⊞		W		50.290	0.313
△			▽		W			50.740	0.356
△	⊗		▽			W		51.295	0.313
△			▽			W		51.350	0.340
△				⊞		W		51.570	0.327
△			▽	⊞			W	52.215	0.326
△			▽	⊞	W			52.800	0.330
			▽	⊞		W		53.260	0.330
	⊗		▽	⊞			W	53.610	0.309
△			▽			W		53.845	0.321
	⊗		▽	⊞	W			54.040	0.310
	⊗		▽			W		54.475	0.312
			▽	⊞	W			54.835	0.296
			▽		W			55.080	0.306
			▽	⊞		W		55.290	0.317
	⊗		▽			W		55.390	0.313
	⊗		▽	⊞		W		55.605	0.304
△					W			55.635	0.305
			▽			W		55.695	0.324
△	⊗		▽		W			56.065	0.310
△						W		56.160	0.309
△	⊗			⊞		W		56.200	0.304
△	⊗		▽	⊞			W	56.255	0.301
△	⊗		▽		W			56.295	0.312
△	⊗		▽	⊞	W			56.655	0.312
△	⊗					W		56.835	0.291
△	⊗					W		57.095	0.279
△	⊗			⊞		W		57.135	0.291
△				⊞		W		57.180	0.319
				⊞		W		57.390	0.306
				⊞		W		58.955	0.285
△	⊗			⊞		W		59.085	0.297
	⊗			⊞	W			59.190	0.297
△	⊗			⊞	W			59.270	0.285
	⊗					W		59.595	0.279
△						W		59.995	0.300
	⊗			⊞	W			60.145	0.281
	⊗					W		60.150	0.289
△	⊗			⊞	W			60.675	0.278
	⊗			⊞		W		60.705	0.289
△	⊗					W		60.975	0.292
						W		61.655	0.267
	⊗				W			65.220	0.238
					W			71.765	0.190

841 tests for object-oriented software (e.g., Pex [113], Randoop [93] or EvoSuite [40]). Which (types of) classes  
842 should be selected for experimenting? Following common practice in many empirical studies (e.g., [5, 98, 15]),  
843 is only using “container classes” acceptable? Arguably, it should depend on what is the target set of classes  
844 for the evaluation. If the proposed testing techniques are aimed *only* at container classes (e.g., [15]), then this  
845 would likely be acceptable. On the other hand, if the goal is to propose a *general* tool for generating unit tests,  
846 then using only container classes would lead to *serious* threats to external validity. But then the question is  
847 which classes should ideally be used? Again, one does not have well defined populations of classes that can be  
848 explicitly targeted and sampled. One possible simple heuristic is to try to maximize the diversity in terms of  
849 the type of classes, their size and complexity, and various other properties that are deemed relevant given the  
850 objective of the randomized algorithm, e.g., number of tasks accessing a lock when investigating deadlocks or  
851 data races [107].

852 As a practical alternative, one could use open source repositories such as SourceForge<sup>4</sup>, and randomly select  
853 a subset of projects for experimenting among the 319,000 that are currently hosted (as for example done by  
854 Fraser and Arcuri [39]). If one wants to evaluate the applicability of a general tool for unit testing, this would  
855 be better than using only container classes or arbitrarily choosing some programs in a non-systematic way (as  
856 it is often the case in the literature). However, even if one randomly samples projects from SourceForge, the  
857 empirical analyses would likely have some sort of bias. For example, open source projects in general may  
858 not be representative of programs developed in industry. Embedded systems and financial applications, for  
859 example, are unlikely to be well represented among these open source projects.

860 Regarding randomized algorithms (in particular search and optimization algorithms), there are specific  
861 and rigorous theoretical reasons for which the choice of artifacts is extremely important. The *No Free Lunch*  
862 theorem states that, on average across all possible problems (i.e., artifacts), all search algorithms have the same  
863 performance [121]. If one does not clearly define which is the *space* of artifacts being targeted, then any  
864 comparison among randomized algorithms is doomed to be arbitrary. For example, consider again the example  
865 of unit testing of object-oriented software. Assume that a case study involves 10 classes, and algorithm  $\mathcal{A}$  is  
866 statistically better on seven of them, whereas algorithm  $\mathcal{B}$  is statistically better on the other three. One could  
867 naively claim that algorithm  $\mathcal{A}$  is *on average* better than  $\mathcal{B}$ . But, maybe, those seven classes for which  $\mathcal{A}$  is  
868 better are all container classes, whereas the other three classes are related to string manipulations (e.g., [4]).  
869 If one had chosen for the case study more classes of this latter type, then the conclusions could be different  
870 (i.e.,  $\mathcal{B}$  would be considered *on average* better than  $\mathcal{A}$ ). Though the problem of choosing *appropriate* artifacts  
871 is intrinsically difficult, it is important for researchers to define their target artifacts as well as possible and  
872 carefully attempt to provide plausible reasons for differences in results across artifacts, such as classes, based  
873 on a thorough analysis of their characteristics.

874 Ideally, when realistic artifacts for a certain type of problems are difficult to find, one would like to be  
875 able to generate large numbers of them automatically in a realistic fashion. However, this requires that the  
876 artifacts have a clear and predictable structure, that there exist heuristics to generate correct and meaningful  
877 instances of such artifacts. If this is possible, one strong advantage is that one can control and vary interesting  
878 properties of the artifacts (e.g., class size, number of test cases) to enable interesting sensitivity analyses and  
879 assess the performance of randomized algorithms as a function of these properties. For example, in the work  
880 of Hemmati *et al.* [59], the authors analyzed different test suite reduction techniques for model-based testing  
881 of large systems. Obtaining real models from industry is difficult, and UML models of real systems are not  
882 common in open source repositories. Although the case study was based on two real industrial systems (e.g.,  
883 one provided by Cisco Systems), to cope with possible threats to external validity, the authors also used a large  
884 set of artificially generated test suites following some specific rules and a randomized construction algorithm.  
885 For example, the number of test cases in the test suites and the fault detection rate were varied in order to assess  
886 their impact on the effectiveness of the resulting selection technique. The aim was to do so while retaining as  
887 much as possible the realism of the test suites in the case studies. Such studies may be considered a type of  
888 simulation and may not generate fully realistic artifacts. But they may provide useful insights into the impact  
889 of some artifact properties on the effectiveness of a randomized algorithm.

890 For some types of software engineering problems, a large number of artifacts can be selected or generated  
891 (e.g., randomly selecting classes to investigate the unit testing of open source software). When evaluating  
892 randomized algorithms in this context one has to make the following decision: Assume a budget for experiments

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<sup>4</sup><http://sourceforge.net/>, accessed November 2011.

893  $b = n \times z$  for each algorithm, where  $n$  represents the times a randomized algorithm is run on each artifact, and  
 894  $z$  is the number of these artifacts. If one considers  $b$  to be fixed (e.g., depending on how long it takes to run  
 895  $b$  experiments), then a practical and important question is how to choose  $n$  and  $z$ ? Two extreme cases would  
 896 be  $(n = 1, z = b)$  and  $(n = b, z = 1)$ , but they would clearly lead to problems in terms of statistical testing  
 897 and external validity, respectively. Researchers have to strike a balance between two objectives: one wants to  
 898 analyze as many artifacts as possible to improve external validity and wishes, at the same time, to retain enough  
 899 runs (i.e.,  $n$ ) to check whether there is a statistically significant difference on any single artifact when applying  
 900 and comparing two randomized algorithms. This would, for obvious reasons, not be possible if  $n = 1$ . Though  
 901 in Section 8 it was suggested as a rule of thumb to use  $n = 1,000$  when possible, in certain circumstances  
 902 this may not be an option. If one has the possibility to analyze a large number  $z$  of artifacts but has practical  
 903 constraints regarding the number of experiments to be run (e.g., having experiments running on a PC for a  
 904 couple of years would not be very practical), then it may be more appropriate to execute less runs, perhaps as  
 905 low as  $n = 30$  or even  $n = 10$ . But going lower than such values would make the use of standard statistical  
 906 tests very difficult and, very likely, depending on the actual effect size and variance, would bring statistical  
 907 power to unacceptably low levels.

908 As discussed in Section 3, there are cases in the literature (e.g., [90, 118]) in which a random instance  
 909 generator is used, but then the algorithms are run only once (i.e.,  $n = 1$ ) on each artifact. For all the reasons  
 910 discussed in this section, in general one would prefer to have a higher number of runs even if that would lead  
 911 to use less artifacts. It is possible that there might be cases in which having  $n = 1$  could be preferable. At  
 912 any rate, in such cases it is recommended to properly clarify why the choice of using  $n = 1$  was made, and to  
 913 inform the reader of the possible validity threats related to statistical power and representativeness of the case  
 914 study.

## 915 10.2 Analysis of Multiple Artifacts

916 If for the addressed research question the considered artifacts can be considered representative of the target,  
 917 it is meaningful to then use statistical tests for evaluating whether algorithm  $\mathcal{A}$  is significantly better than  $\mathcal{B}$   
 918 on all selected artifact instances. However, as it will be shown below, which type of test is used is of the  
 919 highest importance. Using again the same example described before, assume six classes have been selected  
 920 for investigating the unit testing of object-oriented software. Each algorithm is run on each of these six classes  
 921  $n$  times (e.g.,  $n = 30$ ), and average values out of these runs are collected for each class. This makes up a  
 922 total of  $2 \times 6 \times 30 = 360$  runs. Assume that the algorithms are evaluated based on how many test cases they  
 923 generate before reaching full coverage. For the first algorithm, assume that a researcher obtains the following  
 924 average values  $X = \{10k, 20k, 30k, 40k, 50k, 60k\}$ , whereas for the second algorithm she obtains  $Y =$   
 925  $\{12k, 21k, 34k, 41k, 53k, 68k\}$ . The average values are ordered by problem instance where  $k = 1000$ , i.e., in  
 926  $X$ , out of  $n = 30$  runs on the first artifact the average number of test cases run equals 10,000. Further assume  
 927 that the problem instances are ordered by difficulty (i.e., solving the first problem is much easier than solving the  
 928 fifth, because on average it requires to generate/run less test cases). If one wants to evaluate whether there is any  
 929 statistical difference between  $X$  and  $Y$ , an *unpaired test*, such as Mann-Whitney U-test, would yield a  $p$ -value  
 930 equal to 0.699 (e.g., by using the  $R$  [97] command “`wilcox.test(X,Y)`”), thus suggesting the difference is not  
 931 statistically significant. However, this would be technically incorrect since different artifacts present different  
 932 levels of difficulty, and considering all data together at the same time would blur the relative performance of  
 933 the compared algorithms. In other words, a run of an inefficient algorithm on an *easy* problem would likely  
 934 result in a better value than a run of a more efficient algorithm that is run instead on a *difficult* problem. If the  
 935 case study involves artifacts of different levels of difficulty (as it is usually the case, either by design or due to  
 936 random sampling) then it might be challenging to detect any statistical difference with an unpaired test.

937 Alternatively, *paired tests* such as the Wilcoxon rank sum test can be used (e.g., “`wilcox.test(X,Y,paired=TRUE)`”  
 938 in  $R$  [97]). In a paired rank sum test, what is evaluated is whether the differences  $Z_i = Y_i - X_i$  are centered  
 939 around 0, i.e., the null hypothesis is  $Z = 0$ . In that example, it would be  $Z = \{2k, 1k, 4k, 1k, 3k, 8k\}$ , i.e.,  
 940 on average the second algorithm is always better than the first. A Wilcoxon rank sum test here yields  $p$ -value  
 941  $= 0.035$ , which suggests a statistically significant difference among the performance of the two algorithms, a  
 942 result in sharp contrast with the unpaired test results above. This highlights why it is extremely important to use  
 943 paired tests when comparing randomized algorithms on a set of selected artifacts. Another similar approach  
 944 would be to calculate the effect sizes and check whether they are symmetric around the null hypothesis. As-

945 sume for example that the resulting  $\hat{A}_{XY}$  effect sizes are equal to  $ES = \{0.4, 0.4, 0.4, 0.4, 0.4, 0.4\}$  (note,  
946 their actual values are not important as long as they are lower than 0.5). Then a test for symmetry in  $R$  would  
947 be “`wilcox.test(ES, mu=0.5)`”, which would result in a  $p$ -value equal to 0.019.

948 In the above example, the first algorithm is better in six out of six cases, which is a clear case. But typically  
949 results are not that consistent, and several of the compared algorithms may perform best on different artifacts.  
950 For example, assume a case study involving 100 artifacts: if an algorithm fares better on 51 of these, then the  
951 difference among the two would not be statistically significant when using a paired test. Using the example  
952 where an algorithm  $\mathcal{A}$  is better than another  $\mathcal{B}$  on some artifacts and worse on other artifacts, a paired rank sum  
953 test evaluates whether one algorithm is statistically better on a higher number of artifacts.

954 The above discussion on the use of appropriate statistical tests is incomplete as it considers the evaluation  
955 of a randomized algorithm as ternary, i.e., it is either better, equivalent or worse than another one. Consider the  
956 following example: algorithm  $\mathcal{A}$  is better on 60% of the case study, but only by a very limited amount (where  
957 such “better” is defined based on the effect size). On the other hand, on the other 40% of the case study, it  
958 is much worse than algorithm  $\mathcal{B}$ . In this case, blindly applying a paired Wilcoxon rank sum test would lead  
959 to the conclusion that  $\mathcal{A}$  is preferable, whereas a practitioner might prefer to use  $\mathcal{B}$ . Another option could  
960 be to collect standardized effect sizes for each problem instance, and then average them over all problems  
961 instances. This would provide additional information, but it would not solve the problem of fully describing  
962 the relative performance of two randomized algorithms, and would still be strongly dependent on the choice  
963 of the case study. Consider a case with five artifacts and the following  $\hat{A}_{12}$  measures  $\{0.6, 0.6, 0.6, 0.6, 0.1\}$ .  
964 One algorithm is better than the other on four artifacts ( $\hat{A}_{12} = 0.6$ ), but worse on the last one ( $\hat{A}_{12} = 0.1$ ).  
965 If one averages those values on the entire case study, he would obtain  $\hat{A}_{12} = 0.5$ , thus suggesting there is  
966 no difference among the two algorithms! This example illustrates the fact that aggregate statistics on a set of  
967 artifacts are useful to summarize the comparisons of two (or more) algorithms, but only as long as particular  
968 care is taken to handle cases where sharp differences can be observed among artifacts. In general, researchers  
969 should report the performance of the algorithms on each problem instance separately and attempt, as discussed  
970 above, to explain differences. One useful way to show the relative performance of randomized algorithms on a  
971 set of artifacts is to use box-plots of the effect sizes, especially when dealing with many artifacts

## 972 11 Practical Guidelines

973 Based on the above discussions, this section summarizes a set of practical guidelines for the use of statistical  
974 tests in experiments comparing randomized algorithms. Though one would expect exceptions, given the current  
975 state of practice (see Section 3 and the systematic reviews of Ali *et al.* [3] and Kampenes *et al.* [63]), the authors  
976 of this paper believe that it is important to provide practical guidance that will be valid in most cases and enable  
977 higher quality studies to be reported. It is recommendable that practitioners follow these guidelines and justify  
978 any necessary deviation.

979 There are many statistical tools that are available. In the following, all the examples will be provided based  
980 on  $R$  [97], because it is a powerful tool that is freely available and supported by many statisticians. But any  
981 other professional tool would provide similar capabilities.

982 Practical guidelines are summarized below. Notice that often, for reasons of space, it is not possible to  
983 report all the data of the statistical tests. Based on the circumstances, authors need to make careful choices on  
984 what to report.

- 985 • When randomized algorithms are analyzed, clearly specify the number of runs and employed statistical  
986 tests. For example, they can be summarized in a threats to validity section, in which how randomness has  
987 been taken into account should be discussed and justified.
- 988 • On each artifact in the case study, run each randomized algorithm at least  $n = 1,000$  times. If this is not  
989 possible, explain the reasons and report the total amount of time it took to run the entire case study. If for  
990 example 30 runs were performed and the total execution time was just one hour, then it is rather difficult  
991 to justify why a higher number of runs was not used to gain statistical power, lower  $p$ -values, and narrow  
992 the confidence interval of effect size estimates (Section 8).
- 993 • When a large number of artifacts can be used in the case study (e.g., for unit testing of open source  
994 software) but there are constraints in terms of execution time, then it is advisable to execute less runs



995 per artifact (though at least  $n = 10$ ) and use more artifacts (rather than having  $n = 1,000$  but only few  
996 artifacts, see Section 10.1). The objective is to strike a balance between generalization and statistical  
997 power.

- 998 • The choice of artifacts, to which randomized algorithms are applied, has a large impact on the validity  
999 and statistical interpretation of the final results (Section 10.1). Ideally, a large unbiased selection of  
1000 artifacts that are representative of the addressed problem should be used as case study. Even if obtaining  
1001 such artifacts is usually not possible, it is important to always clarify how they were chosen. The aim is  
1002 to allow the reader to properly interpret the results of the statistical analyses when more than one artifact  
1003 is used in a case study.
- 1004 • For detecting statistical differences, use the two-tailed non-parametric Mann-Whitney U-test for interval-  
1005 scale results and the Fisher exact test for dichotomous results (i.e., in the cases of censored data as  
1006 discussed in Section 6). For the former case, in *R* you can use the function “w=wilcox.test(X,Y)” where  
1007 *X* and *Y* are the data sets with the observations of the two compared randomized algorithms. If you  
1008 are comparing a randomized algorithm against a deterministic one, use the one-sample version of the  
1009 test with “w=wilcox.test(X,mu=D)”, where *D* is the resulting performance measure for the deterministic  
1010 algorithm. When there are *a* successes for the first algorithm and *b* successes for the second, one should  
1011 use “f=fisher.test(m)”, where *m* is a matrix derived in this way: “m=matrix(c(a,n-a,b,n-b),2,2)”.
- 1012 • Report all the obtained *p*-values, whether they are smaller than  $\alpha$  or not, and not just whether differences  
1013 are significant. The motivation is for the reader to choose the level of risk that is suitable in her application  
1014 context. When reporting all *p*-values is not possible (e.g., due to space reasons), one could report the  
1015 proportion of significant test results: “*x* out of *y* tests were significant at  $\alpha$  level . . .”.
- 1016 • Always report standardized effect size measures. For dichotomous results, the odds ratio  $\psi$  can be cal-  
1017 culated using Equation 2, where for example  $\rho = 0.5$  (used to address zero occurrence cases [55]). For  
1018 interval-scale results and the  $\hat{A}_{12}$  effect size, the rank sum  $R_1$  used in Equation 1 can be calculated with  
1019 “R1=sum(rank(c(X,Y))[seq\_along(X)])”. It is also strongly advised to report effect size confidence inter-  
1020 vals, e.g., by using a bootstrapping technique. In *R*, there is library *boot* from which the function “boot”  
1021 (to do the sampling) and “boot.ci” (to create a confidence interval) can be used. A confidence interval  
1022 is much easier to use than *p*-values for decision making as potential benefits can be compared to costs  
1023 while accounting for uncertainty.
- 1024 • To help the meta-analyses of published results across studies, report means and standard deviations (in  
1025 case readers for some reasons want to calculate effect sizes in the *d* family). For dichotomous experi-  
1026 ments, always report the values *a* and *b* (so that other types of effect sizes can be computed [55]).
- 1027 • If space permits, provide full statistics for the collected data, as for example mean, median, variance,  
1028 min/max values, skewness, kurtosis and median absolute deviation. Box-plots are also useful to visualize  
1029 them.
- 1030 • When analyzing more than two randomized algorithms, use pairwise comparisons including pairwise  
1031 statistical tests and effect size measures. If the case study can be considered as a statistically valid  
1032 sample, then you can also use a test for symmetry on the null hypothesis for the effect sizes (Sec-  
1033 tion 10.2). For example, if *ES* contains the  $\hat{A}_{12}$  effect sizes for each artifact in the case study, then  
1034 “w=wilcox.test(ES,mu=0.5)” will tell whether one algorithm is better on a *higher number* of artifacts  
1035 (but this would not take into account the *magnitude* of the improvement).
- 1036 • If space permits, state the employed statistical tool and how it was used (there can be subtle differences  
1037 on how the tests are computed).

## 1038 12 Threats to Validity

1039 The systematic review in Section 3 is based on only four sources, from which only 54 out of 246 papers  
1040 were selected. Although this systematic review is larger than the majority of systematic reviews in software

1041 engineering [70], accounting for more sources of information might lead to different results. One can, however,  
1042 safely argue that TSE and ICSE are representative of research trends in software engineering. Furthermore,  
1043 that review is only used as a motivation for providing practical guidelines, and its results are in line with other  
1044 larger systematic reviews [3, 63]. Last, papers sometimes lack precision and interpretation errors are always  
1045 possible.

1046 As already discussed in Section 11, the practical guidelines provided in this paper may not be applicable  
1047 to all contexts. Therefore, in every specific context, one should always carefully assess them. For some spe-  
1048 cific cases, other statistical procedures could be preferable, especially when only few runs of the randomized  
1049 algorithms are possible.

## 1050 **13 Conclusion**

1051 Randomized algorithms (e.g., Genetic Algorithms) are widely used to address many software engineering prob-  
1052 lems, such as test case selection. In this paper, as a first contribution, a systematic review is performed to  
1053 evaluate how the results of randomized algorithms in software engineering are analyzed.

1054 Similar to previous systematic reviews on related topics [3, 63], this review shows that most of the published  
1055 results regarding the use of randomized algorithms in software engineering are missing rigorous statistical  
1056 analyses to support the validity of their conclusions.

1057 To cope with this problem, this paper provides, discusses, and justifies a set of *practical* guidelines targeting  
1058 researchers in software engineering. In contrast to other guidelines in the literature for experimental software  
1059 engineering [120] and other scientific fields (e.g., [89, 64]), the guidelines in this paper are tailored to the  
1060 specific properties of randomized algorithms when applied to software engineering problems, with a particular  
1061 focus on software verification and validation. The use of these guidelines is important in order to develop a  
1062 reliable body of empirical results over time, by enabling comparisons across studies so as to converge towards  
1063 generalizable results of practical importance. Otherwise, as in many other aspects of software engineering,  
1064 unreliable results will prevent effective technology transfer and will inevitably limit the impact of research on  
1065 practice.

1066 Note that there are advanced topics in statistics that have not been discussed in this paper, as for example  
1067 Bayesian data analysis [47]. This paper is not meant to be a complete and ultimate reference for experimenters  
1068 in software engineering, but rather be an essential guide to help them to use fundamental and common statistical  
1069 methods in an appropriate manner.

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