
Algorithm 1 Top-level Design of LS-Sampler.

Input: F : SMT formula; k : the number of samples;**Output:** A : the set of generated, valid samples;

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1:  $F_1 \leftarrow \text{Simplify } F$ ;  
2:  $F_2 \leftarrow \text{Boolean skeleton of } F_1$ ;  
3:  $A \leftarrow \{\}$ ;  
4:  $\text{Initialize}(F_1, F_2)$   
5: for  $iter \leftarrow 1$  to  $k$  do  
6:    $\alpha \leftarrow \text{pickSample}(F_1, \text{prob}, A)$ ;  
7:    $A \leftarrow A \cup \alpha$ ;  
8:   for each  $b_i \in B(F_2)$  do  
9:      $\text{prob}(b_i) \leftarrow \text{updateProb}(x_i, A)$ ;  
10:  end for  
11: end for  
12: return  $A$ ;
```

Algorithm 2 The Initialize Component.

Input: F_1 : Simplified SMT formula; F_2 : Boolean skeleton of F_1 ;**Output:** $prob$: vector of all skeleton variables' sampling probabilities;

```
1: for each integer variable  $v_i \in V(F_1)$  do
2:   if  $v_i \geq lb$  and  $v_i \leq ub$  then
3:      $v_i$  is assigned with a random value in  $[lb, ub]$ ;
4:   else if  $v_i \geq lb$  or  $v_i \leq ub$  then
5:      $v_i$  is assigned with  $lb$  or  $ub$ ;
6:   else if  $v_i$  is unbounded then
7:      $v_i$  is assigned with 0;
8:   end if
9: end for
10: for each  $b_i \in B(F_2)$  do
11:   set  $prob(b_i)$  as 0.5;
12: end for
```

Algorithm 3 The PickSample Component.

Input:

F_1 : Simplified SMT formula;
 F_2 : Boolean skeleton of F_1 ;
 $prob$: vector of all skeleton variables' sampling probabilities;

Output:

α^* : the selected, valid sample;

- 1: $C \leftarrow \{\}$;
 - 2: **for** $j \leftarrow 1$ to λ **do**
 - 3: $\alpha_j \leftarrow$ generate a sample by sampling the value of each skeleton variable b_i according to $prob(b_i)$;
 - 4: $\alpha_j^* \leftarrow \text{DAHS}(F_1, F_2, \alpha_j, prob)$;
 - 5: $C \leftarrow \alpha_j^*$;
 - 6: Calculate $\text{score}(\alpha_j^*)$ for the valid sample α_j^* ;
 - 7: **end for**
 - 8: $\alpha^* \leftarrow$ the valid sample in C with the largest score;
 - 9: **return** α^*
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Algorithm 4 The UpdateProb Component.

Input: b_i : skeleton variable in F_2 ; A : The current set of valid samples;**Output:** $prob(b_i)$: Updated sampling probability for x_i ;1: $B \leftarrow$ the set of all b_i -false samples in A ;2: $prob(b_i) \leftarrow |B|/|A|$;3: **return** $prob(b_i)$

Algorithm 5 The Diversity-aware Heuristic Search Component.

Input:

F_1 : simplified SMT formula;
 F_2 : Boolean skeleton of F_1 ;
 α : the input sample, possibly invalid;
 $prob$: vector of all skeleton variables' sampling probabilities;

Output:

α^* : the modified, valid sample;

```
1:  $\alpha^* \leftarrow \alpha$ ;  
2: while  $\alpha^*$  is invalid do  
3:   if  $DV \neq \emptyset$  then  
4:      $x_i \leftarrow$  the variable with the largest gain in  $DV$ , breaking ties by preferring the variable with the  
       largest priority.  
5:   else  
6:      $c \leftarrow$  a random unsatisfied clause;  
7:      $x_i \leftarrow$  the variable with the largest priority in  $c$ ;  
8:   end if  
9:   if  $x_i$  is pure Boolean variable then  
10:     $\alpha^* \leftarrow \alpha$  with  $x_i$  flipped;  
11:  else if  $x_i$  is LIA atom then  
12:     $a \leftarrow$  the theoretical atom corresponding to  $x_i$ ;  
13:     $op \leftarrow$  a  $cm(v, a)$  operation with the greatest cscore;  
14:  end if  
15: end while  
16: return  $\alpha^*$ ;
```

> $gain(x_i)$: the decrement in the number of unsatisfied clauses by flipping the current truth value of x_i .

> Decreasing variable: x_i is decreasing variable if $gain(x_i) > 0$.

> DV : the current set of all decreasing variables.

> $priority(x_i)$: $(1 - v_i) \times prob(x_i) + v_i \times (1 - prob(x_i))$