software

A Product of Shape & Sequences abstractions

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Introduction

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What do we want to verify ?

Introduction

When we talk about automatic static analysis of program manipulating dynamic data-structure, there are several properties we are interested in.

```
tree *insert(tree *t, int v) {
        tree *m = malloc(sizeof(tree));
2
        m->left = m->right = NULL;
3
       m \rightarrow data = v;
4
       if (!t) {
6
          // Empty case
        } else {
7
          tree \hat{*}c = t:
8
          while (v < c->data && c->left ||
9
                  v >= c->data && c->right)
            if (v \le c > data) {
               c = c -> left;
12
            } else {
13
               c = c - right;
14
            3
15
          if (v \le c > data) {
16
17
            c \rightarrow left = m:
          } else {
18
            c->right = m;
19
20
          return t;
        }
22
23
```

- 1. No ill-pointer (null, ...) dereference "c->"
- Preservation of structural invariants
 "If t is a well-formed binary tree then so
 is the returned value."
- Partial functional correctness
 "If t is a well-formed BST, then the
 returned value r should be a well-formed
 BST containing the same elements as t
 plus value v."

Conclusion

Comparison of existing static analysis over dynamic data structures

Various automatic static analysis over dynamic data-structures have been proposed:

	pointer	structural	partial f ^{al}	
Analysis	doroforonco	invariants	correctness	
	derererence	IIIvallallus	SLL	tree
Pointer analysis	 Image: A set of the set of the	×	×	×
Shape analysis based on				
3-Value logic	 	~	×	×
e.g. [Sagiv et al. TOPLAS, 99]				
Separation logic	 	~	×	×
e.g. [Chang et al. POPL, 08]				
k-limited graphs	~	~	~	×
e.g. [Bouajjani et al, CAV, 10]				

None of these approaches could prove functional correctness of insertion into a binary search tree !

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None of these approaches could prove functional correctness of insertion into a binary search tree !

How to improve the expressiveness of automatic static analysis over dynamic data-structures to prove partial functional correctness?

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Separation Logic based shape analysis

[Chang et al. POPL, 2008] introduces a shape analysis based on abstract interpretation.

It uses a subset of **separation logic** [Reynolds, LICS 02] as an abstract representation for memory states:

• Abstract memory regions are connected with the **separating conjunction**. It expresses that these regions are disjoint

This allows to reason locally

• Inductive data-structures are synthesized by inductive predicates **Example** The predicate **tree**(c), denoting a binary tree:



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Inductive predicates are not expressive enough



 \Longrightarrow This predicate is expressive enough to prove memory safety & structure preservation.

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Problem This not enough for partial functional correctness: tree forgets the content !

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Inductive predicates are not expressive enough



 \Longrightarrow This predicate is expressive enough to prove memory safety & structure preservation.

Problem This not enough for partial functional correctness: tree forgets the content !

[Li et al. SAS, 2015] added set parameters expressing the content of data-structures.

Problem Set parameters express no constraint in respect to order of appearance !

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Sequence parameters

Our solution: Express constraints on the sequence of values stored in the tree. Add a **sequence parameter** to the inductive predicate: tree(c, S).



The specification of the (partial) functional correctness of insert can be expressed as:

$$\begin{cases} \mathsf{tree}(\mathsf{t},S) \\ S = \mathsf{sort}(S) \end{cases} \mathsf{r} = \mathsf{insert}(\mathsf{t},\mathsf{v}) \begin{cases} \mathsf{tree}(\mathsf{r},S') \\ \text{where } S' = \mathsf{sort}(S.[\mathsf{v}]) \end{cases}$$

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Sequence parameters

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Requires to extend the shape analysis to derive precise sequence constraints. Requires an abstract domain to reason about (possibly) sorted sequences.

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An abstract domain reasoning over sequence constraints

To reason on content with order, length constraint, extremal elements, sortedness

A Reduced product between the sequence domain and an existing shape domain

To express constraints over the content of inductive data structures

Evaluation of the analysis in the MemCAD tool

To demonstrate the gain of the expressiveness, the versatility of the approach, and discuss its cost

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Domain description

We build a domain in order to abstract sets of functions from variables to values and sequences of values:

$$\left\{\begin{array}{c} \alpha \mapsto 2\\ \delta \mapsto 1 \end{array}\right\} \left\{\begin{array}{c} S \mapsto 4; 6; 1\\ S_1 \mapsto 4; 6 \end{array}\right\}$$

An **abstract value** $\sigma_s^{\#}$ of the sequence abstract domain $\mathbb{D}_s^{\#}$ consists of:



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$$\operatorname{\mathsf{guard}}_{\scriptscriptstyle\mathcal{S}}: \mathbb{D}^{\#}_{\scriptscriptstyle\mathcal{S}} o \operatorname{seq.}$$
 constraint $\to \mathbb{D}^{\#}_{\scriptscriptstyle\mathcal{S}}$

$$S = S_1 \cdot [\alpha] \land S = \mathbf{sort}(S)$$
$$\land S_1 = \mathbf{sort}(S_1)$$

To assume $S_r = [\alpha]$, \mathbf{guard}_s follows this algorithm:

$$\wedge \mathsf{mset}_S = \{\!\!\{\alpha\}\!\!\} \uplus \mathsf{mset}_{S_1}$$

$$\wedge \operatorname{len}_S = 1 + \operatorname{len}_{S_1} + \operatorname{len}_{S_2}$$

$$\wedge \min_S \leqslant lpha \leqslant \max_S \ \wedge \min_S \leqslant \min_{S_1} \wedge \max_{S_1} \leqslant \max_S$$

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 $S = S_1.[\alpha] \land S = \mathbf{sort}(S)$ $\land S_1 = \mathbf{sort}(S_1)$ $\land S_r = [\alpha]$

$$\wedge \operatorname{mset}_{S} = \{\!\!\{\alpha\}\!\!\} \uplus \operatorname{mset}_{S_{1}}$$
$$\wedge \operatorname{mset}_{S_{r}} = \{\!\!\{\alpha\}\!\!\}$$

$$\begin{split} &\wedge \min_{S} \leqslant \alpha \leqslant \max_{S} \\ &\wedge \min_{S} \leqslant \min_{S_{1}} \wedge \max_{S_{1}} \leqslant \max_{S} \\ &\wedge \min_{S_{r}} = \alpha = \max_{S_{r}} \end{split}$$

- 1. add the new definition in the conjunction
- 2. add content/length/bounds constraints

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- 1. add the new definition in the conjunction
- 2. add content/length/bounds constraints
- 3. fold other definitions

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- 1. add the new definition in the conjunction
- 2. add content/length/bounds constraints
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4. Saturate constraints

$$S = S_1....S_n$$

$$\frac{\forall i, S_i = \mathbf{sort}(S_i) \quad \forall i < j, \mathbf{max}_{S_i} \leqslant \mathbf{min}_{S_j}}{S = \mathbf{sort}(S)}$$

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- 1. add the new definition in the conjunction
- 2. add content/length/bounds constraints
- 3. fold other definitions
- 4. Saturate constraints $S = S_1....S_n$ $\frac{\forall i, S_i = \mathbf{sort}(S_i) \quad \forall i < j, \mathbf{max}_{S_i} \leqslant \mathbf{min}_{S_j}}{S = \mathbf{sort}(S)}$
- 5. detect & remove cyclic constraints

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$$S = S_1 . S_r \land S = \mathsf{sort}(S)$$

$$\land S_1 = \mathsf{sort}(S_1)$$

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To assume $S_r = [\alpha]$, guard_s follows this algorithm:

- 1. add the new definition in the conjunction
- 2. add content/length/bounds constraints
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4. Saturate constraints

$$S = S_1....S_n$$

$$\frac{\forall i, S_i = \text{sort}(S_i) \quad \forall i < j, \max_{S_i} \leq \min_{S_j}}{S = \text{sort}(S)}$$

5. detect & remove cyclic constraints

Theorem: Soundness of guard_s

 $\gamma_{\scriptscriptstyle S}(\operatorname{guard}_{\scriptscriptstyle S}(\sigma_{\scriptscriptstyle S}^{\#},S=E)) \text{ contains all valuations in } \gamma_{\scriptscriptstyle S}(\sigma_{\scriptscriptstyle S}^{\#}) \text{ satisfying } S=E.$

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Abstract lattice operators

•
$$\sqsubseteq_{\mathcal{S}} : \mathbb{D}_{\mathcal{S}}^{\#} \to \mathbb{D}_{\mathcal{S}}^{\#} \to \{ \mathsf{true}, \mathsf{false} \}$$

Abstract inclusion checking, using verify_s

•
$$\sqcup_{\mathcal{S}} : \mathbb{D}_{\mathcal{S}}^{\#} \to \mathbb{D}_{\mathcal{S}}^{\#} \to \mathbb{D}_{\mathcal{S}}^{\#}$$

That tries to infer common definitions from both inputs.

Example
$$\begin{pmatrix} S = S_1.S_2 \\ \land S_3 = [] \end{pmatrix} \sqcup_s \begin{pmatrix} S = S_2.S_3 \\ \land S_1 = [] \end{pmatrix} = (S = S_1.S_2.S_3)$$

• $\nabla_{\mathcal{S}}: \mathbb{D}_{\mathcal{S}}^{\#} \to \mathbb{D}_{\mathcal{S}}^{\#} \to \mathbb{D}_{\mathcal{S}}^{\#}$

That selects the constraints in the left arguments verified in the right one.

Shape analysis with sequence predicates

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Integrating sequence parameters in the shape domain

The tree(c) predicate only synthesizes full binary trees.

To abstract partial trees, the shape domain uses a segment predicate treeseg(1, c).





The shape domain automatically derives treeseg from tree.

The analysis must keep tracks of the content stored in the segment

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Integrating sequence parameters in the shape domain

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To abstract partial trees, the shape domain uses a segment predicate treeseg(1, c).





The shape domain automatically derives treeseg from tree.

The analysis must keep tracks of the content stored in the segment

In order to reason precisely over inductive predicates, the shape analysis relies on:

- Unfold: refines the memory by materializing synthesized memory.
- Fold: extrapolates the memory state to gain generality.

Used to over-approximate two memory states

For each of these operations, the shape domain should **derive the corresponding** sequence constraints to assume or verify.

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Adding sequence parameters to segment predicates



The sequence stored in the tree is: 0 1 2 3 4 5 6 9 10 11 12

The analysis needs to recall the location of the missing sequence in treeseg.

 \implies the segment predicate has **two sequence parameters**: S_1, S_2

One for each side of the missing sequence

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Refining abstract memory state with unfolding

To analyze $if(1){v= l->data}$ with initial state tree(1, S)



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Refining abstract memory state with unfolding

To analyze $if(1){v= l->data}$ with initial state tree(l, S)

1. The numerical constraint $\texttt{l} \neq \texttt{0x0}$ is guarded in the numerical part of the sequence domain.



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Refining abstract memory state with unfolding

To analyze $if(1){v= l->data}$ with initial state tree(1, S)

- 1. The numerical constraint $\texttt{l} \neq \texttt{0x0}$ is guarded in the numerical part of the sequence domain.
- 2. To materialize 1->data, the analysis unfolds the predicate

The abstract memory is replaced by the definition: δ, S_l, S_r are fresh variables The numerical and sequences constraints are guarded in the sequence domain



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Refining abstract memory state with unfolding

To analyze $if(1){v= l->data}$ with initial state tree(1, S)

- 1. The numerical constraint $\texttt{l} \neq \texttt{0x0}$ is guarded in the numerical part of the sequence domain.
- 2. To materialize 1->data, the analysis **unfolds the predicate**

The abstract memory is replaced by the definition: δ , S_l , S_r are **fresh variables** The numerical and sequences constraints are guarded in the sequence domain

• The empty case: Inconsistent with the if assumption \Longrightarrow Discarded



Refining abstract memory state with unfolding

To analyze $if(1){v= l->data}$ with initial state tree(1, S)

- 1. The numerical constraint $1 \neq 0x0$ is guarded in the numerical part of the sequence domain.
- 2. To materialize 1->data, the analysis unfolds the predicate

The abstract memory is replaced by the definition: δ , S_l , S_r are **fresh variables** The numerical and sequences constraints are guarded in the sequence domain

- The empty case: Inconsistent with the if assumption \implies Discarded
- The non-empty case: c->data corresponds to δ .







Refining abstract memory state with unfolding

To analyze $if(1){v= l->data}$ with initial state tree(1, S)

- 1. The numerical constraint $\texttt{l} \neq \texttt{0x0}$ is guarded in the numerical part of the sequence domain.
- 2. To materialize 1->data, the analysis unfolds the predicate

The abstract memory is replaced by the definition: δ, S_l, S_r are **fresh variables** The numerical and sequences constraints are guarded in the sequence domain

- The empty case: Inconsistent with the if assumption \Longrightarrow Discarded
- The non-empty case: c->data corresponds to δ .
- 3. The assignment $\mathbf{v} \leftarrow \boldsymbol{\delta}$ is performed.



The resulting disjunction of abstract states over approximates the original state.

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Folding the abstract state

Fold generalizes the abstract state by rewriting parts of the memory into a predicate. The analysis first checks that some constraints hold in the sequence domain.

Folding an inductive predicate



$$\operatorname{verify}_{\mathcal{S}}(\sigma_{\mathcal{S}}^{\#}, S = S_l.[\delta].S_r) = \operatorname{true}$$



Folding segment and predicates



$$\operatorname{verify}_{\mathcal{S}}(\sigma_{\mathcal{S}}^{\#}, S = S_1.S_0.S_2) = \operatorname{true}$$



Theorem: Soundness of folding

The folded abstract state over-approximates the original one.

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Lattice operators

Inclusion checking

Folds the left input until both are syntactically equal.

Upper bound

Folds both inputs until they are syntactically equal.

Widening

With these operators, we design a sound and automatic static analysis by forward abstract interpretation. And the analysis checks that the final state satisfies to post-condition to prove partial functional correctness.

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Proving the insertion into a BST

After two iterations, the analysis inferred the following loop invraiant:



Finally, the analysis was able to prove that the final state sastifies the post condition:



Experiments
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The analysis described has been implemented in the MemCAD static analyzer available at gitlab.inria.fr/memcad/memcad

For each test, we specify:

- the full inductive predicates,
- the pre- and post-conditions,

Everything else (segment predicates/loop invariants) is inferred by the analysis.

(Q1) Is this analysis precise enough to prove memory safety $({\sf Safe})$ and functional properties $({\sf Fc})$?

 ${\rm (Q2)}$ How significant is the overhead of the combined analysis compared to the baseline?

(Q3) Can this analysis successfully verify real-world C libraries?

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Experiment 1: Classical list & BST programs

	wo/ seq	with seq parameters			
Example	Safe time		e	Fc	
	verified	overhd.	% num	verified	
	Singly lir	iked list			
concat	Safe	2.4x	21.7%	Fc	
deep copy	Safe	1.7 ×	18.1%	Fc	
length	Safe	4.7×	50.0%	Fc	
insert at position	Safe	5.4x	60.2%	Fc	
sorted insertion	Safe	6.1×	47.3%	Fc	
minimum	Safe	7.8×	45.9%	Fc	
insertion sort	Safe	29.0×	46.0%	Fc	
bubble sort	Safe	19.1×	51.5%	Fc	
merge sorted lists	Safe	9.6x	51.4%	Fc	
Binary search trees					
Insertion	Safe	6.0x	38.6%	Fc	
Delete max	Safe	6.2×	48.6%	Fc	
Search (present)	Safe	4x	45.3%	Fc	
BST to list	Safe	3.2×	38.2%	Fc	
list to BST	Safe	11.9×	46.1%	Fc	

Expressiveness

- Prove Fc for complex programs including 3 sorting algorithms
- Sequences improve precision for Safe!

Overhead

- High slowdown for complex programs Up to 30x for insertion sort
- Most of it in the numerical domain Quadratic cost of sortedness checking Length constraints are expensive

• Sequence domain slows down convergence

Needs one more iteration for $\nabla_{\mathcal{S}}$ to stabilize.

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Experiment 2: Real-world libraries

Introduc

We tested MemCAD on real-world list libraries implementing various features:

	Linux	FreeRTOS	GDSL
Circular DLL with distinguished header	Yes	Yes	Yes
Extreme sentinel nodes	No	No	Yes
Intrusive	Yes	Yes	No
Pointer to header	No	Yes	No
Length in header	No	Yes	Yes
Sorted	No	Yes	No

	Linux		FreeRTOS		GDSL	
	wo/ seq	w/ seq	wo/ seq	w/ seq	wo/ seq	w/ seq
Safe	4/4✔	4/4✔	4/4✔	4/4✔	13✔ 1 X (†)	14/14
Fc		4/4✔		4/4✔		12 ✔ 2४ (‡)

- †: Cannot prove Safe for extraction at position.
- ‡: Cannot prove **Fc** for min/max extraction.

Conclusion

Conclusion

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How to improve the expressiveness of static analysis over dynamic data-structures to prove partial functional correctness?

• Design of a novel sequence abstract domain

It leverages existing numerical/set domains to express length/bounds/content constraints.

Integration into a separation logic based shape analysis

The reduced product derives corresponding sequence constraints for unfolding/weakening.

• Implementation in the MemCAD static analyzer

Proves partial functional correctness for complex algorithms on SLL/BST and real world libraries.

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Thank you !

Lemma

If
$$S = S_1 \dots S_n$$
, then
$$S = \operatorname{sort}(S) \Leftrightarrow \forall i, S_i = \operatorname{sort}(S_i) \land \forall i < j, \max_{S_i} \leqslant \min_{S_j}$$

Question The number of constraints in the right-hand side is quadratic! Could we relax it for j := i + 1?

 $\begin{array}{l} \Longrightarrow \mbox{ NO } ! \mbox{ Because of the empty sequence case } ! \\ \mbox{By consistency of the concretization: } \nu_{\mathcal{S}}(S) = \varepsilon \Longrightarrow \left\{ \begin{array}{c} \max_{S} = -\infty \\ \min_{S} = +\infty \end{array} \right. \end{array} \right. \label{eq:max_s}$

$$\text{Consider } \nu_{\mathcal{S}} = \left\{ \begin{array}{ll} S \mapsto 3 \ 1 \\ S_1 \mapsto 3 \\ S_2 \mapsto \varepsilon \\ S_3 \mapsto 1 \end{array} \right\} \begin{array}{ll} \text{We have indeed:} \\ \nu_{\mathcal{S}} \models S = S_1.S_2.S_3 \\ \nu_{\mathcal{S}} \models S_i = \operatorname{sort}(S_i), \quad \forall i \quad \text{But:} \\ \nu_{\mathcal{S}} \models \max_{S_1} \leqslant \min_{S_2} \\ \nu_{\mathcal{S}} \models \max_{S_2} \leqslant \min_{S_3} \end{array} \right.$$

Removing cyclic constraints

Assume the abstract state $\sigma^\#_{\scriptscriptstyle \mathcal{S}}$ contains the following constraints:

$$S = S_1.S'.S$$

$$\land S' = S_3.S''$$

$$\land S'' = S.S_4$$

If we inline definitions over S^\prime and $S^{\prime\prime}$ into the definition of S we obtain:

$$S=S_1.S_3.S.S_4.S_2$$
 The constraints over S,S',S'' are replaced by
$$\left\{ \begin{array}{l} S_1=S_2=S_3=S_4=[]\\ S=S'=S'' \end{array} \right.$$

If one constraint contains at least one atom [α], then the state is reduced to \perp_{s} .

 $S={\rm sort}(S)$ does not count as a cyclic constraint as the implementation of the abstract domain does not represent it as such.

Concatenating inductive predicates

seg-full case



$$\operatorname{verify}_{\mathcal{S}}(\sigma_{\mathcal{S}}^{\#}, S = S_1.S_0.S_2) = \operatorname{true} \xrightarrow{}$$



seg-seg case



$$\begin{split} & \mathsf{verify}_{s}(\sigma_{s}^{\#}, S_{1} = S_{1}'.S_{1}'') = \mathsf{true} \\ & \mathsf{verify}_{s}(\sigma_{s}^{\#}, S_{2} = S_{2}''.S_{2}') = \mathsf{true} \end{split}$$



Segment tree predicate



Hypothesis to derive segment from full predicate

Hypothesis

- The constraint over sequence parameter is only concatenation based
- The argument of each recursive call occurs exactly once in the constraint









S



























Weakening: inclusion test



To verify:

Weakening: inclusion test





To verify: $S_{l,1} = []$ $S_{l,2} = []$ $\alpha_l = \alpha_c$

The constraints are simplified. $\alpha \neq 0$

 $\alpha_l = \alpha_l$ $S_1 = []$ $S_2 = [\delta].S_r$ The numerical ones are checked with verify,

The sequence ones are used for definition of S_1 and S_2

Weakening: inclusion test



fold





To verify:
$$\begin{split} S_{l,1} &= [] \\ S_{l,2} &= [] \\ \alpha_l &= \alpha_c \\ S_1 &= S_{l,1} \\ S_2 &= S_{l,2}. \left[\delta\right].S_r \\ \alpha_t &\neq \texttt{0x0} \end{split}$$
 The constraints are simplified. $\begin{array}{l} \alpha \neq 0 \\ \alpha_l = \alpha_l \\ S_1 = [] \\ S_2 = [\delta].S_r \end{array}$ The numerical ones are checked with **verify**_s The sequence ones are used for definition of S₁ and S₂





$\alpha \leftrightarrow \alpha_{\rm c} \mapsto \alpha_l$	$, \alpha_r$
$S \longleftrightarrow S_0 \mapsto S_l$	$, S_r$
$\alpha \leftrightarrow \alpha_{t} \mapsto \alpha$	$, \alpha$
$] \leftrightarrow S_1 \mapsto []$	$,S_{l}.[\delta]$
$[] \leftrightarrow S_2 \mapsto [\delta].S$	r, []

$$\begin{array}{l} \alpha \leftrightarrow \alpha_{c} \mapsto \alpha_{l} &, \alpha_{r} \\ S \leftrightarrow S_{0} \mapsto S_{l} &, S_{r} \\ \alpha \leftrightarrow \alpha_{t} \mapsto \alpha &, \alpha \\ [] \leftrightarrow S_{1} \mapsto [] &, S_{l} \cdot [\delta] \\ [] \leftrightarrow S_{2} \mapsto [\delta] \cdot S_{l} , [] \end{array}$$

Result:



 $S = \mathbf{sort}(S)$

 \square_{Σ}

 $\wedge \alpha \neq \texttt{0x0}$

$$\begin{split} S &= \mathsf{sort}(S) \land S = S_l.[\delta].S_r \\ S_i &= \mathsf{sort}(S_i) \quad i \in \{l, r\} \\ &\land \alpha \neq \mathsf{0x0} \\ &\land \alpha_l \neq \mathsf{0x0} \\ &\land \forall < \delta \\ &\mathsf{max}_{S_l} \leqslant \delta \leqslant \mathsf{min}_{S_r} \end{split}$$

$$\begin{split} S &= \mathsf{sort}(S) \land S = S_l. [\delta]. S_r \\ S_i &= \mathsf{sort}(S_i) \ i \in \{l, r\} \\ &\land \alpha \neq \mathsf{0x0} \\ &\land \alpha_r \neq \mathsf{0x0} \\ &\land \forall \geqslant \delta \\ &\mathsf{max}_{S_l} \leqslant \delta \leqslant \mathsf{min}_{S_r} \end{split}$$

$$\begin{array}{l} \alpha \leftrightarrow \alpha_{c} \mapsto \alpha_{l} &, \alpha_{r} \\ S \leftrightarrow S_{0} \mapsto S_{l} &, S_{r} \\ \alpha \leftrightarrow \alpha_{t} \mapsto \alpha &, \alpha \\ [] \leftrightarrow S_{1} \mapsto [] &, S_{l} . [\delta \\ [] \leftrightarrow S_{2} \mapsto [\delta] . S_{l} , [] \end{array}$$

Result:



$$S = \operatorname{sort}(S)$$

$$S_1 = S_2 = []$$

$$\wedge \alpha = \alpha_c = \alpha_t \neq 0 \mathrm{x} 0$$

 \square_{Σ}

$$\begin{split} S &= \operatorname{sort}(S) \land S = S_0.S_2 \\ S_i &= \operatorname{sort}(S_i) \quad i \in \{l, r, 1, 2\} \\ S_0 &= S_l \land S_1 = [] \land S_2 = [\delta].S_r \\ \land \alpha = \alpha_t \neq 0 \operatorname{xo} \\ \land \alpha_l = \alpha_c \neq 0 \operatorname{xo} \\ \land \forall < \delta = \min_{S_2} \land \max_{S_1} = -\infty \\ \max_{S_l} &\leq \delta \leqslant \min_{S_r} \end{split}$$

$$\begin{split} S &= \operatorname{sort}(S) \land S = S_1.S_0 \\ S_i &= \operatorname{sort}(S_i) \quad i \in \{l, r, 1, 2\} \\ S_0 &= S_r \land S_1 = S_l.[\delta] \land S_2 = [] \\ \land \alpha = \alpha_t \neq 0 \mathbf{x} 0 \\ \land \alpha_r = \alpha_c \neq 0 \mathbf{x} 0 \\ \land \mathbf{x} \geqslant \delta = \max_{S_1} \land \min_{S_2} = +\infty \\ \max_{S_l} &\leq \delta \leqslant \min_{S_r} \end{split}$$

$$\begin{array}{l} \alpha \leftrightarrow \alpha_{c} \mapsto \alpha_{l} &, \alpha_{r} \\ S \leftrightarrow S_{0} \mapsto S_{l} &, S_{r} \\ \alpha \leftrightarrow \alpha_{t} \mapsto \alpha &, \alpha \\ [] \leftrightarrow S_{1} \mapsto [] &, S_{l} \cdot [\delta] \\ [] \leftrightarrow S_{2} \mapsto [\delta] \cdot S_{l} , [] \end{array}$$

 $S = \operatorname{sort}(S)$ $S_1 = S_2 = []$ $\wedge \alpha = \alpha_c = \alpha_t \neq 0 \operatorname{xo}$

$$\begin{split} S &= \mathsf{sort}(S) \land S = S_0.S_2 \\ S_i &= \mathsf{sort}(S_i) \quad i \in \{l, r, 1, 2\} \\ S_0 &= S_l \land S_1 = [] \land S_2 = [\delta].S_r \\ \land \alpha = \alpha_t \neq 0 \mathsf{x} 0 \\ \land \alpha_l = \alpha_c \neq 0 \mathsf{x} 0 \\ \land \forall < \delta &= \min_{S_2} \land \max_{S_1} = -\infty \\ \max_{S_l} &\leq \delta \leqslant \min_{S_r} \end{split}$$

Result:

$$S_i = \operatorname{sort}(S_i) \quad i \in \{_, 0, 1, 2\}$$

$$\begin{split} S &= \operatorname{sort}(S) \land S = S_1.S_0 \\ S_i &= \operatorname{sort}(S_i) \quad i \in \{l, r, 1, 2\} \\ S_0 &= S_r \land S_1 = S_l.[\delta] \land S_2 = [] \\ \land \alpha = \alpha_t \neq 0 \mathbf{x} 0 \\ \land \alpha_r = \alpha_c \neq 0 \mathbf{x} 0 \\ \land \forall \geqslant \delta = \max_{S_1} \land \min_{S_2} = +\infty \\ \max_{S_l} &\leq \delta \leqslant \min_{S_r} \end{split}$$

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 ${\textstyle \bigsqcup}_{\Sigma}$

$$\begin{split} S &= \mathsf{sort}(S) \\ S_1 &= S_2 = \texttt{[]} \\ \wedge \alpha &= \alpha_{\texttt{c}} = \alpha_{\texttt{t}} \neq \texttt{0x0} \end{split}$$

$$\begin{split} S &= \mathsf{sort}(S) \land S = S_0.S_2 \\ S_i &= \mathsf{sort}(S_i) \quad i \in \{l, r, 1, 2\} \\ S_0 &= S_l \land S_1 = [] \land S_2 = [\delta].S_r \\ \land \alpha = \alpha_t \neq 0 \mathsf{x} 0 \\ \land \alpha_l = \alpha_c \neq 0 \mathsf{x} 0 \\ \land \forall < \delta = \min_{S_2} \land \max_{S_1} = -\infty \\ \max_{S_l} &\leq \delta \leqslant \min_{S_r} \end{split}$$

Result:

$$\begin{split} S_i &= \mathsf{sort}(S_i) \quad i \in \{_, 0, 1, 2\} \\ S &= S_1.S_0.S_2 \end{split}$$

$$\begin{split} S &= \operatorname{sort}(S) \land S = S_1.S_0 \\ S_i &= \operatorname{sort}(S_i) \quad i \in \{l, r, 1, 2\} \\ S_0 &= S_r \land S_1 = S_l.[\delta] \land S_2 = [] \\ \land \alpha = \alpha_t \neq 0 \mathbf{x} 0 \\ \land \alpha_r = \alpha_c \neq 0 \mathbf{x} 0 \\ \land \psi \geqslant \delta = \max_{S_1} \land \min_{S_2} = +\infty \\ \max_{S_l} &\leq \delta \leqslant \min_{S_r} \end{split}$$

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Result:

$$\begin{split} S_i &= \texttt{sort}(S_i) \quad i \in \{_, 0, 1, 2\} \\ S &= S_1.S_0.S_2 \\ \alpha_{c}, \alpha_{t} \neq \texttt{0x0} \end{split}$$

$$S = \operatorname{sort}(S)$$

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Result:

$$\begin{split} S_i &= \mathsf{sort}(S_i) \quad i \in \{_, 0, 1, 2\} \\ S &= S_1.S_0.S_2 \\ \alpha_c, \alpha_t \neq \mathsf{0x0} \\ \max_{S_1} &\leqslant v \leqslant \min_{S_2} \end{split}$$

$$S = \operatorname{sort}(S)$$

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$$\begin{cases} \text{tree}(t, S) \\ \land S = \text{sort}(S) \\ \end{cases}$$

if (t== NUL) {
// Empty case
}else{
 ptrec c= t;
 while(•v< c>d & c>>1 |
 vy> c>>d & c>>1 |
 vy> c>>d & c>>1 |
 c = c>>1 \\ belse {
 c = c->r;
 }
 if (v< c>>d) {
 c = c->r;
 }
 if (v< c>>d) {
 c = c->r = m;
 pelse{
 c->r = m;
 return t;
 }

 $\{\mathbf{tree}(\mathtt{t},\mathbf{sort}(S.[\mathtt{v}]))\}$



$$\{ tree(t, sort(S.[v])) \}$$



$$\begin{cases} \text{tree}(t, S) \\ \land S = \text{sort}(S) \end{cases} \\ \text{if} (t=* \text{NUL}) \{ // Empty \ case \} \text{else}(t) \\ \text{ptrec}(t) \\ \text{ptrec}(t) \\ \text{vs}(t) \\ \text{vs}(t) \\ \text{vs}(t) \\ \text{vs}(t) \\ \text{vs}(t) \\ \text{ss}(t) \\ \text{vs}(t) \\$$

```
\{ tree(t, sort(S.[v])) \}
```



$$\begin{cases} \text{tree}(t, S) \\ \land S = \text{sort}(S) \\ \end{cases}$$

if (t== NUL) {
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ptrec c= t;
while (v< c->d & c->1||
v>c->d & c->r ;
c = c->1e
belse {
c = c->r;
}
if (v< c->d) {
c->r = m;
}
return t;
} \end{cases}

 $\{ tree(t, sort(S.[v])) \}$






Exemple: insertion in binary search tree





 \Longrightarrow Invariant found after two iterations !

Exemple: insertion in binary search tree





 \Longrightarrow Invariant found after two iterations !