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# Une Table d'Association d'Intervalles Fusionnable

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**ABSTRACT.** *This article describes an efficient persistent mergeable data structure for mapping intervals to values. We call this data structure rangemap. We provide an example of application where the need for such a data structure arises (abstract interpretation of programs with pointer casts). We detail different solutions we have considered and dismissed before reaching the solution of rangemaps. We show how they solve the initial problem. We then describe their implementation and, as a conclusion, mention further work we would like to do.*

**RÉSUMÉ.** *Cet article décrit une structure de données représentant efficacement des tables d'associations persistantes indexées par des intervalles, ayant la propriété supplémentaire d'être fusionnable. Nous nommons (en anglais) rangemap cette structure de données. Nous donnons un exemple de circonstances dans lesquelles se rencontre le besoin d'associer de cette façon des valeurs à des intervalles (analyse par interprétation abstraite de programmes comportant des conversions de pointeurs). Nous détaillons différentes solutions envisagées puis écartées avant d'arriver à la solution des rangemaps. Nous montrons ensuite comment ceux-ci résolvent le problème initial. Enfin, nous décrivons leur implémentation avant d'évoquer les travaux en cours et futurs que nous nous proposons de mener à ce sujet.*

**KEYWORDS:** *Rangemaps, interval maps, Patricia trees, abstract interpretation, sharing*

**MOTS-CLÉS :** *Rangemaps, table d'association d'intervalles, arbres Patricia, interprétation abstraite, partage.*

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## 1. Introduction

### *The problem at hand*

The problem we are concerned with in this article is the representation of persistent maps indexed by intervals. The expected solution is a data structure that allows to retrieve the value associated to an interval when this interval has previously been used as the key in an insertion; an additional constraint is that the representation must also be efficient for finding what an interval is mapped to when this specific interval is not employed as a key in the map. In this case, the data structure must allow to retrieve all bindings that intersect the required interval, and let these partial results be combined to the programmer's liking.

As a concrete example, consider the modelization of the contents of a char array during the abstract execution of a C program. Assume that such platform characteristics as endianness and word size are known. The C program may take the address of any cell in this array and cast this address to an `int*`. If the run-time architecture allows it, the program may then use this pointer to write an `int` (on a 32-bit architecture, the `int` occupies four consecutive cells in the original char array). If the same pointer (as cast to `int*`) is now dereferenced, it is desirable to recover exactly the same abstract value that was previously written there. If a `char*` pointing to one of the four aforementioned cells is dereferenced, the result should be that the value read is a part of the stored `int`. And lastly, if an 8-byte `double` is read from the same location, the data structure should be able to indicate that the bits read are made partly from the `int`, and partly from other values, each of which may for instance have been written as a `char` previously.

In a concrete (say, hardware) implementation, the memory can be considered as an unadorned char array. A multi-byte memory access reads – or writes – several consecutive bytes, and that's all there is to it. In this case, there is no need to consider the memory as a map whose keys are intervals. But in the case of abstract values, an unfortunate loss of precision would occur if the same approach was employed. Consider as an example of abstract value for a 32-bit word the pair  $\{1, 258\}$ . Projecting this word-level abstract value in abstract values for the component bytes, we obtain the set  $\{1, 2\}$  for the least significant byte,  $\{0, 1\}$

for its neighbor, and  $\{0\}$  for the two most significant bytes. Now trying to read back a word from the same location in memory, it appears that the possible values for the data word are  $\{1, 2, 257, 258\}$ , an unacceptable approximation. Obviously, the component bytes of a data word can not be stored independently without concern for the implicit relationship between them.

For maximum generality, the data structure should not try to decide how to read a single char from a stored int, or how to recompose a double from several smaller values. Instead, it should be generic and call user-provided functions to recompose values when appropriate. Specifically, the structure should be generic enough that, by providing the right functions, little, big, and unknown endiannesses can all be accommodated.

Finally, C being an imperative programming language, the analysis may involve merging together several execution branches where the same char array has been modified in different ways. For good performance, the data structure should be mergeable, in the exact same sense as in Okasaki and Gill's [10]. We informally define mergeability as the following property:

**Property 1 (Mergeability).** *When iterating in parallel on two instances of the data structure, a divide-and-conquer approach allows to consider separately the component substructures. Additionally, the results obtained for these substructures can be cached, and have a good chance to be useful later, when processing other instances that are only slightly different from the initial ones.*

The mergeability property is named in reference to the merge operation that takes two trees  $t_1$  and  $t_2$ , and builds a tree where each key contains the merge of the values associated to this key in  $t_1$  and  $t_2$ . The merge function can quickly check the subtrees that appear at the same level in  $t_1$  and  $t_2$  for physical identity. If these subtrees are physically equal, the computation of their merge is immediate (the result subtree is in this case identical to the arguments). This can happen for instance when  $t_2$  was created by slightly modifying  $t_1$ , or, if hash-consing [2, 3] is used to ensure maximal sharing, at every opportunity.

## *Contents of the article*

This article shows the development of an efficient solution to the problem we have exposed. A starting point for the reflection is the Patricia tree structure, summarized in Sect. 2. We show in Sect. 3 how we considered various other solutions based upon AVL trees (Sect. 3.1) and Patricia trees (Sect. 3.2 and Sect. 3.3). This eventually leads us to describe the data structure answering our needs in Sect. 4. We have a prototype implementation of rangemaps, from which we have extracted relevant technical details in Sect. 5. Considerations about the future of this work conclude this article in Sect. 6.

## **2. Reminder: Patricia trees**

Morrison's Patricia trees [9] provide a great technical solution for mapping integers to values, when mergeability is important. In fact, Patricia trees are rather unique in this respect, and they will serve as a natural guide in our reflection.

Patricia trees are used to implement maps when there exists a natural lexicographical order on the keys. When used with integers as keys, the lexicographical order used is the comparison of the keys' binary representations. Either the little-endian or the big-endian representation can be used as long as the same choice is used consistently. If the big-endian representation is chosen, and assuming we have to deal only with positive integers (both assumptions we will make in the remainder of this article), the lexicographical order on binary representations coincides with the usual order of integers.

During the lookup of a key  $k$  in a Patricia tree, each node, starting from the root, tests one bit in the binary representation of  $k$  in the lexicographical order. When, for a given prefix, all keys present in the tree have the same value for the next bit, the comparison of this bit is skipped, so that in general, it takes about  $\log_2(n)$  comparisons to get the value associated to a key in a map of  $n$  bindings. There never are any rebalancing in Patricia trees: the nodes of the tree are hierarchized according to the lexicographical order that has been fixed in advance. Patricia trees can therefore be unbalanced (a worst-case example is a

map where the keys are 1, 2, 4, 8, 16... This map is represented as a comb). However, the height of a Patricia tree is bounded by the base-2 logarithm of the difference between the values of its smallest and largest key. If arbitrary 32-bit integers are used as keys, a Patricia tree can never have an height higher than 32 (and a lookup never require more than 32 comparisons).

When maps indexed by a type key different from `int` are required, it is still sometimes possible to employ Patricia trees. It may be a simple matter of tagging each object of type `key` at creation with a unique integer (e.g. an `id` field in a record type). This makes it possible to use Patricia trees for representing maps from keys to values, by using the `id` field as the actual key during lookups.

However, these maps may be expected to provide primitives that give access to the original keys, e.g. a function `mapi: (key -> 'a -> 'b) -> 'a t -> 'b t`. This may require to keep in memory a reverse map from ids to keys. Alternatively, it is possible to store in the Patricia tree, associated to the id, the original key together with the value. In effect, the latter method overloads each Patricia tree ever built so that it contains a copy of the reverse map for all keys that appear in it. The latter method is more memory-efficient when each possible key is used less than once on average. The former method is more memory efficient when each possible key is used many times (but note that a separate hashtable incurs overhead that needs to be amortized, and that applying this method may require the use of a weak hashtable).

### 3. Towards mergeable binary trees indexed by intervals

Consider now the problem of mapping non-overlapping intervals of integers to values. A concrete use case for this structure was provided in Sect. 1. The same property of mergeability that Patricia trees have is expected of this data structure.

#### *A data structure indexed by intervals*

Mapping an interval to a value is not the same thing as mapping all integers in the interval to this value. The former can sometimes be

used as a concise representation of the latter in a kind of Run-Length Encoding, as in diets [4]. In general though, the data structure should not make assumptions about what happens when a new binding added to a map overlaps with some of the bindings already in place. Behaviors that may be useful are to remove the overwritten binding completely, or more generally to alter the contents of the untouched parts of the original binding to reflect the fact that they were part of a larger binding that was partially overwritten. Providing these behaviors implies for instance that the binding  $0..31 \rightarrow \{22\}$  is not the same thing as 32 individual bindings  $0 \rightarrow \{22\}, 1 \rightarrow \{22\}, \dots$

The solution data structure should, for instance, be able to answer queries such as finding all bindings that intersect a given interval. Being able to do this is necessary for lookups, but also at the time of adding a new binding, in order to maintain the invariant that the intervals in the map are disjoint.

While it is possible to associate a unique integer to use as an id to any interval, this solution, applied naively, would store the intervals in the map without respect for their natural order, and as a consequence, the whole tree would have to be explored in order to find the bindings that intersect a given interval.

### **3.1. *Using AVL trees with the natural interval order***

Note that the intervals used as keys in a given map are guaranteed to be disjoint. The first idea is therefore to order them according to their natural order, which happens to be total among the keys of a single tree. Many data structures have been proposed for representing maps indexed by totally ordered keys as trees. In most of them, the tree structure can be taken advantage of to quickly look up a range of bindings. As an example, the Map module in the OCaml standard library (implemented with AVL trees [1]) can easily be augmented with functions for looking up all bindings that intersect the query interval. The resulting data structure, unfortunately, is no more mergeable than Map usually is. Balancing operations interfere with sharing.

### 3.2. Trying to use Patricia trees indexed by intervals

Since, for us, mergeability is an important criterion, and since Patricia trees have a reputation for being mergeable, it is natural to try to representing the map as a big-endian Patricia tree, using for instance the lower bound of each interval as its id. It may be possible to make this representation work, but mergeability again suffers unexpectedly. Consider indeed the example on Fig. 1.

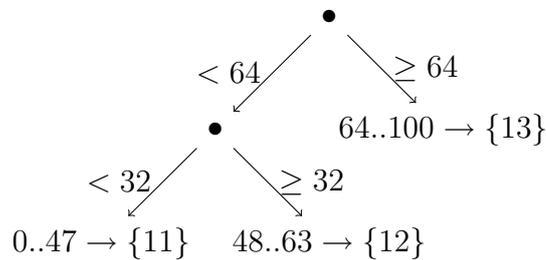


Figure 1: Interval map as a Patricia tree, using the minimum bound as id

In Fig. 1, the binding  $0..47 \rightarrow \{11\}$  is stored in the leftmost binding, but it contains information that may interfere with a differently located binding in another tree.

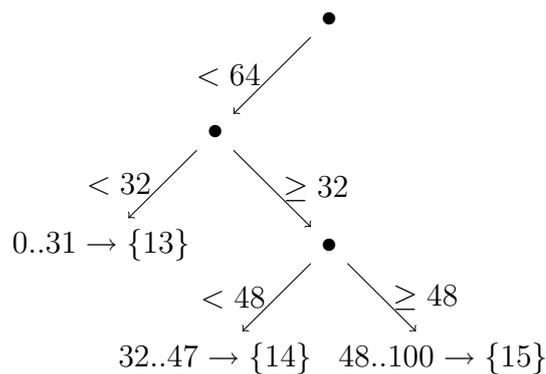


Figure 2: Another interval map represented as a Patricia tree

When computing the merge of the two trees from Fig. 1 and Fig. 2, the divide-and-conquer approach does not work! The 0..47 binding in the left-hand-side subtree of Fig. 1 contains information that is relevant for computing other branches of the result. The computation of the merge of such affected subtrees can therefore not be cached, since it depends on external factors. This choice of representation is simply not mergeable in the sense of Okasaki and Gill [10].

### 3.3. *Trying to use Patricia trees indexed by integers*

We will now say a word about a representation that is not the one that we are proposing, but that seems to us a valid alternative. Patricia trees can be made to omit the nodes whose children both eventually lead to the same leaf anyway (making them even closer than they already are to Binary Decision Diagrams working on the bits of the key). We have not encountered this optimization in the literature, perhaps because it requires comparing the values that are bound to the keys, which may be expensive outside the context of hash-consing. This optimization can be done by changing the `Leaf` constructor to contain a prefix instead of a key (meaning that in this tree, all keys with this prefix are bound to the same value). Then, it is only a matter of systematically replacing applications of the `Node` constructor by a “smart constructor” function which checks whether the left and right subtrees are both `Leaf` with the same value, and builds a `Leaf` in this case.

With this optimization enabled, Patricia trees can efficiently represent identical bindings to long consecutive sequences of keys. It becomes feasible to represent the  $0..47 \rightarrow \{12\}$  binding as a binding from each of the integers  $0, 1, \dots, 47$  to an identical something. This something should of course contain the bound value  $\{12\}$ , but also the interval that serves as key of this binding, so that an access inside the interval (say, to the value bound to the interval  $8..15$ ) is allowed to recover the information that this binding is part of a sequence that goes from 0 to 47. In particular, adding a new binding to the interval  $8..15$  in this tree should either transform the existing  $0..47 \rightarrow \dots$  binding into three bindings  $0..7, 8..15, 16..47$  or into a single  $8..15$  binding depending on the desired semantics for overlapping writes. In both cases, it

is necessary to have the information that the current binding at 8..15 is really a sub-part of a binding to a larger interval, so that this binding can be completely modified or removed. It would be necessary to make use of zippers[8] in order to navigate efficiently from the bindings at 8..15 to the adjoining 40 bindings that used to be related to them.

As a foreseen drawback with this approach, note that the factoring of identical bindings suggested here only works well when the binary representations of the integers contained in the key interval are characterized by a few common prefixes. A key such as 0..47 would require two actual bindings in the tree to be represented (for 0..32 and 33..47). A worst-case interval such as 0..62 would require 6 actual bindings to represent (for 0..31, 33..47, 48..55, 56..59, 60..61 and 62..62).

#### **4. The solution we propose**

We propose to build maps from intervals to values, with the same mergeable quality that Patricia trees display for integer-indexed maps. As a first difference from Patricia trees, but similarly to diets [4], in our proposal bindings are recorded on the nodes, whereas Patricia trees record bindings at the leaves. Diets are a data structure to represent sets, when a total order, and successor and predecessor functions are available for the elements – for instance, integers. Diets are efficient when long sequences of consecutive elements commonly occur. On the other hand, our proposal is a data structure for maps indexed by intervals of integers, but using this structure to map intervals to a boolean gives an implementation for sets of integers which it is instructive to compare to diets.

##### **4.1. Basic idea**

Let us assume for simplicity that we are only interested in representing interval-indexed maps in which the keys, in addition to being non-overlapping, are contiguous and always cover the same definition interval. We will always use the interval 0..100 in the examples. For consistency with this invariant, the function for creating a new map cre-

ates a map with a single binding from the key  $0..100$  to the provided value: `val new_map : 'a -> 'a tree.`

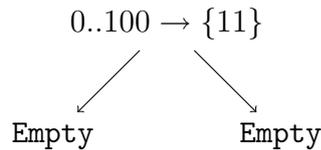


Figure 3: Tree for the map  $0..100 \rightarrow \{11\}$

Such a map is represented in our data structure by a single node with empty subtrees (see Fig. 3). From this point onwards, for the sake of readability, when both subtrees of a node are empty, we omit them from the figure.

A function `add` allows to change part of an existing map:

```
val add : int*int -> 'a -> 'a tree -> 'a tree
```

Let us consider what happens when calling `add (20,30) {12}` on the initial tree created above. The resulting map has bindings  $0..19 \rightarrow \{11\}$ ,  $20..30 \rightarrow \{12\}$ , and  $31..100 \rightarrow \{11\}$ . In fact, we must not confuse the bindings at  $0..19$  and  $31..100$  for bindings containing the value  $\{11\}$ : they are both remaining parts of a binding that originally spanned the interval  $0..100$  and has been partially overwritten. In order to make this distinction explicit, we denote the map as  $0..19 \rightarrow \{11\}_{0..100}$ ,  $20..30 \rightarrow \{12\}$ , and  $31..100 \rightarrow \{11\}_{0..100}$

It is obvious how to arrange these bindings in a tree for easy retrieval: with the lower bindings on the left-hand-side and the higher bindings on the right-hand-side. What is not obvious is deciding which node goes on top so that the tree ends up balanced or nearly balanced.

One possibility is to record the height of the trees and to build trees that are balanced by construction. This amounts to using AVL trees [1] with the natural order on intervals, which we proposed as an ad-hoc solution in Sect. 3.1. Unfortunately, the re-balancing operations cause the creation of physically different trees that contain the same sets of bindings, that is, loss of mergeability.

Another possibility, which preserves mergeability, relies on the same idea that underlies Patricia trees. In Patricia trees, there is a static hierarchy for deciding which node goes above the other, and this static hierarchy ensures trees are balanced or almost balanced without any re-balancing operations. We similarly define a static ordering on intervals that tells which node must be placed above the others. Intuitively, the interval containing the multiple of the largest power of two is put at the top. Like the ordering in Patricia trees, this ordering has the bounded chain length property (the bound is the 2-logarithm of the definition interval's width, give or take a couple of units).

We now define the ordering more formally.

**Definition 1** (Rank of an interval). The rank of an interval  $I$  is defined as:

$$\begin{aligned} \text{rank}(I) = & (k \mid \exists x \in I \ x \bmod 2^k = 0 \wedge \\ & (\forall y \in I \forall k' \ y \bmod 2^{k'} = 0 \Rightarrow k' \leq k)) \end{aligned}$$

In particular, it follows from Def. 1 that any interval containing 0 has any rank, because  $\forall k, \ 0 \bmod 2^k = 0$ . We take as convention that the interval containing 0 will always have the highest rank of all intervals contained in a tree (this special value will be denoted as  $\infty$ ). From the definition of a rank, we can now define a strict partial order on our intervals.

**Definition 2** (Strict partial order over intervals,  $\succ_i$ ). Let  $I_1$  and  $I_2$  be intervals.

$$I_1 \succ_i I_2 \iff \text{rank}(I_1) > \text{rank}(I_2)$$

This strict partial order has the additional property that two contiguous intervals are always comparable.

**Lemma 1** (Comparability of adjacent intervals). *Let  $I_1$  and  $I_2$  be (non-equal) adjacent intervals. Either  $I_1 \succ_i I_2$  or  $I_2 \succ_i I_1$ .*

*Proof.* Let  $I_1 = a_1..b_1$  and  $I_2 = a_2..b_2$  with  $a_2 = b_1 + 1$ .  
 Now assume  $\text{rank}(I_1) = \text{rank}(I_2)$ , i.e.  $\exists n_1, n_2, k, n_1 2^k \in I_1 \wedge n_2 2^k \in I_2$  with  $n_1 < n_2$ .

We know that  $\exists n, n_1 \leq 2n \leq n_1 + 1 \leq n_2$  (one of two consecutive integers is even). Hence  $\exists n, n_1 2^k \leq 2n 2^k = n 2^{k+1} \leq n_2 2^k$ . Either  $I_1$  or  $I_2$  contains the value  $n 2^{k+1}$  as they are contiguous, therefore either  $\text{rank}(I_1)$  or  $\text{rank}(I_2)$  is  $k + 1$ , contradicting our first assumption.  $\square$

Fig. 4 shows the tree representation ordered with the rank function for the following bindings:  $0..19 \rightarrow \{11\}_{0..100}$ ,  $20..30 \rightarrow \{12\}$ ,  $31..100 \rightarrow \{11\}_{0..100}$ .

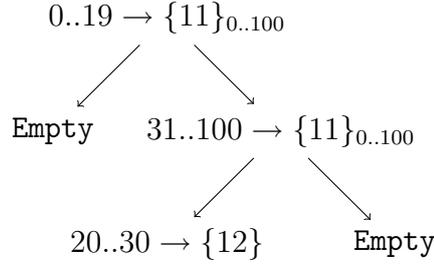


Figure 4: Ordering nodes according to rank

In Fig. 4, the interval  $0..19$  is at the root because it contains 0. The interval containing 0 is always at the root by convention. The interval  $31..100$  contains  $64 = 2^6$  whereas the interval  $20..30$  contains  $24 = 3 * 2^3$ , therefore the former goes above the latter. The  $20..30$  binding ends up as left child of the  $31..100$  binding.

#### 4.2. Pretty well mergeable

In Sect. 3.2, we claimed that Patricia trees in which lower bounds of intervals were used as ids did not fit the mergeability constraint because during a recursive descent on two separate trees, corresponding subtrees would have different definition domains. Trying to merge the trees in Fig. 1 and Fig. 2, for instance, one encounters the problem that the

corresponding leftmost subtrees contains the bindings for 0..47 in one tree and 0..31 in the other. To merge these subtrees in practice, it is necessary to borrow the contents of the 32..47 range from the context of the second tree.

The attentive reader may have noticed that the solution we are proposing appears to suffer from a similar problem. During a recursive descent of separate rangemaps (for instance in the context of a merge operation), the definition domains for encountered subtrees may differ too. This is in fact unavoidable, as the partitioning of the definition domain into intervals may not match at all between the two trees. With rangemaps, it may too be necessary to patch the narrower subtree to extend its definition domain to the same size as the other, in effect borrowing bindings from its context.

The important difference is that in rangemaps, with the interval ordering that we defined, the “context” in which it is necessary to look for bindings to borrow is limited. Specifically, only the most immediate ancestor node from which we descended to the right, or the most immediate ancestor from which we descended to the left, to the exclusion of any other, need to be borrowed from. Both on the left-hand-side and on the right-hand-side, there is at most one binding to move temporarily to the narrowest subtree to equalize it. Because of the way the interval ordering works, it is never necessary to look further than this parent.

To illustrate this claim, consider the example of two corresponding rank 5 subtrees  $t_1$  and  $t_2$  (let us assume each subtree’s root binding contains 32). The subtree that reaches the farthest to the right,  $t_1$ , can not span past 63. On the other hand,  $t_2$ ’s parent is the binding that contains 64, and therefore, it is not necessary to look elsewhere than in this parent node to get a piece of binding that equalizes the definition domain of  $t_2$  with that of  $t_1$ .

By contrast, in the solution from Sect. 3.2, an arbitrary number of bindings may have to be borrowed to equalize the definition domains.

### 4.3. Relative subtrees

An orthogonal optimization, mentioned here for completeness, is to make all subtrees relative. The arrows between the nodes carry offsets that must be tracked when traversing the tree. The benefit obtained in exchange for this additional complexity is that sharing (see also Sect. 5.5) becomes possible within a single, repetitive map, in addition to the sharing between distinct but similar maps that other tree representations usually allow. Note that this optimization is not specific to rangemaps and can be adapted to most kinds of trees with numerical keys.

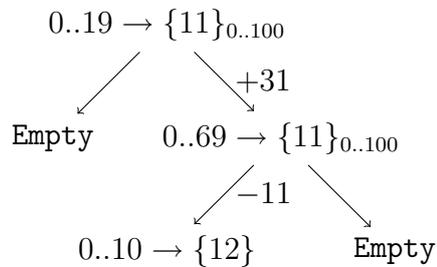


Figure 5: The same map as in Fig. 4, represented with relative subtrees

### 4.4. Automatic stitching of identical adjacent bindings

Yet a different, complementary optimization for compact representation of repetitive trees is to automatically stitch adjacent bindings to the same value into a single, wider binding. This is in the spirit of what Erwig proposed for diets [4]. However, because our data structure is a map, requirements for stitching are more sophisticated. Values must be identical, but also be stored with the same width, and the rightmost one must start where the leftmost one ends.

To make this optimization easily applicable, we store the information about the original span of the binding (that we denoted as a subscript  $\{11\}_{0..100}$  in previous examples) in the form of a binding width (for this example, 101) and an offset (for this example, 0).

An example of binding that can be stitched to this one is  $101..201 \rightarrow \{11\}$ . It starts where the  $0..100$  binding finishes, and it contains the same value (with the same width). With the (width, offset) representation, the criterion for recognizing that it is stitchable is that values, widths, and offsets are identical for both bindings, and in addition, that the stitching point (101) is congruent to offset modulo width (that is, the point of stitching is actually a point where a value ends and a new one can start). Stitching occurs in particular when adding a new node to an already existing rangemap: this is illustrated in Fig. 7.

In the use case of abstract interpretation of C programs, it is clear why it is undesirable to omit the last condition above: on a little-endian architecture, a binding  $0..0 \rightarrow \{1, 258\}_{0..3}$  may be the abstract result of taking the first byte of the concrete 32-bit value 1. The binding  $1..3 \rightarrow \{1, 258\}_{0..3}$  may be the abstract result of taking the last three bytes of the concrete 32-bit value 258. Stitching these two bindings together into a single binding  $0..3 \rightarrow \{1, 258\}$  would be incorrect: the value contained in these four bytes in a concrete execution, 257, would not be represented by the abstract value  $\{1, 258\}$  resulting from the stitching.

On the other hand, when the stitching occurs at a point where a binding ends and another starts, no incorrectness results from stitching them together. Indeed, these bindings are still considered as different bindings after the stitching has occurred.

In effect, the optimization proposed in this section consists in enforcing the invariant that “no two adjacent stitchable bindings coexist in the rangemap”. Therefore, whenever a binding is added, or changed, in a rangemap, the adjoining bindings must be checked for stitching possibilities.

## 5. Implementation notes

We have implemented proof-of-concept rangemaps, and we hope to soon be able to substitute this implementation to the existing, ad-hoc interval-indexed maps in the value analysis of Frama-C [5]. This section describes the OCaml implementation. Most functions in the implementation of rangemaps follow a divide-and-conquer pattern. There-

fore, they can be cached, in the hope that partial results from previous similar computations can be reused. In the context of Frama-C’s value analysis, rangemaps can be expected to exhibit sharing both because of the existence of maps that are slight variations of each other, and because of repetitive bindings within a single map. Caching allows to take advantage of spatial sharing to gain in execution time.

### 5.1. *Datatype*

Rangemaps are trees built from the following algebraic datatype:

```

type t =
  | Empty
  | Node of Int.t *
    (* max: min is always implicitly zero *)
    Int.t * t *
    (* offset_left * subtree_left *)
    Int.t * t *
    (* offset_right * subtree_right *)
    Int.t * Int.t * V.t
    (* offset * modulo * value *)

```

Tree nodes carry the following information:

- the length of the interval (max+1);
- where (offset\_left, offset\_right) and what (subtree\_left, subtree\_right) its left and right children are. These offsets are computed as the difference between the lower bound of the child and that of the parent;
- the data bound to the interval, i.e a value repeated each modulo starting from offset.

### 5.2. *Construction of values*

Construction of values of the type `t` is done exclusively using so-called “smart constructors”. This enforces various invariants including

that of Sect. 4.4. Only the module implementing the rangemaps has direct access to the algebraic constructors ( $\tau$  is abstract but could as well have been declared private).

Huet’s zippers [8] are also used in the implementation to allow efficient navigation from a node to its neighbors. Another use of zippers takes place in the stitching phase of Sect. 4.4: they represent the context in which the subtrees of the node to be stitched should be re-attached (see also Fig. 7).

We will concentrate here on the `add_binding` function of the module, which internally calls a smart `make_node`. Their signatures, as implemented, are:

```

val add_binding :
    int64 -> int64 -> int64 -> Int.t -> Int.t ->
    V.t -> t -> int64 * t
(* [current_tree_offset] ->
   [min] -> [max] -> [off] -> [modu] -> [value] ->
   [tree] -> [new_current_tree_offset] * [current_tree]
*)

val make_node :
    int64 -> Int.t -> Int.t -> t -> Int.t -> t ->
    Int.t -> Int.t -> V.t -> int64 * t
(* [current_tree_offset] -> [max] ->
   [offset_left_subtree] -> [left_subtree] ->
   [offset_right_subtree] -> [right_subtree] ->
   [off] -> [modu] -> [value] ->
   [current_new_tree_offset] * [new_tree]
*)

```

Let us illustrate how the smart `add_binding` operates on the rangemap shown in Fig. 6, which represents the following sequential intervals:  $0..19 \rightarrow \{11\}$ ,  $20..30 \rightarrow \{12\}$ ,  $31..65 \rightarrow \{11\}$ ,  $66..80 \rightarrow \{14\}$ ,  $81..88 \rightarrow \{15\}$ ,  $89..100 \rightarrow \{13\}$  with respective ranks  $\infty$  (by convention), 3 ( $24 = 3 * 2^3$ ), 6 ( $64 = 2^6$ ), 5 ( $80 = 5 * 2^4$ ), 3 ( $88 = 11 * 2^3$ ), 5 ( $96 = 3 * 2^5$ ). As an added hypothesis, we suppose all

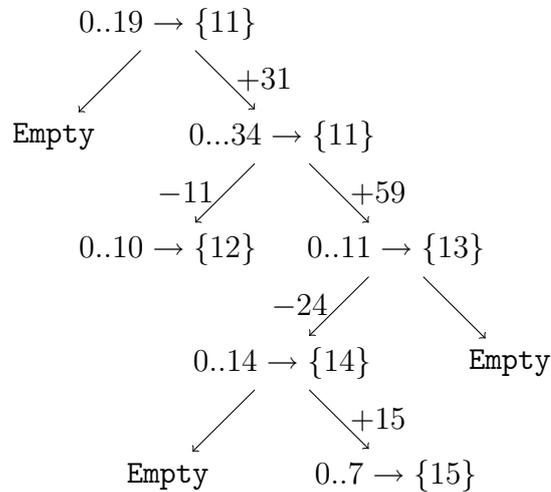


Figure 6: Initial tree before addition of a binding

off and modu are given in such a way that the adjacent intervals of this rangemap can be stitched together provided they hold the same value.

Suppose that we now want to add the following binding  $31..65 \rightarrow \{14\}$  to this rangemap of Fig. 6. The operation can be decomposed as follows (see also Fig. 7):

- 1) Find the correct spot according to rank (Def. 1) where the new node should be;
- 2) See if it can be stitched together with some node of its subtrees;
- 3) Call `make_node` and `rezip` if needed, stitch if needed.

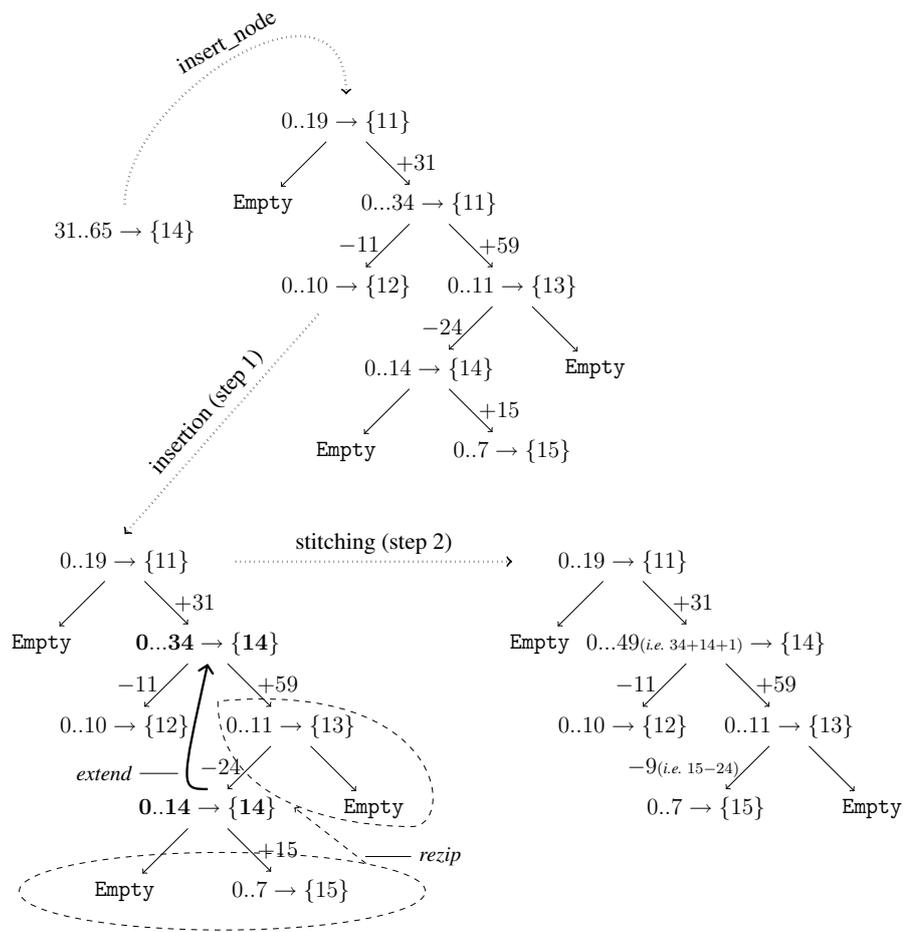


Figure 7: Inserting a node: stitching illustrated

### 5.3. Merging rangemaps

From the get-go, we were aiming at a mergeable structure and our module naturally contains a suitable function with the following immediate signature:

```
val merge: int64 -> t -> int64 -> t -> int64 * t
(* [offset_t1] -> [t1] -> [offset_t2] -> [t2] ->
   [current_new_tree_offset] * [new_tree]
*)
```

This function, apart from its primary importance, is not unnecessarily complicated to implement. Actually, it makes extensive use of the `make_node` function and of recursive calls to itself. A simplified version can be informally stated as follows, assuming  $n_1$  and  $n_2$  are the current nodes of the two trees  $t_1$  and  $t_2$  to be merged:

1) If  $I_{n_1} \cap I_{n_2} = \emptyset$ :

Let  $n_{max}$  be the highest ranked node between  $n_1$  and  $n_2$  and  $n_{min}$  the other one, and  $t_{max}$  and  $t_{min}$  the respective trees they belong to. Let also  $subt_{max}^+$  be the subtree  $n_{min}$  should be included in: it is the left subtree of  $t_{max}$  if  $max(n_{min}) < min(n_{max})$ , the right one otherwise. Let  $subt_{max}^-$  be the other unchanged subtree of  $t_{max}$ .

Merge  $subt_{max}^+$  and  $t_{min}$  into a new tree  $t'$ .

Make a new tree from  $n_{max}$ ,  $t'$  and  $subt_{max}^-$ .

2) Otherwise, let  $I = I_{n_1} \cap I_{n_2}$  and compute the value(s) it contains according to the ones contained in  $I_{n_1}$  and  $I_{n_2}$ . Let  $I_{n_1}^<, I_{n_1}^> = I_{n_1} \setminus I$ . These two new intervals represent the lower part of  $I_{n_1}$  not in  $I$  and the upper part of  $I_{n_1}$  not in  $I$ . Let similarly  $I_{n_2}^<, I_{n_2}^> = I_{n_2} \setminus I$ .

Add  $I_{n_1}^<$  to the left subtree of  $t_1$  and  $I_{n_1}^>$  to the right subtree of  $t_1$ . Do the same for  $I_{n_2}^<, I_{n_2}^>$  and  $t_2$ .

Merge both new left subtrees and merge both new right subtrees.

Make a new smart tree with  $I$  and the results of the previous recursive calls.

Note that the values mapped to the intervals are not changed except when the two trees have overlapping intervals.

#### 5.4. *Caching*

As noted in Sect. 4.4, the implemented functions often need to access the rightmost and leftmost bindings of a subtree (i.e. those directly on the right and left-hand side of the current node if the interval is looked at linearly). This is right now naively done by recursively descending the subtree. Another solution would consist in borrowing ideas from monoid caching trees [7] and have fingers [6] pointing at the rightmost left and leftmost right children.

However, this solution was not chosen. We chose to save the two words necessary at each node to record the rightmost and leftmost bindings. In our context, the space savings can more efficiently be used to create caches for high-level operations, even if in the case of a cache miss, the operation takes a little longer because of the logarithmic access to these leftmost and rightmost bindings.

#### 5.5. *Sharing*

The representation of relative subtrees described in Sect. 4.3 allows maximal sharing on subtrees of the data structure: rangemaps are actually DAGs and not trees. The implemented version is on the right-hand side of Fig. 8. Note that Empty subtrees are not drawn as shared although they actually are: this is automatically done by OCaml and not in the implementation of rangemaps.

The trees in Fig. 8 represent shared and unshared rangemaps for the sequential interval:  $0..19 \rightarrow \{11\}$ ,  $20..29 \rightarrow \{12\}$ ,  $30..49 \rightarrow \{11\}$ ,  $50..59 \rightarrow \{12\}$ ,  $60..100 \rightarrow \{11\}$  with respective ranks  $\infty, 3, 5, 3, 6$ .

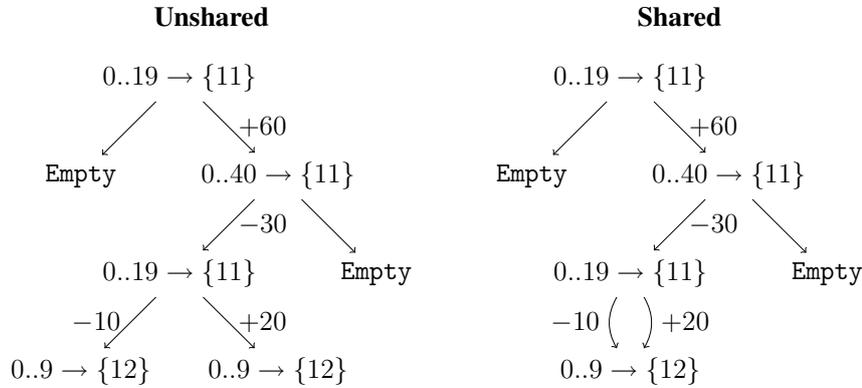


Figure 8: Unshared vs. shared version of the same tree

## 6. Conclusion and further work

In this article, we have introduced rangemaps, a data structure to represent persistent maps indexed by intervals. We have shown the unique properties of rangemaps and highlighted details of the current prototype implementation.

The next step is to replace the existing ad-hoc data structure for representing char arrays in Frama-C’s value analysis<sup>1</sup> by rangemaps. We expect this will increase the efficiency of the value analysis both in terms of memory used and speed. We are looking forward to the possibility to run benchmarks comparing the current ad-hoc representation and the representation based on rangemaps.

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1. The simplest way to handle heterogeneous pointer casts in an abstract-interpretation based static analyzer is to decide from the start to treat every memory location as a char array. In fact, the value analysis treats every memory location as a bit array as part of its treatment for bitfields in C.

gile Prevosto et Julien Signoles de nous avoir prêté main forte lors de la relecture des versions antérieures de ce document.

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