# ESAMPLER: Boosting Sampling of Satisfying Assignments for Boolean Formulas via Derivation

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#### Abstract

Boolean satisfiability (SAT) plays a key role in diverse areas such as spanning planning, inference, data mining, testing and optimization. Apart from the classical problem of checking Boolean satisfiability, generating random satisfying assignments has attracted significant theoretical and practical interests over the past years. In practical applications, usually a large number of satisfying assignments for a given Boolean formula are needed, the generation of which turns out to be a computational hard problem in both theory and practice. In this work, we propose a novel approach to derive a large set of satisfying assignments from a given one in an efficient way. Our approach is based on an insight that flipping the truth values of properly chosen variables of a satisfying assignment could result in satisfying assignments without invoking computationally expensive SAT solving. We propose a derivation algorithm to discover such variables for each given satisfying assignment. Our approach is orthogonal to the previous techniques for generating satisfying assignments and could be integrated into the existing SAT samplers. We implement our approach as an open-source tool ESAMPLER using two representative state-of-the-art samplers (QUICKSAMPLER and UNIGEN3) as the underlying satisfying assignment generation engine. We conduct extensive experiments on various publicly available benchmarks and apply ESAMPLER to solve Bayesian inference. The results show that ESAMPLER can efficiently boost the sampling of satisfying assignments of both QUICKSAMPLER and UNIGEN3 on a large portion of the benchmarks and is at least comparable on the others. ESAMPLER performs considerably better than QuickSampler and UniGen3, as well as another state-of-the-art sampler SearchTreeSampler.

Keywords: Boolean satisfiability, Constraint-based sampling, SAT solving

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#### 1. Introduction

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Boolean satisfiability, also known as SAT, concerns determining whether a given Boolean formula is satisfiable. There have been strong theoretical and practical interests in the SAT problem, which has played a key role in diverse areas spanning planning, inferencing, data mining, testing and optimization [1, 2]. Apart from the classical problem of checking Boolean satisfiability, generating random satisfying assignments has attracted significant theoretical and practical interests over the years [3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15]. In several practical applications, a large number of satisfying assignments for a given Boolean formula are needed. For instance, simulation-based verification 10 is a commonly adopted technique to test hardware design. In this scenario, the simulated behavior is compared with the expected behavior where any mismatch 12 is flagged as an indication of a bug [12, 13]. It is a common practice to generate 13 a large number of stimuli satisfying a given set of constraints in the form of 14 Boolean formulas. These constraints typically arise from various sources such 15 as application-specific knowledge and environmental requirements. Another application scenario is the generation of adversarial examples for adversarial train-17 ing [16, 17]. Adversarial training is a widely adopted technique to improve the 18 robustness of neural networks against adversarial attacks where a large number 19 of adversarial inputs (e.g., images) would be generated explicitly or implicitly. 20 For instance, to adversarially train a binarized neural network [18, 19], adver-21 sarial images could be generated by encoding a binarized neural network as a 22 Boolean formula based on which satisfying assignments are sampled [20, 21]. 23

Sampling satisfying assignments for a given Boolean formula is, however, challenging. Cook has shown in 1971 that the SAT problem is **NP**-complete [22]. In recent years, we have seen a tremendous progress in SAT solving, supported by techniques such as conflict-driven clause learning (CDCL [23, 24, 25]), yielding powerful solvers such as CryptoMiniSAT [26]. However, generating a large number of satisfying assignments is still computationally prohibitive and often infeasible in practical settings [27, 28].

In this work, we develop ESAMPLER, aiming for boosting the generation of a large number of satisfying assignments efficiently for a given Boolean formula. The general strategy is to use an existing sampler to produce a seed sample as a satisfying assignment, from which we derive more satisfying assignments by flipping some variables of the given Boolean formula. Clearly, naively flipping variables may yield unsatisfying assignments. To tackle this problem, we propose a novel derivation procedure which explores the semantics of the Boolean formula under the seed sample, so that the resulting assignments can be guaranteed to satisfy the Boolean formula. The advantage of our approach lies in that it can be integrated with the existing SAT samplers, so would enjoy considerably wider applicability.

To demonstrate our approach, we implement a sampler ESAMPLER based on the two types of state-of-the-art sampler QUICKSAMPLER [28] and UNI-GEN3 [29]. We carry out extensive experiments on the publicly available benchmarks from UNIGEN [30] which include hundreds of Boolean formulas from

real-world testing and verification applications, and apply ESAMPLER to solve
Bayesian inference of the plan recognition problems [31]. Our experimental results show that ESAMPLER is able to effectively boost the sampling of satisfying
assignments of both QUICKSAMPLER and UNIGEN3 on a large portion of the
benchmarks and is at least comparable on the others. Consequently, ESAMPLER
considerably performs better than QUICKSAMPLER and UNIGEN3, as well as
the another state-of-the-art sampler SEARCHTREESAMPLER (STS in short) [32].
The experimental results confirm the efficacy of our derivation approach.

Our main contributions can be summarized as follows.

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- We introduce a novel approach for deriving a large set of satisfying assignments from a given seed. It is generic and could be integrated with the existing samplers. To the best of our knowledge, it is the first work to generate satisfying assignments from a given seed.
- We implement an integrated sampler ESAMPLER based on two state-ofthe-art samplers. Our tool is available at https://github.com/ESampler/ Esampler.
- We conduct extensive experiments on hundreds of real-world benchmarks. The results show that our derivation approach is effective and consequently ESAMPLER performs considerably better than the three state-of-the-art samplers QuickSampler, STS and UniGen3.

**Related Work.** Various techniques have been proposed to tackle the problem 66 of the satisfying assignment generation for Boolean formulas [33]. Binary decision diagrams (BDD) and Markov Chain Monte Carlo (MCMC) algorithms 68 such as simulated annealing and Metropolis-Hastings are widely used for generating satisfying assignments [9, 34, 35]. These techniques usually provide 70 theoretical guarantees of uniformity but are limited in scalability and efficiency. Therefore, heuristics are proposed to speed up at the cost of theoretical guar-72 antees of uniformity [36, 37, 34]. Another class of satisfying assignment generation techniques with theoretical guarantees of uniformity is based on hash-74 ing [38, 39, 40, 41, 42, 30, 43, 29]. Hashing-based techniques add hash functions 75 (e.g., XOR of a random subset of variables) to the Boolean formula in order 76 to partition the search space uniformly and then randomly pick a satisfying 77 assignment from a randomly chosen cell. These algorithms are also limited in scalability and efficiency. In comparison, our approach primarily aims for effi-79 ciency, using fewer solver calls to generate a large number of solutions. We also ຂດ provide a parameter to balance the uniformity of the generated samples and the 81 efficiency of the procedure. Although we do not provide a theoretical guarantee of uniformity, the experimental results demonstrate that our approach is able 83 to produce solutions nearly uniformly when the maximal number of solutions per seed is set in a reasonable range. 85

SAT samplers aiming to quickly generate a large number of assignments have recently been proposed. Both QUICKSAMPLER [28] and STS [32] share

the same goal as our work, namely, fast generation of a larger number of assignments. QuickSampler works as follows. Given a Boolean formula Φ, it first constructs a random assignment v and then uses the MaxSAT solver [44] 90 to solve the MaxSAT problem with the hard constraint  $\Phi$  and soft constraint 91  $\Psi$ , where  $\Psi$  is the conjunction of literals x if v(x) = 1 or  $\neg x$  if v(x) = 0. Solving 92 the MaxSAT problem yields a satisfying assignment v' of  $\Phi$  that is close to the random assignment v. After that, QUICKSAMPLER iteratively flips the value 94 of each variable x in the satisfying assignment v' to find another close satisfy-95 ing assignment  $v'_x$  using the MaxSAT solver, where the soft constraint asserts 96 the satisfying assignment v' except for the flipped variable x, and the original Boolean formula together with the flipped variable is used as hard constraint. For each flipped variable x, the difference  $\delta_x$  between two satisfying assignments 99 v' and  $v'_x$  is computed. All such differences are combined and applied to mu-100 tate the satisfying assignment v' to generate a large number of assignments. 101 However, the assignments generated by QUICKSAMPLER may not satisfy the 102 Boolean formula, hence follow-up checkings are needed. In contrast, our ap-103 proach only mutates proper variables by which the formula is guaranteed to 104 be satisfied. STS explores the tree of variable assignments in a breadth-first 105 way with the MiniSat SAT solver [45] as an oracle. During this procedure, it generates pseudosolutions, which are partial assignments to the variables that 107 can be completed to full satisfying assignments. However, it has to invoke SAT solvers multiple times during the breadth-first exploration. In contrast, ESAM-109 PLER does not require SAT solving when generating satisfying assignments from 110 a seed. 111

Technically, our derivation procedure aims to generate a large set of satisfying assignments from a given seed, and is orthogonal to the existing SAT samplers. It can be integrated into the existing samplers to improve their efficiency as we demonstrated using QUICKSAMPLER and UNIGEN3.

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Sampling satisfying assignments is also closely related to the model-counting problem which counts the number of satisfying assignments for a Boolean formula. Model-counting techniques have been used for sampling satisfying assignments (e.g., SPUR [46]) while satisfying assignment sampling techniques can also be used for model-counting (e.g., STS [32] and APPROXCOUNT [35]).

This article is an extended version of [47], but with substantial new material. In particular, we apply ESAMPLER to boost another uniform sampler UNIGEN3 and carry out more experiments (cf. Section 5.4), which show the generality and wide applicability of ESAMPLER to diverse seed generation samplers. We also apply ESAMPLER for inference of Bayesian networks and report experimental results on the real-world plan recognition problems (cf. Section 6), showing a significant improvement of our approach ESAMPLER over the samplers QUICK-SAMPLER, STS and UNIGEN3.

**Outline.** The remainder of this paper is organized as follows. In Section 2, we briefly revisit related concepts of Boolean formulas. We present our derivation procedure in Section 3, and show how to integrate it into existing SAT samplers in Section 4. We report evaluation results in Section 5. We apply ESAMPLER

#### 2. Preliminaries

We first recap some basic notions and notations which are used in this work.

**Boolean formulas**. Let us fix a set of Boolean variables  $\mathcal{V}$ . A literal l is either a Boolean variable  $x \in \mathcal{V}$  or its negation  $\neg x$ . We denote by var(l) the variable x used in the literal l, namely,  $var(x) = var(\neg x) = x$ .

A Boolean formula  $\Phi$  is a Boolean combination of literals using logical-AND  $(\land)$  and logical-OR  $(\lor)$  operators. As a convention, we assume that Boolean formulas are given in the conjunctive normal form (CNF)  $\bigwedge_{j=1}^{m}\bigvee_{i=1}^{n_{j}}l_{i}^{j}$ , where for each  $1 \leq j \leq m$  and  $1 \leq i \leq n_{j}$ ,  $l_{i}^{j}$  is a literal, and  $\bigvee_{i=1}^{n_{j}}l_{i}^{j}$  is referred to a clause for each  $1 \leq j \leq m$ . Given a Boolean formula  $\Phi$  and a literal l, let  $\Phi_{l}$  denote the set of clauses that contain the literal l. For each clause  $\phi = \bigvee_{i=1}^{n_{j}}l_{i}^{j}$ , we assume that all literals in  $\phi$  are distinct, and denote by  $|\phi|$  the number  $n_{j}$  of literals in the clause  $\phi$ .

Assignments. An assignment is a function  $v: \mathcal{V} \to \{0,1\}$  which assigns a Boolean value to each Boolean variable  $x \in \mathcal{V}$ . Given a Boolean formula  $\Phi$  and an assignment v, v is a satisfying assignment of  $\Phi$ , denoted by  $v \models \Phi$ , if the Boolean formula  $\Phi$  evaluates to 1 under the assignment v. A partial assignment is a partial function  $v: \mathcal{V} \to \{0,1\}$  such that for each  $x \in \mathcal{V}$ , v(x) is a Boolean value if x is defined in v, otherwise x is undefined in v.

For each assignment v, variable  $x \in \mathcal{V}$  and value  $i \in \{0, 1\}$ , we denote by  $v[x \mapsto i]$  the assignment that agrees with v except for the variable x, i.e., for each variable  $y \in \mathcal{V}$ ,

$$v[x \mapsto i](y) = \begin{cases} v(y), & \text{if } y \neq x; \\ i, & \text{otherwise.} \end{cases}$$

Satisfiability and maximum satisfiability. Given a Boolean formula  $\Phi$ , the satisfiability problem (SAT) is to determine whether a satisfying assignment of  $\Phi$  exists or not. If  $\Phi$  is satisfied, then a solution is produced as a witness. It is well-known that the SAT problem is **NP**-complete [22].

Given a pair of Boolean formulas  $(\Phi, \Psi)$ , the maximum satisfiability problem (MaxSAT) is to find a satisfying assignment that satisfies the Boolean formula  $\Phi$  and meanwhile maximizes the number of satisfied clauses in  $\Psi$ . The clauses in  $\Phi$  are usually called hard constraints, while the clauses in  $\Psi$  are called soft constraints. It is easy to see that the MaxSAT problem is at least **NP**-hard and can be solved by the state-of-the-art solvers such as Z3 [44].

In this work, by solvers we mean tools that are able to produce one satisfying assignment of the (Max)SAT problem whilst by samplers we mean those that are able to generate more than one satisfying assignments.

**Independent support.** Given a Boolean formula  $\Phi$ , an *independent support* Supp of  $\Phi$  [30], is a set of variables such that for each pair of satisfying assignments (v, v') of  $\Phi$ , if v(x) = v'(x) holds for all variables  $x \in \text{Supp}$ , then

v(y) = v'(y) holds for all variables  $y \in \mathcal{V} \setminus \text{Supp}$ . Intuitively, the truth values of the independent support  $\text{Supp}_{\Phi}$  uniquely determine the truth values of the other variables. In other words, flipping the truth value of any variable  $y \in \mathcal{V} \setminus \text{Supp}$  in the satisfying assignment v only will make the resulting assignment  $v[y \mapsto \neg v(y)]$  fail to satisfy  $\Phi$ .

It is easy to see that any superset of an independent support of  $\Phi$  is also an independent support. There are tools, such as MIS and SMIS [48], that are able to compute minimal and minimum independent supports for Boolean formulas, where *minimal* means removing any variable from the independent support X will lead to a non-independent support, and *minimum* means there does not exist any independent support whose size is smaller. Remark that the problem of deciding whether a set of variables is a minimal independent support of a Boolean formula  $\Phi$  is  $\mathbf{DP}$ -complete [49], where  $\mathbf{DP} := \{A - B \mid A, B \in \mathbf{NP}\}$ .

#### 3. Derivation Procedure

In this section, we first present a motivating example which exemplifies the key insight behind our approach for efficiently generating a large number of satisfying assignments. We then provide a derivation procedure which is able to derive more satisfying assignments from a seed by flipping the truth values of properly chosen variables without invoking computationally expensive SAT solving. The derivation procedure is the basis for efficiently generating a large number of satisfying assignments, and can be integrated into other samplers.

#### 3.1. Motivating Example

To exemplify the key insight behind our approach, let us consider the following Boolean formula

$$\Phi_e \equiv (\neg a \lor b \lor c) \land (a \lor \neg c \lor \neg d) \land (\neg b \lor c) \land (b \lor d).$$

Suppose we have already obtained one satisfying assignment v (called seed) of  $\Phi_e$  with v(a) = v(b) = v(c) = v(d) = 1. We can observe that the clause  $\neg a \lor b \lor c$  (resp.  $b \lor d$ ) contains two literals b and c (resp. b and d) whose values are 1 under the assignment v. Moreover, the common literal b does not appear in the other clauses, namely,  $a \lor \neg c \lor \neg d$  and  $\neg b \lor c$ . By flipping the value v(b) of the variable b in the assignment v, we can obtain a new assignment  $v[b \mapsto \neg v(b)]$ , which is also a satisfying assignment of  $\Phi_e$ .

However, by flipping the value v(c) of the variable c in the assignment v, the new assignment  $v[c \mapsto \neg v(c)]$  is not a satisfying assignment of  $\Phi_e$ . This is because the clause  $\neg b \lor c$  contains only one literal c whose value is 1 under the assignment v. After flipping the value v(c) of the variable c in the assignment v, the clause  $\neg b \lor c$  is no more satisfied.

This simple observation suggests that, for a seed v, we may identify proper variables (such as b but not c in the above example) so that when the value of one such variable is flipped it is still a satisfying assignment. Furthermore, the

## Algorithm 1 Deriving satisfying assignments from a seed

```
1: procedure DERIVATION(\Phi, v, MaxNum, Supp)
         Derived = \{v\};
 2:
 3:
         Queue = [v]:
         while Queue \neq \emptyset \land | \texttt{Derived} | \leq \texttt{MaxNum do}
 4:
             v = Queue.DEQUEUE();
 5:
             L = \{x \mid v(x) = 1\} \cup \{\neg x \mid v(x) = 0\};
 6:
             for all l \in L \land var(l) \in Supp do
 7:
                  if \forall \bigvee_{i=1}^{m} l_i \in \Phi_l, \exists i. (1 \leq i \leq m \land l \neq l_i \land l_i \in L) then
 8:
                       x = var(l);
 9:
                       v' = v[x \mapsto \neg v(x)];
10:
                       if v' \not\in \texttt{Derived then}
11:
                           Derived = Derived \cup \{v'\};
12:
                           Queue. ENQUEUE (v');
13:
14:
                       end if
                  end if
15:
             end for
16:
         end while
17:
         return Derived:
18:
19: end procedure
```

new satisfying assignments can be used as seeds to derive more satisfying assignments. This often allows generation of a larger number of satisfying assignments without invoking computationally expensive SAT solving.

## 3.2. Derivation Algorithm

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In this subsection, we present a derivation procedure for deriving new satisfying assignments from a given seed. Given a Boolean formula  $\Phi$ , a seed v, an independent support Supp of  $\Phi$ , and the maximal number MaxNum of expected satisfying assignments, the procedure Derivation in Algorithm 1 iteratively derives new satisfying assignments from the seed v until no new satisfying assignment can be found or the number of generated satisfying assignments hits the threshold MaxNum. It returns the set of generated satisfying assignments including the original seed v.

To start, Algorithm 1 initializes the set Derived for recording all the generated satisfying assignments (Line 2) and the queue Queue for storing the seeds (Line 3). It then repeats the following procedure until no new satisfying assignments can be found or the number of the generated satisfying assignments exceeds the threshold MaxNum (While-loop).

For each seed v in Queue (Line 5), it first identifies all the literals whose value is 1 under the assignment v (Line 6). After that, for each literal  $l \in L$  whose variable  $\operatorname{var}(l) \in \operatorname{Supp}$  (Line 7), it checks, for each clause  $\bigvee_{i=j}^m l_j$  that contains the literal l (i.e.,  $\bigvee_{i=j}^m l_j \in \Phi_l$ ), whether  $\bigvee_{i=j}^m l_j$  contains a distinct literal  $l_i$  whose value is also 1, i.e.,  $l_i \in L$  (Line 8). If this is the case, we can

deduce that the assignment  $v[x \mapsto \neg v(x)]$  obtained from the assignment v by flipping the variable x = var(l) is also a satisfying assignment of  $\Phi$ . Therefore, we extract the variable x from the literal l (Line 9) and construct the assignment  $v' = v[x \mapsto \neg v(x)]$  (Line 10). If the assignment v' has not been generated before, it is inserted to Derived and Queue (Lines 12 and 13).

One may notice that only variables in Supp are considered for flipping (Line 7). In general, we can take all the variables into account for flipping. However, as mentioned before (cf. Section 2), flipping variables outside of Supp will definitely lead to unsatisfying assignments. Therefore, it suffices to consider variables from Supp for flipping. Due to this, the values of each variable outside of Supp are the same in all the generated satisfying assignments from a given seed.

We remark that the derivation procedure DERIVATION could alternatively be presented as a recursive procedure which invokes itself when a new satisfying assignment is generated, or equivalently, use a stack rather than a queue to store the generated seeds. Intuitively, using the queue Queue to store the seeds, our algorithm works in a breadth-first fashion, while the other two ways would follow a depth-first fashion. We adopt the current way because it is more efficient than the other two ways.

**Theorem 1.** Given a Boolean formula  $\Phi$ , a seed v and an independent support Supp of  $\Phi$ , the set Derived returned by Algorithm 1 contains only satisfying assignments of  $\Phi$ . Moreover, these assignments agree on the variables outside of Supp.

PROOF. We show that the set Derived returned by Algorithm 1 contains only satisfying assignments of  $\Phi$  by applying induction on the sequence  $v_0v_1\cdots$  of the assignments added into Derived. The base case is trivial as the seed  $v_0$  satisfies the Boolean formula  $\Phi$ . We prove the inductive step below.

Suppose  $v_0, v_1 \cdots v_{k-1}$  have been added into the set Derived and the inductive step adds the assignment  $v_k$  into the set Derived. Then,  $v_k$  must be added due to one v of the previously added satisfying assignments  $v_0, v_1 \cdots v_{k-1}$ . There necessarily exists a literal l such that x = var(l) and  $v_k = v[x \mapsto \neg v(x)]$ .

To show that  $v_k$  satisfies  $\Phi$ , it is sufficient to prove that  $v_k$  satisfies all the clauses of  $\Phi$ . Let us consider a clause  $\bigvee_{i=j}^{m} l_j$  of  $\Phi$ ,

- If  $\bigvee_{i=j}^{m} l_j$  does not contain the literal l, then by applying induction hypothesis, v satisfies the Boolean formula  $\Phi$  and hence v satisfies the clause  $\bigvee_{i=j}^{m} l_j$ . Since  $v_k = v[x \mapsto \neg v(x)]$  and x = var(l), the truth of the clause  $\bigvee_{i=j}^{m} l_j$  does not change when the value of x in v is flipped. Therefore, we get that the assignment  $v_k$  satisfies the clause  $\bigvee_{i=j}^{m} l_j$ .
- If  $\bigvee_{i=1}^m l_i$  contains the literal l, then there exists another literal  $l_i \in \{l_1, \dots, l_m\}$  such that  $l_i \neq l$  and  $l_i \in L = \{x \mid v(x) = 1\} \cup \{\neg x \mid v(x) = 0\}$ . From  $l_i \in L = \{x \mid v(x) = 1\} \cup \{\neg x \mid v(x) = 0\}$ , we deduce that the literal  $l_i$ , hence the clause  $\bigvee_{i=1}^m l_i$ , holds under the assignment  $v_k$ .

```
(\neg a \lor b \lor c) \land (a \lor \neg c \lor \neg d) \land
\Phi_e:
                                                                           (\neg b \lor c)
                                                                                                      (b \vee d)
             (0 \lor 1 \lor 1)
                                           (1 \lor 0 \lor 0)
                                    \wedge
                                                                              (0 \lor 1)
                                                                                                      (1 \lor 1)
v_1:
             flip b and d respectively \downarrow
                                            (1 \lor 0 \lor
                                                              0)
             (0 \lor 0 \lor 1)
                                    Λ
                                                                                                      (0 \lor 1)
v_2:
                                                                              (1 \lor 1)
                                                                              (0 \lor 1)
v_3:
              (0 \lor 1 \lor 1)
                                            (1 \lor 0 \lor
                                                              1)
                                                                      Λ
                                                      flip a \downarrow \downarrow
             (1 \lor 1 \lor 1)
                                            (\mathbf{0} \vee \mathbf{0} \vee
                                                              1)
                                                                              (0 \lor 1)
v_4:
                                                                      \wedge
```

Figure 1: Derivation steps of the motivating example

**Example 1.** Recall the motivating example  $\Phi_e$ . Suppose the input seed is  $v_1$  with  $v_1(a) = v_1(b) = v_1(c) = v_1(d) = 1$  and the independent support  $Supp = \{a, b, d\}$ . The derivation steps are shown in Figure 1. At the beginning of the first iteration of the while-loop,  $v = v_1$  and  $L = \{a, b, c, d\}$ .

- 1. Suppose the variable a is chosen for flipping (Line 7). Since the clause  $a \lor \neg c \lor \neg d$  does not have any literals other than a that occur in L, Algorithm 1 will not flip the variable a.
- Next, the variable b is chosen for flipping (Line 7). Since the clause ¬a ∨ b ∨ c contains the literal c, the clause b ∨ d contains the literal d, and both literals c and d occur in L, Algorithm 1 will flip the variable b (Line 9) and produce a new satisfying assignment v<sub>2</sub> = v<sub>1</sub>[b → 0] (Line 10).
- 3. Finally, the variable d is chosen for flipping (Line 7). Since the clause  $b \lor d$  contains literal b that occurs in L, Algorithm 1 will flip the variable d (Line 9) and produce a new satisfying assignment  $v_3 = v_1[d \mapsto 0]$  (Line 10).

At the end of the first iteration of the while-loop,  $\operatorname{Derived} = \{v_1, v_2, v_3\}$  and  $\operatorname{Queue} = [v_2, v_3]$ . After entering the second iteration of the while-loop,  $v = v_2$ ,  $\operatorname{Queue}$  (resp. L) becomes  $[v_3]$  (resp.  $\{a, \neg b, c, d\}$ ). By applying similar steps as above, the satisfying assignment  $v_2$  is regenerated but will not be inserted to  $\operatorname{Derived}$  or  $\operatorname{Queue}$ .

At the end of the second iteration of the while-loop,  $Derived = \{v_1, v_2, v_3\}$  and  $Queue = [v_3]$ . After entering the third iteration of the while-loop,  $v = v_3$ ,  $Queue \ (resp. \ L) \ becomes \ \emptyset \ (resp. \ \{a,b,c,\neg d\})$ . By applying similar steps as above, Algorithm 1 will flip the variable a and produce a new satisfying assignment  $v_4 = v_3[a \mapsto 0]$ . In the end, no more new satisfying assignments can be generated and Algorithm 1 returns the set  $\{v_1, v_2, v_3, v_4\}$ .

#### 4. ESAMPLER

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In this section, we show that our derivation procedure is of generic nature in the sense that it can be integrated with other samplers. The basic idea is to generate seeds by invoking an existing sampler as an iterator, which returns a

## Algorithm 2 Integrated our derivation procedure into an existing sampler

```
1: procedure INTEGRATEDSAMPLER(Sampler,Φ,T,MaxPerSeed,Supp,RT,DT)
       Solutions = \emptyset:
 2:
 3:
       Derivable = false:
       Round = 0;
 4:
       Iterator = Sampler(\Phi, Supp);
 5:
 6:
       repeat
          v = Iterator.next();
 7:
          if v == \text{Null then}
 8:
              break:
 9:
          end if
10:
          if v \in Solutions then
11:
              continue;
12:
          end if
13:
14:
          if Derivable == true \lor Round < RT then
              Derived = Derivation(\Phi, v, MaxPerSeed, Supp);
15:
              Solutions = Solutions \cup Derived;
16:
              if |Derived| > DT then
17:
                 Derivable = true:
18:
19:
              else
20:
                 Round = Round + 1:
              end if
21:
22:
          else
              Solutions = Solutions \cup \{v\};
23:
          end if
24:
       until T is satisfied
25:
       return Solutions;
26:
27: end procedure
```

unique satisfying assignment each time. For each seed, we derive more satisfying assignments by invoking our derivation procedure. However, our derivation procedure may not be effective on some Boolean formulas. Therefore, we propose a heuristic to determine whether our derivation procedure is able to derive a large number of satisfying assignments or not. If it can derive a large number of satisfying assignments, we apply the derivation procedure for each satisfying assignment generated by the sampler, otherwise we disable it.

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Our idea is formalized as the procedure INTEGRATEDSAMPLER in Algorithm 2, which takes, as input, an off-the-shelf sampler Sampler, a Boolean formula  $\Phi$ , a threshold T as the termination condition, the maximum number MaxPerSeed of satisfying assignments per seed, an independent support Supp of the Boolean formula  $\Phi$ , two thresholds RT and DT to determine whether our derivation procedure is able to derive a large number of satisfying assignments, and returns a set Solutions of satisfying assignments of the formula  $\Phi$ .

The procedure Integrated Sampler first initializes the set Solutions, the

Boolean flag Derivable, the counter Round and the iterator Iterator of the sampler using the independent support Supp and Boolean formula  $\Phi$  (Lines 2– 5), where the Boolean flag Derivable and counter Round are used to determine if our derivation procedure is able to derive a large number of satisfying assignments. Then, it repeats the following procedure until the threshold T is hit.

During each iteration, INTEGRATED SAMPLER first invokes the iterator to get a satisfying assignment v, where v is Null if  $\Phi$  is unsatisfiable or the iterator cannot find new satisfying assignments. If v is Null, it breaks the loop (Line 9). If v already exists in Solutions, it skips this loop (Line 12). Otherwise it checks if the Boolean flag Derivable is true or the number Round of iterations is less than the threshold RT.

- If neither holds, the derivation procedure is considered to be not able to derive a large number of satisfying assignments and will be skipped;
- Otherwise, the derivation procedure is invoked to generate more satisfying assignments which are added to the set Solutions (Lines 15–16). If the number of satisfying assignments generated by the derivation procedure exceeds the threshold DT, we consider that the derivation procedure is able to derive a large number of satisfying assignments and set the Boolean flag Derivable to true (Line 18). Otherwise, we increase the counter Round by one. In general, we probe the effectiveness of the derivation procedure by checking the number of satisfying assignments generated by the derivation procedure in the first RT iterations. In our experiments, we found few rounds are sufficient to detect for each benchmark whether a large number of satisfying assignments can be derived from a seed. In fact, on some benchmarks, the derivation algorithm can derive a few satisfying solutions from a seed in the beginning, but no solution could be derived afterwards. Thus, a small DT value can be used to avoid Derivable being set to true on these benchmarks, while it will not change on other benchmarks. Based on these observations, we set RT = 3 and DT = 16.

By Theorem 1, we obtain that

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Theorem 2. The set Solutions returned by Algorithm 2 contains only satisfying assignments of  $\Phi$ . 352

#### 5. Implementation and Evaluation

We implement Algorithms 1—2 as an open-source tool ESAMPLER in C++, with QUICKSAMPLER as the underlying seed generator. QUICKSAMPLER takes a Boolean formula and its independent support as inputs, and outputs a set of assignments. However, as mentioned above, assignments produced by QUICK-SAMPLER may be duplicated or not satisfy the formula. As we focus on satisfying assignments of each Boolean formula in this work, we modify it so that duplicated and unsatisfying assignments are omitted. To demonstrate the generic nature of Algorithm 1 for deriving satisfying assignments from a seed, we also implement Algorithm 2 with UNIGEN3 as the underlying seed generator in our tool ESAMPLER. In contract to QUICKSAMPLER, UNIGEN3 only produces satisfying assignments for each given Boolean formula and the satisfying assignments are sampled uniformly at random with theoretical guarantees.

ESAMPLER takes a Boolean formula in the DIMACS [50] format and other required options as inputs, and outputs a set of satisfying assignments for the given Boolean formula. To reduce the memory usage of storing the satisfying assignments, we only store and output the satisfying assignments for the variables in the given independent support. Indeed, the truth values of the independent support determine those of the other variables, thereby the satisfying assignments can be easily completed.

In the rest of this work, we denote by ESAMPLER+QS (resp. ESAMPLER+UG) our tool ESAMPLER using QUICKSAMPLER (resp. UNIGEN3) as the underlying seed generator.

We mainly compare ESAMPLER+QS with three state-of-the-art tools QUICK-SAMPLER, STS and UNIGEN3 [29]. As done by [28], for a fair comparison, we modify STS so that the additional independent support information can be used by STS. To show the generic nature of Algorithm 1, we also compare ESAMPLER+UG with UNIGEN3.

Benchmarks. To evaluate the performance, we conducted extensive experiments. Industrial testing and verification instances are typically proprietary and unavailable for published research. Therefore, we conducted experiments on the publicly available benchmarks from UNIGEN [30], which consist of 370 Boolean formulas in the DIMACS format and the independent supports thereof. Indeed, the independent supports of most Boolean formulas could be computed using MIS [48] in few seconds. These benchmarks come from four classes of problem instances:

- 1. ISCAS89: constraints arising from ISCAS89 circuits with parity conditions on randomly chosen subsets of outputs and next-state variables;
- 2. SMTLib: bit-blasted versions of SMTLib benchmarks;
- 3. ProgSyn: constraints arising from automated program synthesis; and
- 4. BMC: constraints arising in bounded model checking of circuits.

Note that the accompanied independent supports of these benchmarks may contain variables that are not involved in the corresponding Boolean formulas; such variables are removed from the independent supports in our experiments. We remark that our approach also works without the given independent supports, in which case the independent support of a Boolean formula contains all the involved variables.

Since it does not make any sense to compute solutions for unsatisfiable Boolean formulas or the satisfiability cannot be solved, we checked the satisfiability of all these Boolean formulas with a timeout of one hour per Boolean formula using Z3 [51]. There are two unsatisfiable formulas (79.sk\_4\_40 and 36.sk\_3\_77), and four unsolvable formulas (logcount.sk\_16\_86, log2.sk\_72\_391,

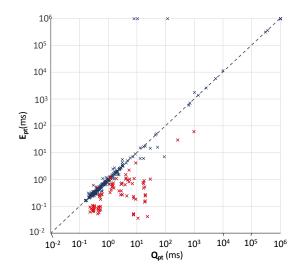


Figure 2: ESAMPLER+QS vs. QUICKSAMPLER

xpose.sk\_6\_134, and listReverse.sk\_11\_43). These formulas are not considered here, leaving 364 Boolean formulas.

Experiment setup. In our experiments, for each sampler and each Boolean formula, we run the sampler once on the Boolean formula. Though samplers are non-deterministic, results on a large number of Boolean formulas are sufficient to demonstrate the performance of ESAMPLER. The maximal number MaxPerSeed of satisfying assignments per seed is 10,000 and the maximal number T of satisfying assignments to compute is 1,000,000, unless the recent 10 assignments/pseudosolutions already exist. As aforementioned, we set RT=3 and DT=16 for ESAMPLER. For STS and QUICKSAMPLER, we use their default parameter settings. All the experiments were conducted on Intel Xeon E5-2620 v4  $2.10 \, \text{GHz}$  CPU with  $256 \, \text{RAM}$  GB and the one-hour timeout.

## 5.1. Comparison of ESAMPLER +QS and QUICKSAMPLER

Figure 2 shows the scatter plot comparing the average execution time per satisfying assignment between ESAMPLER+QS and QUICKSAMPLER on all the 364 formulas. Timeout occurred along the top or right border; the red color indicates that Derivable is set true by Algorithm 2, namely, it determines that our derivation procedure is able to derive a large number of satisfying assignments. Points below (resp. above) the diagonal line indicate that ESAMPLER+QS performs better (resp. worse) than QUICKSAMPLER.

The comparison of QuickSampler and ESampler+QS for a representative subset of the benchmarks is reported in Table 1. Columns benchmark, #Vars and #Cls respectively show the name, numbers of variables and clauses in each Boolean formula. Columns  $Q_t$  and  $E_t$  (resp.  $Q_{pt}$  and  $E_{pt}$ ) give the total execution time in thousand seconds (ks) (resp. execution time per satisfying assignment in milliseconds (ms)) of QuickSampler and ESampler+QS, respec-

Table 1: Comparison of QUICKSAMPLER and ESAMPLER+QS

| Benchmark               | $\# \mathrm{Vars}$ | #Cls   | $Q_t(ks)$ | $Q_n$           | $Q_{pt}(ms)$ | $E_t(ks)$ | $E_n$     | $E_{dn}$                | $E_{pt}(ms)$ | $\frac{Q_{pt}}{E_{pt}}$ |
|-------------------------|--------------------|--------|-----------|-----------------|--------------|-----------|-----------|-------------------------|--------------|-------------------------|
| s27_new_15_7            | 17                 | 43     | 0.00      | 48              | 1.39         | 0.00      | 48        | 42                      | 0.54         | 2.56                    |
| blasted_case.54         | 203                | 725    | 0.20      | 691,127         | 0.30         | 0.20      | 664,548   | 0                       | 0.30         | 0.99                    |
| 20.sk_1_51              | 15,475             | 60,994 | 3.94      | 491,074         | 8.02         | 1.67      | 1,520,152 | $\sim$ 1,520k           | 1.10         | 7.31                    |
| s35932_7_4              | 17,849             | 44,425 | 4.22      | 245,506         | 17.17        | 0.63      | 1,270,247 | $\sim$ 1,270k           | 0.50         | 34                      |
| blasted_case.126        | 302                | 1,129  | 0.34      | 1,007,411       | 0.34         | 0.34      | 1,022,991 | 0                       | 0.33         | 1.03                    |
| blasted_case.40         | 245                | 650    | 0.41      | 1,149,017       | 0.35         | 0.41      | 1,149,017 | 0                       | 0.36         | 0.99                    |
| s349_3_2                | 198                | 469    | 0.24      | 1,008,386       | 0.24         | 0.07      | 1,142,757 | $\sim$ 1,088k           | 0.06         | 3.81                    |
| 56.sk_6_38              | 4,842              | 17,828 | 1.97      | 1,004,037       | 1.96         | 1.18      | 1,093,080 | $\sim$ 1,092k           | 1.08         | 1.81                    |
| blasted_case.107        | 618                | 1,661  | 0.82      | $1,\!149,\!017$ | 0.72         | 0.84      | 1,149,017 | 0                       | 0.73         | 0.98                    |
| s832a_15_7              | 693                | 2,017  | 0.53      | 1,001,732       | 0.53         | 0.52      | 1,000,093 | 4                       | 0.52         | 1.01                    |
| $s420\_new\_7\_4$       | 312                | 770    | 0.35      | $1,\!117,\!085$ | 0.31         | 0.08      | 1,048,576 | $\sim 1,043 \mathrm{k}$ | 0.07         | 4.18                    |
| blasted_case.124 $\mid$ | 133                | 386    | 0.23      | 1,039,563       | 0.22         | 0.22      | 1,008,715 | 0                       | 0.22         | 1.02                    |
| s35932_15_7             | 17,918             | 44,709 | 4.29      | 145,499         | 29.46        | 1.34      | 1,270,247 | $\sim$ 1,270k           | 1.06         | 27                      |
| blasted_case.207        | 824                | 2,128  | 1.02      | 1,149,017       | 0.89         | 0.98      | 1,149,017 | 0                       | 0.86         | 1.04                    |
| blasted_case.120        | 284                | 851    | 0.41      | 1,113,780       | 0.37         | 0.40      | 1,044,731 | 0                       | 0.38         | 0.97                    |
| 63.sk_3_64              | 7,242              | 24,379 | 4.04      | 917,681         | 4.41         | 0.30      | 1,200,120 | $\sim$ 1,200k           | 0.25         | 17                      |
| s420_7_4                | 312                | 770    | 0.32      | 1,058,100       | 0.31         | 0.10      | 1,366,784 | $\sim 1,363 \text{k}$   | 0.07         | 4.14                    |
|                         |                    |        |           |                 |              |           |           |                         |              |                         |

tively. Columns  $Q_n$  and  $E_n$  show the total numbers of satisfying assignments generated by QuickSampler and ESampler+QS, respectively. Column  $E_{dn}$  gives the numbers of satisfying assignments generated by our derivation procedure. The last column provides the ratio of execution time per satisfying as-

signment between QUICKSAMPLER and ESAMPLER+QS, depicting the speedup of ESAMPLER+QS. We can observe when our derivation procedure works, it can produce more satisfying assignments (e.g., 20.sk\_1\_51 and s35932\_7\_4) than QUICKSAMPLER in the same time budget, while when it does not work well, it often does not produce any satisfying assignments (e.g., blasted\_case.54 and blasted\_case.40). Note that, since QUICKSAMPLER is a randomized approach, QUICKSAMPLER and ESAMPLER+QS may produce different satisfying assignments when our derivation procedure does not work, although ESAMPLER+QS is built on QUICKSAMPLER.

Summary. ESAMPLER+QS and QUICKSAMPLER respectively failed on 11 and 7 benchmarks due to the failures of MaxSAT solving. The difference between the numbers of the failed benchmarks indicates that the soft constraints generated randomly slightly affect MaxSAT solving. When ESAMPLER+QS determined that the derivation procedure can generate a large number of satisfying assignments, ESAMPLER+QS performed better than QUICKSAMPLER on almost all the benchmarks. While ESAMPLER+QS determined that our derivation procedure was not able to generate a large number of satisfying assignments, ESAMPLER+QS was comparable to QUICKSAMPLER. Specifically, ESAMPLER+QS was faster than QUICKSAMPLER on 227 benchmarks. It was 1.66× faster on average and more than 5× faster on 41 benchmarks, while it was 1.2 times slower on 16 benchmarks.

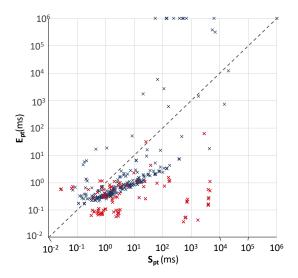


Figure 3: ESAMPLER+QS vs. STS

# 5.2. Comparison of ESAMPLER+QS and STS

Figure 3 shows the scatter plot comparing the average execution time per satisfying assignment between ESAMPLER+QS and STS on all the 364 formulas. Recall that timeout occurred along the top or right border, the red color

Table 2: Comparison of STS and ESAMPLER+QS

| Benchmark        | # Vars | #Cls   | $S_t(ks)$ | $S_n$       | $S_{pt}(ms)$ | $\mid E_t(ks) \mid$ | $E_n$     | $E_{dn}$            | $E_{pt}(ms)$ | $\frac{S_{pt}}{E_{pt}}$ |
|------------------|--------|--------|-----------|-------------|--------------|---------------------|-----------|---------------------|--------------|-------------------------|
| s27_new_15_7     | 17     | 43     | 0.00      | 48          | 0.85         | 0.00                | 48        | 42                  | 0.54         | 1.57                    |
| blasted_case.54  | 203    | 725    | 1.45      | 961,782     | 1.51         | 0.20                | 664,548   | 0                   | 0.30         | 5.06                    |
| 20.sk_1_51       | 15,475 | 60,994 | 3.60      | 151,948     | 23.69        | 1.67                | 1,520,152 | $\sim$ 1,520k       | 1.10         | 21                      |
| s35932_7_4       | 17,849 | 44,425 | 3.49      | 800         | 4,361        | 0.63                | 1,270,247 | $\sim$ 1,270k       | 0.50         | 8,757                   |
| blasted_case.126 | 302    | 1,129  | 0.92      | 1,000,006   | 0.92         | 0.34                | 1,022,991 | 0                   | 0.33         | 2.78                    |
| blasted_case.40  | 245    | 650    | 1.53      | 1,000,000   | 1.53         | 0.41                | 1,149,017 | 0                   | 0.36         | 4.30                    |
| s349_3_2         | 198    | 469    | 0.31      | 1,000,028   | 0.31         | 0.07                | 1,142,757 | $\sim 1,088 { m k}$ | 0.06         | 4.94                    |
| 56.sk_6_38       | 4,842  | 17,828 | 1.99      | 1,000,048   | 1.99         | 1.18                | 1,093,080 | $\sim$ 1,092k       | 1.08         | 1.84                    |
| blasted_case.107 | 618    | 1,661  | 3.60      | 558,950     | 6.44         | 0.84                | 1,149,017 | 0                   | 0.73         | 8.82                    |
| s832a_15_7       | 693    | 2,017  | 1.55      | 1,000,018   | 1.55         | 0.52                | 1,000,093 | 4                   | 0.52         | 2.97                    |
| s420_new_7_4     | 312    | 770    | 0.72      | 1,000,001   | 0.72         | 0.08                | 1,048,576 | $\sim$ 1,043k       | 0.07         | 9.68                    |
| blasted_case.124 | 133    | 386    | 0.32      | 1,000,013   | 0.32         | 0.22                | 1,008,715 | 0                   | 0.22         | 1.47                    |
| s35932_15_7      | 17,918 | 44,709 | 3.50      | 800         | 4,380        | 1.34                | 1,270,247 | $\sim$ 1,270k       | 1.06         | 4,140                   |
| blasted_case.207 | 824    | 2,128  | 3.60      | $276,\!250$ | 13.03        | 0.98                | 1,149,017 | 0                   | 0.86         | 15                      |
| blasted_case.120 | 284    | 851    | 1.59      | 1,000,000   | 1.59         | 0.40                | 1,044,731 | 0                   | 0.38         | 4.13                    |
| 63.sk_3_64       | 7,242  | 24,379 | 3.60      | 148,050     | 24.31        | 0.30                | 1,200,120 | $\sim$ 1,200k       | 0.25         | 97                      |
| s420_7_4         | 312    | 770    | 0.74      | 1 000 038   | 0.74         | 0.10                | 1.366.784 | $\sim$ 1.363 $k$    | 0.07         | 0 00                    |

indicates that Derivable is set true by Algorithm 2, and points below the diagonal line indicate that ESAMPLER+QS performs better than QUICKSAMPLER, and vice versa.

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Table 2 reports the performance of STS and ESAMPLER+QS for the same

representative subset of the benchmarks. Column  $S_t$  (resp.  $S_{pt}$ ) gives the total execution time in thousand seconds (ks) (resp. execution time per satisfying assignment in milliseconds (ms)) of STS. Column  $S_n$  shows the total number of satisfying assignments generated by STS for each Boolean formula. The last column provides the ratio of execution time per satisfying assignment between STS and ESAMPLER+QS, depicting the speedup of ESAMPLER+QS.

Summary. STS failed on 1 benchmark because the underlying SAT solver
Minisat failed to solve the Boolean formula, while ESAMPLER+QS failed on 11
benchmarks. In general, ESAMPLER+QS performed better than STS on most
benchmarks. It was faster on 316 benchmarks (5.47× faster on average and
more than 10× faster on 93 benchmarks), while it was 1.2 times slower on only
benchmarks.

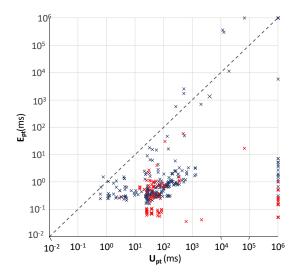


Figure 4: ESAMPLER+QS vs. UNIGEN3

## 5.3. Comparison of ESAMPLER+QS and UNIGEN3

Figure 4 shows the scatter plot comparing the average execution time per satisfying assignment between ESAMPLER+QS and UNIGEN3 on all the 364 formulas. Almost all the points are below the diagonal line, indicating ESAMPLER+QS significantly outperforms UNIGEN3.

Table 3 reports the performance of UNIGEN3 and ESAMPLER+QS on the same representative subset of benchmarks. Column  $U_t$  (resp.  $U_{pt}$ ) gives the total execution time in thousand seconds (ks) (resp. execution time per satisfying assignment in milliseconds (ms)) of UNIGEN3. Column  $U_n$  shows the total number of satisfying assignments generated by UNIGEN3 for each Boolean formula. The last column provides the ratio of execution time per satisfying assignment between UNIGEN3 and ESAMPLER+QS, depicting the speedup of ESAMPLER+QS.

Table 3: Comparison of UNIGEN3 and ESAMPLER+QS

| 539.9                   | 0.07         | $\sim$ 1,363k         | 1,366,784 | 0.10      | 37.79        | 95,260  | 3.60      | 770    | 312    | s420_7_4         |
|-------------------------|--------------|-----------------------|-----------|-----------|--------------|---------|-----------|--------|--------|------------------|
| 300                     | 0.25         | $\sim$ 1,200k         | 1,200,120 | 0.30      | 75.01        | 48,004  | 3.60      | 24,379 | 7,242  | 63.sk_3_64       |
| 182.9                   | 0.38         | 0                     | 1,044,731 | 0.40      | 69.5         | 51,799  | 3.60      | 851    | 284    | blasted_case.120 |
| 279                     | 0.86         | 0                     | 1,149,017 | 0.98      | 239.92       | 15,026  | 3.60      | 2,128  | 824    | blasted_case.207 |
| _                       | 1.06         | $\sim$ 1,270k         | 1,270,247 | 1.34      |              | 0       | 3.60      | 44,709 | 17,918 | s35932_15_7      |
| 182.1                   | 0.22         | 0                     | 1,008,715 | 0.22      | 40.28        | 89,376  | 3.60      | 386    | 133    | blasted_case.124 |
| 519.9                   | 0.07         | $\sim 1,043 { m k}$   | 1,048,576 | 0.08      | 36.39        | 98,934  | 3.60      | 770    | 312    | s420_new_7_4     |
| 52.17                   | 0.52         | 4                     | 1,000,093 | 0.52      | 27.13        | 132,705 | 3.60      | 2,017  | 693    | s832a_15_7       |
| _                       | 0.73         | 0                     | 1,149,017 | 0.84      | ı            | 0       | 3.60      | 1,661  | 618    | blasted_case.107 |
| 32.01                   | 1.08         | $\sim$ 1,092k         | 1,093,080 | 1.18      | 34.57        | 104,149 | 3.60      | 17,828 | 4,842  | 56.sk_6_38       |
| 415.8                   | 0.06         | $\sim 1,088 \text{k}$ | 1,142,757 | 0.07      | 24.95        | 144,279 | 3.60      | 469    | 198    | s349_3_2         |
| 198.5                   | 0.36         | 0                     | 1,149,017 | 0.41      | 71.46        | 50,380  | 3.60      | 650    | 245    | blasted_case.40  |
| 141.4                   | 0.33         | 0                     | 1,022,991 | 0.34      | 46.67        | 77,185  | 3.60      | 1,129  | 302    | blasted_case.126 |
| _                       | 0.50         | $\sim$ 1,270k         | 1,270,247 | 0.63      | ı            | 0       | 3.60      | 44,425 | 17,849 | s35932_7_4       |
| 46.56                   | 1.10         | $\sim$ 1,520k         | 1,520,152 | 1.67      | 51.21        | 70,312  | 3.60      | 60,994 | 15,475 | 20.sk_1_51       |
| 75.87                   | 0.30         | 0                     | 664,548   | 0.20      | 22.76        | 158,168 | 3.60      | 725    | 203    | blasted_case.54  |
| 38.57                   | 0.54         | 42                    | 48        | 0.00      | 20.83        | 48      | 0.00      | 43     | 17     | s27_new_15_7     |
| $\frac{U_{pt}}{E_{pt}}$ | $E_{pt}(ms)$ | $E_{dn}$              | $E_n$     | $E_t(ks)$ | $U_{pt}(ms)$ | $U_n$   | $U_t(ks)$ | #Cls   | # Vars | Benchmark        |

Summary. UNIGEN3 failed on 40 benchmarks. Recall that ESAMPLER+QS failed on 11 benchmarks. No matter whether or not ESAMPLER+QS determined that the derivation procedure was able to generate a large number of satisfying assignments, ESAMPLER+QS performed significantly better than UNIGEN3 on almost all the benchmarks. Specifically, ESAMPLER+QS was faster than UNI-

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GEN3 on 348 benchmarks. It was 69.8× faster on average and more than 100× faster on 194 benchmarks, while it was 1.2 times slower on only 7 benchmarks.

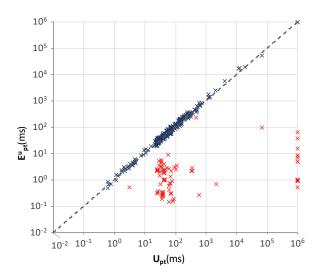


Figure 5: ESAMPLER+UG vs. UNIGEN3

## 5.4. Comparison of ESAMPLER+UG and UNIGEN3

Figure 5 shows the scatter plot comparing the average execution time per satisfying assignment between UNIGEN3 and ESAMPLER+UG on all the 364 formulas. Almost all the red points are below the diagonal line while almost all the blue points are close to the diagonal line. indicating that ESAMPLER+UG significantly outperforms UNIGEN3 on the benchmarks on which Derivable was set to true by Algorithm 2, while it was still comparable on other benchmarks.

Table 4 reports the performance of UniGen3 and ESampler+UG on the same representative subset of benchmarks. Column  $E^u_t$  gives the total execution time in thousand seconds (ks) of ESampler+UG, while column  $E^u_{pt}$  gives the execution time per satisfying assignment in milliseconds (ms). Column  $E^u_n$  shows the total number of satisfying assignments generated by ESampler+UG for each Boolean formula, while column  $E^u_{dn}$  gives the numbers of satisfying assignments generated by the derivation procedure. The last column provides the ratio of execution time per satisfying assignment between UniGen3 and ESampler+UG, depicting the speedup of our algorithm.

Summary. UNIGEN3 failed on 40 benchmarks while ESAMPLER+UG failed on 25 benchmarks. Specifically, ESAMPLER+UG was faster than UNIGEN3 on 207 benchmarks. It was  $2.19\times$  faster on average and more than  $10\times$  faster on 85 benchmarks, while there were only 12 benchmarks on which it was at least  $1.2\times$  slower. The results demonstrate the generic nature of Algorithm 1 for deriving satisfying assignments from a seed using UNIGEN3 as the underlying seed generator in our tool ESAMPLER.

Table 4: Comparison of UNIGEN3 and ESAMPLER+UG

| 129.95                    | 0.292            | $\sim 1,093 \text{k}$ | 1,096,960 | 0.32        | 37.79        | 95,260  | 3.60      | 770    | 312    | s420_7_4          |
|---------------------------|------------------|-----------------------|-----------|-------------|--------------|---------|-----------|--------|--------|-------------------|
| 80.39                     | 0.934            | $\sim$ 3,959k         | 3,960,000 | 3.70        | 75.02        | 48,004  | 3.60      | 24,379 | 7,242  | 63.sk_3_64        |
| 1.25                      | 55.6             | 0                     | 64,745    | 3.60        | 69.5         | 51,799  | 3.60      | 851    | 284    | blasted_case.120  |
| 1.00                      | 239.9            | 0                     | 15,054    | 3.61        | 239.9        | 15,026  | 3.61      | 2,128  | 824    | blasted_case.207  |
| 4109                      | 15.92            | $\sim$ 229k           | 230,032   | 3.66        | 65455        | 55      | 3.60      | 44,709 | 17,918 | s35932_15_7       |
| 1.05                      | 38.27            | 0                     | 94,080    | 3.60        | 40.27        | 89,376  | 3.60      | 386    | 133    | blasted_case.124  |
| 154.47                    | 0.236            | $\sim 1,083 \text{k}$ | 1,086,720 | 0.26        | 36.39        | 98934   | 3.60      | 770    | 312    | s420_new_7_4      |
| 0.98                      | 33.4             | ယ                     | 130,075   | 3.60        | 27.13        | 132,705 | 3.60      | 2,017  | 693    | s832a_15_7        |
| ,                         | 1                | 0                     | 0         | 3.60        | 1            | 0       | 3.60      | 1,661  | 618    | blasted_case.107  |
| 10.81                     | 3.2              | $\sim 1,194 \text{k}$ | 1,194,765 | 3.82        | 34.57        | 104,149 | 3.60      | 17,828 | 4,842  | 56.sk_6_38        |
| 10.35                     | 2.41             | $\sim$ 974k           | 1,001,811 | 2.42        | 24.95        | 144,279 | 3.60      | 469    | 198    | s349_3_2          |
| 0.92                      | 48.59            | 0                     | 46,327    | 3.60        | 71.46        | 50,380  | 3.60      | 650    | 245    | blasted_case.40   |
| 0.96                      | 56.01            | 0                     | 74,111    | 3.60        | 46.7         | 77,185  | 3.60      | 1,129  | 302    | blasted_case.126  |
| 1                         | 67.16            | $\sim$ 59 $k$         | 60,005    | 4.03        | ·            | 0       | 3.60      | 44,425 | 17,849 | s35932_7_4        |
| 17.46                     | 2.93             | $\sim$ 1,319k         | 1,320,000 | 3.87        | 51.2         | 70,312  | 3.60      | 60,994 | 15,475 | 20.sk_1_51        |
| 1.25                      | 18.21            | 0                     | 197,693   | 3.60        | 22.8         | 158,168 | 3.60      | 725    | 203    | blasted_case.54   |
| 2.20                      | 0.01             | 42                    | 48        | 0.00        | 0.02         | 48      | 0.00      | 43     | 17     | $s27\_new\_15\_7$ |
| $\frac{U_{pt}}{E_{pt}^u}$ | $E_{pt}^{u}(ms)$ | $E_{dn}^u$            | $E_n^u$   | $E_t^u(ks)$ | $U_{pt}(ms)$ | $U_n$   | $U_t(ks)$ | #Cls   | # Vars | Benchmark         |
|                           |                  |                       |           |             |              |         |           |        |        |                   |

# 5.5. Execution Time vs Number of Satisfying Assignments

To see the relation between the execution time and the number of satisfying assignments, we evaluate ESAMPLER on four randomly chosen benchmarks by varying the execution time and counting the number of satisfying assignments. Figure 6(a) and Figure 6(b) respectively show the plots of results on the

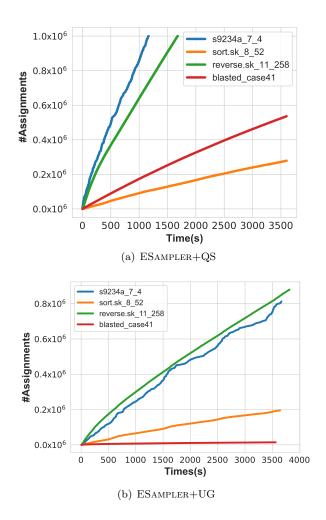


Figure 6: Time vs. #assignments of ESAMPLER

four randomly chosen benchmarks using ESAMPLER+QS and ESAMPLER+UG, where the x-axis is the execution time (in seconds) and the y-axis is number of satisfying assignments (#assignments). We can observe that the number of satisfying assignments for each benchmark is almost linear in the execution time. These results demonstrate the effectiveness of our derivation procedure.

## 5.6. Testing Uniformity

Similar to QUICKSAMPLER, ESAMPLER does not provide a guarantee of uniformity. Remark that UNIGEN3 provides a theoretical guarantee of uniformity based on hashing, at the cost of sampling efficiency. We empirically show

that the uniformity of the solutions can be controlled by adjusting the maximal number of solutions per seed, i.e., the parameter MaxNumPerSeed. We run both ESAMPLER+QS and ESAMPLER+UG on a randomly selected benchmark (i.e., 27.sk\_3\_32) on which our derivation procedure works, where duplicated solutions are recorded to measure uniformity and the mutation phase of QUICKSAMPLER is disabled to be more precise.

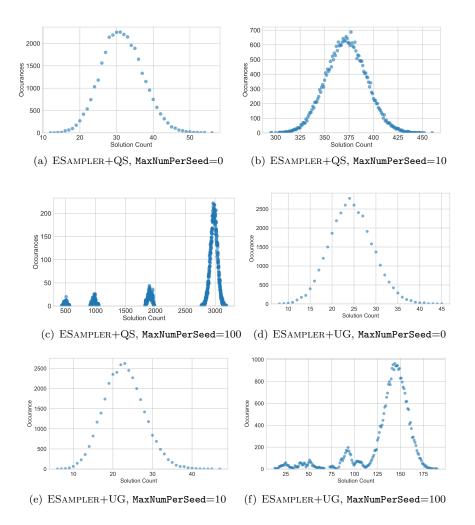


Figure 7: Distributions of solutions

Figure 7 depicts the distributions of solutions ESAMPLER+QS and ESAMPLER+UG when MaxNumPerSeed is set to 0, 10 and 100, where (x, y) denotes that there are y unique solutions each of which occurs x times. We can observe that the smaller the parameter MaxNumPerSeed is, the closer the distribution

is to the normal distribution, meaning that the solutions generated by our tool are actually close to uniform when MaxNumPerSeed is chosen properly.

## 6. Application to Bayesian Inference

In this section, to further show the effectiveness and efficiency of ESAMPLER in real-world applications, we apply ESAMPLER to Bayesian inference, namely, computing the posterior probability of a query given evidence in a Bayesian network [52, 53].

**Bayesian inference**. A Bayesian network is a tuple (V, E, T), where  $V = \{X_1, \dots, X_n\}$  is a finite set of nodes each of which represents a discrete random variable,  $E \subseteq V \times V$  is a finite set of edges each of which represents dependence between two random variables. (V, E) forms a directed acyclic graph (DAG), and T is a finite set of conditional probability tables (CPTs) each of which encodes the conditional probability distribution of a random variable. Given a random variable X, value a, and a partial assignment v of some other random variables, the Bayesian inference is to compute the posterior probability  $\Pr(X = a \mid v)$ .

Bayesian inference is a well-known  $\#\mathbf{P}$ -complete problem [54]. Sang et al. [31] proposed an encoding from the Bayesian inference problem to the model-counting problem of Boolean formulas ( $\#\mathbf{SAT}$ ), which we leverage to solve Bayesian inference.

Bayesian inference to #SAT. Given a Bayesian network (V, E, T) and a Bayesian inference query  $\Pr(X = a \mid v)$ , Sang et al. [31] use chance variables to encode entries in CPTs and state variables to encode the values of the nodes, based on which two Boolean formulas  $\Phi_1 \wedge \Phi_2 \wedge \Phi_3$  and  $\Phi_1 \wedge \Phi_2$  can be constructed, where  $\Phi_1$  encodes the Bayesian network (V, E, T),  $\Phi_2$  encodes the partial assignment v and  $\Phi_3$  encodes X = a. With satisfying assignments of  $\Phi_1 \wedge \Phi_2 \wedge \Phi_3$  and  $\Phi_1 \wedge \Phi_2$ , one can calculate (or approximate) the posterior probability in the Bayesian network. Suppose  $n_1$  (resp.  $n_2$ ) denotes the number of the discovered satisfying assignments of  $\Phi_1 \wedge \Phi_2 \wedge \Phi_3$  (resp.  $\Phi_1 \wedge \Phi_2$ ), the approximate posterior probability is  $\frac{n_1}{n_2}$ . Furthermore, if  $n_1$  is exact, then  $\frac{n_1}{n_2}$  gives an upper-bound of the posterior probability; on the other hand, if  $n_2$  is exact, then  $\frac{n_1}{n_2}$  gives a lower-bound.

Performance of Bayesian inference. To evaluate ESAMPLER for Bayesian inference, we use the QUICKSAMPLER sampler as the seed generation engine to solve Bayesian inference of the plan recognition problems provided by [31]. There are 11 plan recognition problems given as Bayesian networks on which we compute the posterior probability for each random variable, resulting in 11,326 Bayesian inference queries. For each query, we sample satisfying assignments until the 10 recently generated assignments already exist. Solved by ESAMPLER in 65,122 seconds, the calculated probabilities of variables are shown in Table 5, where the columns (Var ID) show the indices of the random variables, and the columns (Prob) show the calculated posterior probabilities of the random variable. For the sake of brevity, we show first hundred variables in problem tire-3,

Table 5: Calculated probabilities of variables in problem tire-3.

|        | 0. 0                  | r   |                       |        | г                     | or objective or |                       |
|--------|-----------------------|-----|-----------------------|--------|-----------------------|-----------------|-----------------------|
| Var ID | $\operatorname{Prob}$ | Var | $\operatorname{Prob}$ | Var ID | $\operatorname{Prob}$ | Var ID          | $\operatorname{Prob}$ |
| 5      | 0.9391                | 43  | 0.0335                | 64     | 0.4827                | 83              | 0.0000                |
| 12     | 0.0000                | 44  | 0.2407                | 66     | 0.0000                | 85              | 0.5720                |
| 19     | 0.0000                | 45  | 0.0000                | 67     | 0.0000                | 86              | 0.0000                |
| 20     | 0.6307                | 50  | 0.6962                | 68     | 0.2452                | 87              | 0.0000                |
| 21     | 0.0000                | 51  | 0.0000                | 69     | 0.0000                | 88              | 0.9085                |
| 25     | 0.8177                | 52  | 0.5173                | 71     | 0.0000                | 89              | 0.0000                |
| 26     | 0.4827                | 53  | 0.9645                | 72     | 0.2470                | 90              | 0.2159                |
| 27     | 0.9997                | 54  | 0.2358                | 74     | 0.1287                | 91              | 0.5366                |
| 31     | 0.9997                | 55  | 0.7548                | 75     | 0.5021                | 92              | 0.1862                |
| 33     | 0.5172                | 56  | 0.1253                | 76     | 0.2502                | 94              | 0.2452                |
| 35     | 0.0000                | 58  | 0.9837                | 77     | 0.0000                | 95              | 0.9342                |
| 36     | 0.0000                | 59  | 0.9977                | 78     | 0.0499                | 96              | 0.9903                |
| 38     | 0.5028                | 61  | 0.2670                | 79     | 0.1691                | 98              | 0.2981                |
| 40     | 0.3693                | 62  | 0.0000                | 80     | 0.0253                |                 |                       |
| 41     | 0.4972                | 63  | 0.0000                | 82     | 0.0000                |                 |                       |

and the results are omitted if the posterior probability is 1. We notice that the reported probabilities in Table 5 are approximation of the exact posterior probabilities when the sampler fails to generate all the possible satisfying assignments of a Bayesian inference query. Remark that computing exact posterior probabilities are computational hard (#P-completeness).

Comparison of samplers on Bayesian inference. To compare the efficiency of ESAMPLER (i.e., ESAMPLER+QUICKSAMPLER), STS, QUICKSAMPLER and UNIGEN3 in solving Bayesian inference problems, we test them on 11 randomly chosen formulas from the plan recognition problems, each of which is aimed to compute 100,000 satisfying assignments within 10 minutes. The other settings are the same as in Section 5.

The results are reported in Table 6, where the last three columns provide the ratio of the execution time per satisfying assignment for QuickSampler, STS, UniGen3 to ESampler respectively, measuring the speedup of ESampler. All the samplers are able to generate satisfying assignments except that UniGen3 failed on 4 benchmarks (log-1, log-4, log-5 and tire-1). For the sake of brevity, we only report the number of satisfying assignments generated by ESampler. We can observe that ESampler outperforms the other three samples on Bayesian inference. On average, ESampler is 13.8, 18.7 and 556.3 times faster than QuickSampler, STS and UniGen3, respectively.

#### 7. Conclusion

We have proposed a novel approach to derive a large set of satisfying assignments from a seed assignment without invoking computationally expensive SAT solving. Our approach is orthogonal to the previous techniques and could be integrated into the existing SAT samplers. We have also developed a new tool ESAMPLER, based on the recent samplers QUICKSAMPLER and UNIGEN3

Table 6: Comparison of ESAMPLER, QUICKSAMPLER, STS and UNIGEN3 on benchmarks derived from Bayesian inferences of plan recognition problems

| Benchmark | #Vars | $\#\mathrm{Cls} \mid E_t(s)$ | $E_n$   | $E_{dn}$ | $E_{pn}(ms)$ | $\frac{Q_{pt}}{E_{pt}}$ | $\frac{S_{pt}}{E_{pt}}$ | $\frac{U_{pt}}{E_{pt}}$ |
|-----------|-------|------------------------------|---------|----------|--------------|-------------------------|-------------------------|-------------------------|
| 4step     | 165   | 418   17.8                   | 66,935  | 0        | 0.27         | 1.01                    | 1.78                    | 36.47                   |
| 5step     | 177   | 475   3.01                   | 80,033  | ~80k     | 0.04         | 7.69                    | 11.52                   | 273.8                   |
| log-1     | 939   | 3,785   10.4                 | 160,016 | 160k     | 0.07         | 16.31                   | 72.99                   | -                       |
| log-2     | 1,377 | 24,777   53.9                | 110,011 | 110k     | 0.49         | 12.67                   | 50.01                   | 251.8                   |
| log-3     | 1,413 | 29,487   166                 | 170,017 | 170k     | 0.98         | 4.78                    | 24.14                   | 511.3                   |
| log-4     | 2,303 | 20,963   18.2                | 120,012 | 120k     | 0.15         | 32.74                   | 423.8                   | -                       |
| log-5     | 2,701 | 29,534   958                 | 10,001  | 100k     | 95.81        | 4,355                   | 2.99                    | -                       |
| tire-1    | 352   | 1,038   9.4                  | 130,347 | ~130k    | 0.07         | 6.81                    | 17.01                   | -                       |
| tire-2    | 550   | 2,001   15.8                 | 160,016 | 160k     | 0.10         | 6.63                    | 10.72                   | 731.7                   |
| tire-3    | 578   | 2,004   13.7                 | 140,014 | 140k     | 0.10         | 11.31                   | 16.65                   | 3,496                   |
| tire-4    | 812   | 3,222   20.1                 | 120,012 | 120k     | 0.17         | 5.92                    | 13.64                   | 5,009                   |

as the seed generator. The extensive experiments on publicly available benchmarks and application on Bayesian inference confirmed the effectiveness and efficiency of our approach.

In future, we plan to further improve the performance of the tool ESAMPLER and extend our derivation approach to SMT formulas, as well as their practical applications.

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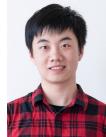
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