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# Improving Performance of the Datalog Engine in Flix

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# **Abstract**

# Acknowledgments

► Write acknowledgments here... ◀

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# Chapter 1

## Introduction

Datalog is a simple yet powerful declarative logic programming language. Programs written in a logic programming language are collections of logical statements and rules. This allows the programmer to solve problems by stating facts and rules about the problem domain rather than by giving step-by-step algorithmic instructions.

The Flix programming language is a new functional, declarative, and strongly typed programming language on the JVM. One of its main features is the first-class support for Datalog, allowing programmers to write Datalog programs directly in Flix.

The Datalog engine in Flix is implemented as a library using semi-naïve evaluation as its evaluation strategy, as most modern Datalog engines do. The Datalog engine in Flix has two parts: it compiles the given Datalog program into an intermediate representation, RAM (Relational Algebra Machine), represented as Flix values, and then it interprets the RAM program statement by statement. The semi-naïve evaluation is reflected in the way the RAM program is constructed. The generated RAM program is highly parallelizable, but it requires a concurrent data structure.

The goal of this thesis is to improve the performance of the Datalog engine in Flix. The main contribution of this thesis is the parallelization of the evaluation, which includes three parts: (1) a benchmark suite is proposed for evaluating the performance of the engine, (2) a new parallel statement is added to RAM, and (3) a concurrent data structure is implemented to enable the interpretation of the parallel statement. Specifically, a concurrent B+-tree is implemented to replace the Red-black tree used in the current implementation. The thesis statement is as follows: A concurrent B+-tree would make Datalog evaluation in Flix more efficient.

The rest of the thesis is organized as follows. In Chapter 2, we give a brief introduction to Datalog, including the syntax and semantics of the language, as well as the semi-naïve evaluation and how it leads to parallelism. In Chapter 3, we first look at how the semi-naïve algorithm is implemented in Flix, specifically how the RAM program is constructed and interpreted. We then examine how parallelism can be introduced. Lastly, we discuss the benchmark suite. In Chapter 4, we look at the concurrent B+ tree implementation that enables the parallelism. In Chapter 5, we evaluate the performance of the trees we implement and the engine using these trees.

# Chapter 2

## Datalog

### 2.1 Datalog Syntax

A Datalog program consists of *facts*, which are statements that are held to be true, and *rules*, which describe how to deduce new facts from known facts. One can think of facts as rows in a relational database table and rules as queries that can be run on the database or views that can be materialized.

**Example 2.1.** The following database stores the edges of a directed graph by means of facts of the form `edge(v1, v2)`, meaning that there is an edge from vertex `v1` to vertex `v2` in the graph:

```
edge("a", "b").  
edge("b", "c").  
edge("c", "d").
```

Below are two rules to compute the transitive closure of a graph stored by means of `edge` facts:

```
tc(X, Y) :- edge(X, Y).  
tc(X, Y) :- edge(X, Z), tc(Z, Y).
```

Intuitively, the first rule above says that if there is an edge from vertex `X` to vertex `Y`, then `(X, Y)` belongs to the transitive closure of the graph. The second rule says that if there is an edge from `X` to `Z` and there exists a vertex `Y` such that `(Z, Y)` belongs to the transitive closure, then `(X, Y)` belongs to the transitive closure as well.

We now give the formal syntax of Datalog programs. A *Datalog rule*  $r$  is of the form:

$$A_0 : -A_1, \dots, A_n.$$

where  $n \geq 0$ , the  $A_i$ 's are *atoms*, and every atom is of the form  $p(t_1, \dots, t_m)$ , where  $p$  is a *predicate symbol* and the  $t_i$ 's are *terms*. A *term* is either a *variable* or a *constant*. In the example above, `edge` and `tc` are predicate symbols, `"a"`, `"b"`, `"c"`, `"d"` are constants, and `X`, `Y`, `Z` are variables. Every variable appearing in  $A_0$  (the *head* of a rule) also appears in at least one of the  $A_1, \dots, A_n$  (the *body* of



a rule). This requirement is called *safety* and is used to avoid rules yielding infinite sets of facts from a finite set of facts. The intuitive meaning of the Datalog rule above is that if the atoms  $A_1, \dots, A_n$  are true, then the atom  $A_0$  is also true. A *fact* is a rule with an empty body, i.e., a rule of the form  $A_0 : -$ , which can be written as  $A_0$ .

A *Datalog program* is a finite set of Datalog rules. The *definition* of a predicate symbol  $p$  in a program  $P$ , denoted  $def(p, P)$ , is the set of rules of  $P$  having  $p$  in the head atom. Recall that a database can be seen as a finite set of facts. In the context of logic programming, all the knowledge (facts and general rules) is usually contained in a single logic program. We consider two sets of Datalog rules:

1. A set of facts  $D$  that represent tuples (rows) of a database.
2. A Datalog program  $P$  whose rules define new relations (or "views") from the database.

$D$  is called the *Extensional Database* (EDB) and  $P$  is called the *Intensional Database* (IDB). We will refer to  $D$  as the database and  $P$  as the Datalog program. Thus, predicate symbols are partitioned into two disjoint sets: *base* (or EDB or *extensional*) and *derived* (or IDB or *intensional*). The definition of base predicate symbols is stored in  $D$ . Base predicate symbols can appear in the body of rules in  $P$  but not in the head. Derived predicate symbols cannot appear in  $D$  and their definition is in  $P$ . We will use  $P_D$  to denote  $P \cup D$ .

## 2.2 Datalog Semantics

We give the least-fixpoint semantics of Datalog, which defines the outcome of the program in an operational way, based on repeatedly applying the rules of the program until no new facts can be derived and thus a fixpoint is reached. It is well known that this semantics is equivalent to the model-theoretic semantics as well as the proof-theoretic semantics [1].

The fixpoint semantics is given in terms of an operator called the *immediate consequence operator*, which derives new ground atoms starting from known ground atoms, using the rules of the program. Namely,  $T_{P_D}$  takes as input a set of ground atoms  $I$ , applies the rules, and returns as output a set of ground atoms  $T_{P_D}(I)$ , called the *immediate consequences* of  $I$  w.r.t.  $P_D$ . We say that a set of ground atoms  $I$  is a *fixpoint* of  $T_{P_D}$  if  $T_{P_D}(I) = I$ .

It is easy to see that  $T_{P_D}$  is monotonic, that is, if  $I_1 \subseteq I_2$ , then  $T_{P_D}(I_1) \subseteq T_{P_D}(I_2)$  for any sets of ground atoms  $I_1$  and  $I_2$ . By the Knaster-Tarski theorem, since  $T_{P_D}$  is monotonic, it has a least fixpoint (that is, a fixpoint that is included in any other fixpoint), which we denote as  $lfp(T_{P_D})$ . The fixpoint semantics of  $P_D$  is given by the least fixpoint  $lfp(T_{P_D})$ .

**Example 2.2.** Consider the Datalog program in Example 2.1. Applying  $T_{P_D}$  yields the following:

$$\begin{aligned}
I_1 &= T_{P_D}(\emptyset) = \{edge("a", "b"), edge("b", "c"), edge("c", "d")\}, \\
I_2 &= T_{P_D}(I_1) = I_1 \cup \{tc("a", "b"), tc("b", "c"), tc("c", "d")\}, \\
I_3 &= T_{P_D}(I_2) = I_2 \cup \{tc("a", "c"), tc("b", "d")\}, \\
I_4 &= T_{P_D}(I_3) = I_3 \cup \{tc("a", "d")\}, \\
I_5 &= T_{P_D}(I_4) = I_4.
\end{aligned}$$

Thus,  $I_4$  is the least fixpoint of  $T_{P_D}$ .

## 2.3 Datalog Evaluation

The fixpoint semantics immediately leads to an algorithm to evaluate Datalog programs, called *naïve evaluation*. To define the algorithm, we first define a function called *Eval-Rule* that takes as input a rule  $r$  and a set of ground atoms  $I$  containing the relations of the predicates appearing in the body of the rule, and returns the set of ground atoms that are the immediate consequences of  $I$  w.r.t.  $r$ . The function *Eval-Rule* can be defined precisely using relational algebra, but we omit the details here, as the intuition is simple and clear.

**Example 2.3.** Let  $r$  be the second rule in Example 2.1 and  $I = I_3$  in Example 2.2, calling *Eval-Rule* on rule  $r$  and relations *edge* and *tc* yields the following:

$$Eval\text{-}Rule(r, edge, tc) = \{tc("a", "c"), tc("b", "d"), tc("a", "d")\}.$$

We then define another function called *Eval* that evaluates a set of rules whose predicates in the head are the same. Consider a program  $P$  and a database  $D$ . Let  $R_1, \dots, R_n$  be the base relations and  $P_1, \dots, P_m$  be the derived relations. For each derived predicate symbol  $p_i$ , we define  $Eval(p_i, R_1, \dots, R_n, P_1, \dots, P_m)$  as the union of calling *Eval-Rule* over all rules with  $p_i$  in the head.

Now, given a Datalog program  $P$  and a database  $D$ , where the base predicate symbols are  $r_1, \dots, r_n$ , the derived predicate symbols are  $p_1, \dots, p_m$ , and the base relations are  $R_1, \dots, R_n$ , Algorithm 2.3 performs the evaluation of  $P$  over  $D$ .

One shortcoming of the naïve evaluation is that in each iteration, all tuples computed in the previous iterations are recomputed. For example, in Example 2.3, the tuples  $tc("a", "c")$  and  $tc("b", "d")$  are already computed in the previous iteration (they are already in  $I_3$ ). The problem is that in the naïve evaluation, in each iteration, the tuples of the derived relations from earlier iterations are used in *Eval-Rule*. The solution to this problem is called *semi-naïve evaluation*.

We start by introducing an incremental version of the *Eval-Rule* function. Consider a rule  $r$  of the form

$$A_0 : -A_1, \dots, A_n.$$

---

**Algorithm 1** Naïve Evaluation

---

```
1: for  $i \leftarrow 1$  to  $m$  do
2:    $P_i \leftarrow \emptyset$ 
3: repeat
4:   for  $i \leftarrow 1$  to  $m$  do
5:      $P'_i \leftarrow P_i$ 
6:   for  $i \leftarrow 1$  to  $m$  do
7:      $P_i \leftarrow \text{Eval}(p_i, R_1, \dots, R_k, P'_1, \dots, P'_m)$ 
8: until  $P_i = P'_i$  for all  $1 \leq i \leq m$ 
9: return  $P_1, \dots, P_m$ 
```

---

and assume to have a relation  $R_i$  and an “incremental” relation  $\Delta R_i$  for each atom  $A_i$  in the body of the rule. We define the incremental version of *Eval-Rule* as follows:

$$\begin{aligned} \text{Eval-Rule-Incr}(r, R_1, \dots, R_n, \Delta R_1, \dots, \Delta R_n) = \\ \bigcup_{1 \leq i \leq n} \text{Eval-Rule}(r, R_1, \dots, R_{i-1}, \Delta R_i, R_{i+1}, \dots, R_n). \end{aligned}$$

We can now define the *semi-naïve evaluation* as shown in Algorithm 2.3.

---

**Algorithm 2** Semi-naïve Evaluation

---

```
1: for  $i \leftarrow 1$  to  $m$  do
2:    $\Delta P_i \leftarrow \text{Eval-rule}(p_i, R_1, \dots, R_k, \emptyset, \dots, \emptyset)$ 
3:    $P_i \leftarrow \Delta P_i$ 
4: repeat
5:   for  $i \leftarrow 1$  to  $m$  do
6:      $\Delta P'_i \leftarrow \Delta P_i$ 
7:   for  $i \leftarrow 1$  to  $m$  do
8:      $\Delta P_i \leftarrow \text{Eval-rule-incr}(p_i, R_1, \dots, R_k, P_1, \dots, P_m, \Delta P'_1, \dots, \Delta P'_m)$ 
9:      $\Delta P_i \leftarrow \Delta P_i - P_i$ 
10:  for  $i \leftarrow 1$  to  $m$  do
11:     $P_i \leftarrow P_i \cup \Delta P_i$ 
12: until  $\Delta P_i = \emptyset$  for all  $1 \leq i \leq m$ 
13: return  $P_1, \dots, P_m$ 
```

---

It is obvious that the semi-naïve evaluation is parallelizable. There are two levels of parallelism in Algorithm 2.3. First, each invocation of *Eval-Rule-Incr* can be done in parallel, that is, the body of the loop from line 9 to line 12. Second, as per the definition of *Eval-Rule-Incr*, the invocations of *Eval-Rule* called by *Eval-Rule-Incr* can also be done in parallel. This is the main idea of the parallelization of this thesis. Also, the body of the loop from line 1 to line 4 can be done in parallel.

## Chapter 3

# Datalog Evaluation in Flix

In Flix, the Datalog engine is implemented as a library. When a Flix program is executed, the Datalog rules are compiled into an intermediate representation called RAM (Relational Algebra Machine), which is then interpreted statement by statement. The RAM program is constructed in a way that reflects the semi-naïve evaluation algorithm. In this chapter, we first look at how the semi-naïve algorithm is implemented in Flix, specifically how the RAM program is constructed and interpreted. We then examine how parallelism can be introduced. Lastly, we discuss the benchmark suite.

### 3.1 RAM

We will not introduce the definition of RAM as Flix code here. Instead, we will give examples of stringified RAM programs. We will denote  $P_i, \Delta P_i, \Delta P_i'$  in Algorithm 2.3 as *Full* relations, *Delta* relations, and *New* relations.

**Example 3.1.** Consider the Datalog program in Example 2.1. We have one base predicate `edge` and one derived predicate `tc`. The initialization part of the semi-naïve evaluation algorithm (lines 1 to 4) will be translated into the following RAM program:

```
// tc(x, y) :- edge(x, y).
search 0$ ∈ edge do
  project (0$[0], 0$[1]) into Δtc
end;
// tc(x, z) :- tc(x, y), edge(y, z).
search 0$ ∈ tc do
  query {1$ ∈ edge | 1$[0] == 0$[1]} do
    project (0$[0], 1$[1]) into Δtc
  end
end;
merge Δtc into tc;
```

The invocation of *Eval-Rule* for the derived predicate `tc` is translated into the above two RAM statements. The translation is on a rule-by-rule basis and is mechanical. Variables are represented by integers followed by a dollar sign and they hold a tuple of a relation. Each atom in the body is translated into a *search*. The *search* statement is used to iterate over the tuples of a relation. The *query* statement is like *search* but with a filter of the form `x[0]==a, x[1]==b, x[2]==c, ...`. The difference between using a *query* and using a *search* with *if* conditions is that the former is more efficient, as the tuples are stored using trees and the query is interpreted using range queries on the trees that avoid unnecessary walking over the tree. We will see more about this in Chapter 4.

The main loop of the semi-naïve evaluation algorithm (lines 5 to 16) will be translated into the following RAM program:

```

until( $\Delta tc == \emptyset$ ) do
  purge  $\Delta tc'$ ;
  //  $tc(x, y) :- Edge(x, y).$ 
  //  $tc(x, z) :- tc(x, y), Edge(y, z).$ 
  search  $0\$ \in \Delta tc$  do
    query { $1\$ \in Edge \mid 1\#[0] == 0\#[1]$ } do
      if( $(0\#[0], 1\#[1]) \notin tc$ ) then
        project ( $0\#[0], 1\#[1]$ ) into  $\Delta tc'$ 
      end
    end
  end
  end;
  merge  $\Delta tc'$  into  $tc$ ;
   $\Delta tc := \Delta tc'$ 
end

```

In this program, `tc` is the only derived predicate. Calling *Eval-Rule-Incr* for `tc` includes the translation of both rules. However, there is no derived predicate in the body of the first rule, so the first rule is translated to an empty statement. For the second rule, the derived predicate `tc` appears in one atom in the body, so the *search* statement is used to iterate over the *New* relation  $\Delta tc$ . In Algorithm 2.3, line 11 is used to compute the *New* relations by set difference. In the implementation, it is done by the *if* operation to check if the tuple is already in the *Full* relation.

We will show one more example of a more complex Datalog program, where derived predicates appear more than once in the body of a rule, which will make the translation of *Eval-Rule-Incr* more interesting.

**Example 3.2.** The following Datalog program performs a point-to analysis of variables in a C program [4]. The instructions of the program can be translated into *EDB* relations as follows:

```

AddressOf(a, b) representing  $a = \&b$ 
Assign(a, b) representing  $a = b$ 
Load(a, b) representing  $a = *b$ 
Store(a, b) representing  $*a = b$ 

```

The point-to analysis can now be done using a Datalog program storing pairs of variables `a` and `b` in the intentional relation `PointsTo` if `a` may point to `b`. The program contains the following four rules with the same head:

```

PointsTo(x, y) :- AddressOf(x, y).
PointsTo(x, y) :- Assign(x, z), PointsTo(z, y).
PointsTo(x, w) :- Load(x, z), PointsTo(z, y), PointsTo(y, w).
PointsTo(x, y) :- Store(z, w), PointsTo(z, x), PointsTo(w, y).

```

We pay attention to the load rule (the third rule). *Eval-Rule* (line 2 of Algorithm 2.3) for this rule will be translated into the following single RAM statement:

```

search 0$ ∈ Load do
  query {1$ ∈ PointsTo | 1$[0] == 0$[1]} do
    query {2$ ∈ PointsTo | 2$[0] == 1$[1]} do
      project (0$[0], 2$[1]) into ΔPointsTo
    end
  end
end
end;

```

Because there are two derived predicates in the body, *Eval-Rule-Incr* (line 11 of Algorithm 2.3) for this rule will be translated into two RAM statements. This corresponds to the union operation in the definition of *Eval-Rule-Incr*:

```

search 0$ ∈ Store do
  query {1$ ∈ ΔPointsTo | 1$[0] == 0$[0]} do
    query {2$ ∈ PointsTo | 2$[0] == 0$[1]} do
      if((1$[1], 2$[1]) ∉ PointsTo) then
        project (1$[1], 2$[1]) into ΔPointsTo'
      end
    end
  end
end;
search 0$ ∈ Store do
  query {1$ ∈ PointsTo | 1$[0] == 0$[0]} do
    query {2$ ∈ ΔPointsTo | 2$[0] == 0$[1]} do
      if((1$[1], 2$[1]) ∉ PointsTo) then
        project (1$[1], 2$[1]) into ΔPointsTo'
      end
    end
  end
end;
end;

```

## 3.2 Parallelization

It is now clear that each statement translated from a rule can be executed in parallel. Consider Example 3.1; the two *search* statements in the *Eval-Rule-Incr* for the load rule can be executed in parallel. In fact, the assign rule will be translated into one statement, the load rule and the store rule will both be translated into two statements. All five statements can be executed in parallel. We mentioned in the previous chapter that there are two levels of parallelism in the semi-naïve

evaluation algorithm. This nested parallelism is automatically spread out in the way the RAM program is constructed.

The first thing to do is to add a new type of statement to RAM, called `Par` (Parallel), in contrast to the existing `Seq` (Sequential) statement:

```
pub enum RamStmt[v] {
  case Insert(RelOp[v])
  case Merge(RamSym[v], RamSym[v])
  case Assign(RamSym[v], RamSym[v])
  case Purge(RamSym[v])
  case Seq(Vector[RamStmt[v]])
  case Par(Vector[RamStmt[v]])
  case Until(Vector[BoolExp[v]], RamStmt[v])
  ...
}
```

One thing to note is that in the previous text we refer to `search` and `query` as statements, whereas in the actual implementation they are cases of the `RelOp` enum, along with `If` and `Project`. All of them belong to the category of the `Insert` statement.

We then change the compiler to generate as many `Par` statements as possible. The new RAM program will be of the following form in general (we use `||` to denote parallel statements and `;` to denote sequential statements):

```
search 0$ ∈ ... do
  ...
end ||
search 0$ ∈ ... do
  ...
end ||
... ;
merge Δ... into ... ||
merge Δ... into ... ||
... ;
Until ... do
  search 0$ ∈ ... do
    ...
  end ||
  search 0$ ∈ ... do
    ...
  end ||
  ... ;
  merge Δ...' into ... ||
  merge Δ...' into ... ||
  ... ;
  Δ... := Δ...' ||
  Δ... := Δ...' ||
  ... ;
end
```

When interpreting the RAM program, we will fire up one thread for each statement in a `Par` statement. The problem is then how to perform concurrent operations on the relations. We have mentioned that the tuples of a relation are stored using trees. More specifically, the tuples are stored in a red-black tree, where the keys are the tuples themselves, and the values are either `Unit` or the corresponding lattice value if lattice is used in the relation.

### 3.3 Red-Black Trees

After the parallelization, there will be more than one thread trying to insert tuples into the same relation, for instance, in Example 3.1. However, red-black trees do not support concurrent insertions. One naïve solution is to include a mutex lock in the red-black tree and change all insertion methods to acquire the lock at the beginning and release the lock at the end. This solution works but is not efficient, as usually, there is more than one rule for deriving a predicate in a realistic Datalog program, and the thread will be blocked by the lock. We will not demonstrate the evaluation of this solution in later sections, but we mention that it sees 1.0x - 1.2x speedup on the realistic tests in our benchmark.

In this section, we will give a brief introduction to red-black trees. More specifically, we will look at what range query is and how it has made Datalog evaluation more efficient.

A red-black tree is a specialized binary search tree data structure. Compared to other self-balancing binary search trees, the nodes in a red-black tree hold an extra bit called "color" representing "red" and "black," which is used when re-organizing the tree to ensure that it is always approximately balanced.

Binary search trees are used to store ordered data. In our case, the data is the tuples of a relation (serving as keys), possibly associated with a lattice value (serving as values). In a binary search tree, each node stores a key and a value, and the left child of a node has a key less than the node's key, and the right child has a key greater than the node's key. This sorted property allows for an efficient range query.

A range query is a query that retrieves data in a given range. In our case, the keys are tuples, and they are sorted first by the first element, then by the second element, and so on. As we have illustrated in Example 3.1, the query statement is always of the form  $x[0]==a, x[1]==b, x[2]==c, \dots$ . When we are doing a range query, if at the current node, the first element of the key is less than  $a$ , or if the first element of the key is equal to  $a$  but the second element of the key is less than  $b$ , we can skip the left child of the current node, and so on. Likewise, we can skip the right child for some node. Only when the equality is met for all the elements in the query condition, we will visit both the left child and the right child.

### 3.4 Benchmark

We have implemented a small benchmark suite that contains four files, 13 programs in total, to evaluate the performance of the Datalog engine in Flix.



- `Closure.flix` contains four realistic programs that compute either the transitive closure or something similar (e.g., strongly connected component) on graphs.
- `Pointer.flix` contains two programs that perform point-to analysis as shown in Example 3.1. The difference between the two programs is in the order of the variables in the rule. We will discuss this in Chapter 6.
- `Intersect.flix` and `Propagate.flix` contain four and three programs respectively that are rather artificial. For example, the third program in `Propagate.flix` is as follows:

```
pub def propaget3(): Int32 =
  let p = #{
    A(slowId(x) + 1) :- Z(x), A(x).
    B(slowId(x) + 1) :- Z(x), B(x).
    C(slowId(x) + 1) :- Z(x), C(x).
    ...
    Z(slowId(x) + 1) :- Z(x), if x < 6.
  };
  let f = #{
    Z(0).
    A(0).
    B(0).
    C(0).
    ...
  };
  query p, f select x from A(x)
  |> Vector.length
```

`SlowId` is an identity function that takes a long time to compute by calling a recursive Fibonacci function inside. In the semi-naïve evaluation, each rule will produce a new one-element tuple for each relation, stopping at 6. Because we only have one rule for each predicate, adding parallelism should significantly speed up the evaluation, no matter using the naïve mutex lock solution or using an advanced concurrent tree. Indeed, when evaluating this program on a 14-core machine, we see a 6x speedup using the mutex lock solution.

## Chapter 4

# B-Trees

We have discussed in previous sections that the tuples of a relation are stored using red-black trees and we want to change to a concurrent data structure to facilitate parallelism. The new data structure should act like a set and support efficient range queries.

The B-tree and its variants are widely used and studied in database systems. Many papers have described how to perform concurrent operations on B-trees. In this section, we will give a brief introduction to B-trees and B+ trees, and then we will introduce a new variant of B-trees called  $B^{\text{link}}$  trees that support concurrent reading and writing. In the next chapter, we will evaluate the performance of the Datalog engine using B+ trees and  $B^{\text{link}}$  trees.

### 4.1 B-Trees and B+ Trees

The B-tree is a multi-way search tree. A  $m$ -way search tree is a tree where each node has at most  $m$  children, hence storing at most  $m - 1$  keys. The keys in a node are sorted, and the keys in the children of a node are in the range defined by the keys of the node.

A B-tree of order  $m$  is a tree which satisfies the following properties:

- Every node has at most  $m$  children.
- Every node, except for the root and the leaves, has at least  $\lceil m/2 \rceil$  children.
- The root node has at least two children unless it is a leaf.
- All leaves appear on the same level.
- A non-leaf node with  $k$  children contains  $k - 1$  keys.

In B-trees, every node stores a range of keys and their corresponding values. Each internal node's keys, including the root node, act as separation values which divide its subtrees, as shown in figure 4.1 For example, if an internal node has 3 child nodes (or subtrees) then it must have 2 keys:

$a_1$  and  $a_2$ . All values in the leftmost subtree will be less than  $a_1$ , all values in the middle subtree will be between  $a_1$  and  $a_2$ , and all values in the rightmost subtree will be greater than  $a_2$ .

The fourth property means that B-tree is a balanced tree. The second property ensures that each node is at least half full. This implies that two half-full nodes can be joined to make a legal node, and one full node can be split into two legal nodes (with an element pulled out and put into the parent). These properties make it possible to delete and insert new values into a B-tree and adjust the tree to preserve the B-tree properties.

A B+ tree is a variant of a B-tree where the internal nodes do not store any pointers to records, thus all pointers to records are stored in the leaf nodes. In addition, a leaf node may include a pointer to the next leaf node to speed sequential access. This makes range queries easy to implement. Since in a B+ tree, values are not stored in internal nodes, all keys appear twice in the tree: once in an internal node and once in a leaf node. This leads to a design choice to be made. In figure 4.1, if we denote all the keys stored in a subtree at pointer  $P_i$  as  $V_i$ , then any  $v \in V_i$  can either satisfy  $K_{i-1} < v \leq K_i$  or  $K_{i-1} \leq v < K_i$ . We prefer the former because it makes the implementation of search consistent. When performing a search for some  $k$  inside a node, we return the index of the first key that is greater than or equal to  $k$ . This index will always be the correct index of the pointer array to follow.

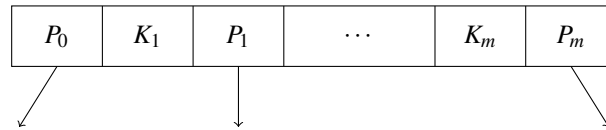


Figure 4.1: A B+ tree node

Compared to binary search trees, such as red-black trees, B-trees and its variants are more efficient in terms of I/O operations. B-trees are widely used in database systems to store indexes. The records in a database are stored in disk and pointers to the records are stored as values in the B-tree. The tree itself, serves as an index to the records, is also stored in disk. In our case, values are pointers to tuples in memory, and the tree is stored in memory as well. In binary search trees, there is a key and two pointers in each node, while in B-trees,  $m$  keys are paired with only  $m + 1$  pointers. This means two things. First, since a key is paired with less pointers, B-trees use less memory. Second, the tree is shallower than a binary search tree so less pointers are followed to find a key. And since the keys are stored consecutively as an array in a node, the keys are more likely to be in the cache when search inside a node. This means that B-trees are more cache friendly than binary search trees.

Compared to B-trees, B+ trees have fewer pointers in the internal nodes. If we use a fixed size memory for a node, then B+ trees can hold more keys in one node, causing the tree to be shallower. And because we only store pointers to records in the leaf node, the structure of the leaf node is almost identical to the structure of the internal node. This also makes range query easier and efficient since we only need to traverse the leaf nodes and the leaf nodes are linked together. The definition of B+ tree in this text is as follows:

```

enum BPlusTree[k: Type, v: Type, r: Region](Region[r], Ref[Node[k, v, r], r])
enum Node[k: Type, v: Type, r: Region] {
  case Inner(Ref[Int32, r], Array[k, r], Array[Node[k, v, r], r])
  case Leaf(Ref[Int32, r], Array[k, r], Array[v, r], Ref[Node[k, v, r], r])
  case Empty
}

```

Flix is a functional-first programming language that encourages the use of immutable data structures, but it has rich support for imperative programming with destructive updates to mutable data. It uses its effect system to separate pure and impure code. In particular, It uses the region concept to track the use of mutable memory. In Flix, all mutable memory belongs to a region that is tied to its lexical scope. When execution leaves the lexical scope of a region, all memory in that region becomes unreachable.

Flix has two basic types of mutable memory: References and Arrays. Higher-level mutable data structures, such as the B+ tree here, are built on top of Refs and Arrays.

A B+ tree contains a region, which tracks the lifetime of the tree, and a reference to the root node of the tree. A region is needed because other mutable data, such as the nodes of the tree, should have the same lifetime as the tree, so they need to be allocated using the same region as the tree. Therefore, the constructor will take a region as an argument, as well as the configuration of the tree, i.e. how many children can a node have at most. A new tree can then be created as follows:

```

def main(): Int32 =
  region r1 {
    let t = BPlusTree.empty(r1, 32);
    BPlusTree.size(t)
  }

```

A node in the B+ tree can be either an inner node or a leaf node. The `Empty` case serves as `Null` in other languages. The structure of the inner node is similar to that of the leaf node, they both contain an array of keys, an array of child nodes or values respectively, and an integer that stores the number of keys that currently exist in the node. The leaf node contains an extra pointer to the right sibling.

## 4.2 Insertion in B+ Trees

We now give the insertion algorithm for a B+ tree in algorithm 3, which will be used in the evaluation in the next chapter and is also the base for the  $B^{\text{link}}$  tree. For convenience, the variable *keys* and *vals* are used to represent the key and value arrays in the node that is currently examined in the context.

The insertion algorithm is a recursive algorithm. It first descends to the leaf node where the key should be inserted. The descending function simply calls the search function to find the child node to follow and pushes the current node and the index of the child into a stack, and then repeatedly calls itself on the child node until it reaches a leaf node. The search function is a binary search

---

**Algorithm 3** Insertion in B+ Tree

---

```
1: function INSERT( $t, k, v$ )
2:    $root \leftarrow root(t)$ 
3:    $leaf, stack \leftarrow descendToLeaf(root, k)$ 
4:    $i \leftarrow searchNode(k, leaf)$ 
5:   if  $i = num(keys)$  or  $k \neq keys[i]$  then
6:      $insertLeaf(k, v, i, stack, leaf, root)$ 
7:   else
8:      $keys[i] \leftarrow k$ 
9:   function INSERTLEAF( $k, v, i, node, stack, root$ )
10:     $k', v' \leftarrow insert\ k\ and\ v\ into\ keys\ and\ vals$ 
11:     $\triangleright k'$  and  $v'$  are the overflowed key and value if the array is full
12:    if  $k'$  and  $v'$  are null then
13:      increase the number of keys in the leaf by 1
14:    else
15:       $newNode, upKey \leftarrow splitLeaf(node, k', v')$ 
16:      put  $k'$  and  $v'$  into  $newNode$ 
17:      if  $stack$  is empty then
18:         $newRoot \leftarrow create\ a\ new\ inner\ node$ 
19:        put  $upKey$ ,  $leaf$ , and  $newNode$  in  $newRoot$ 
20:         $root \leftarrow newRoot$ 
21:      else
22:         $parent, i' \leftarrow pop(stack)$ 
23:         $insertInner(upKey, newNode, i', parent, stack, root)$ 
24:   function INSERTINNER( $k, n, i, node, stack, root$ )
25:    same as  $insertLeaf$ , with the function of  $splitLeaf$  replaced by  $splitInner$ 
```

---

function that returns the index of the first key that is greater than or equal to the key to be inserted, as we have discussed above.

The insertion function then checks if the key already exists in the leaf node, i.e. if the search reaches does not exceed the number of keys in the leaf node and the key is equal to the key at the index. If it exists, the value is updated. Otherwise, the key and value are inserted into the leaf node. The insertion may then cause the node to be split if the node is full. The split function will split the node into two nodes and return the key that should be inserted into the parent node. Recursive insertions are then performed using the stack. If we propagate all the way up until the stack is empty, then a new root node is created. There is one difference between splitting a leaf node and an inner node. As discussed, because values are only stored in leaf nodes in B+ trees, all keys appear twice in the tree. When splitting the leaf node, the middle key goes both into the new node and into the parent, while when splitting the inner node, the middle key leaves the current level and goes into the

parent. This satisfy the sorted property ( $K_{i-1} < v \leq K_i$ ) of the B+ tree.

### 4.3 B<sup>link</sup> Trees

There are many variants of B-trees that are designed to support concurrent reading and writing. In this project, we mainly studied the B<sup>link</sup> tree proposed by Lehman and Yao [3]. We pick this work because it appears implementable in Flix and it suits our need well. It focuses on concurrent reading and insertion while deletion is poorly supported – and we do not need deletion in our case. Although this work is old, it is a classic work in the topic and there are some programs in the real world that use B<sup>link</sup> trees as the underlying data structure, with some improvements, e.g. PostgreSQL.

The idea of the B<sup>link</sup> is to add another key-pointer pair to each inner node and an additional key to each leaf node. The structure of a node in a B<sup>link</sup> tree is shown in figure 4.2. The additional key is called the *high key*, i.e., the largest key in the subtree rooted at that node. For an inner node, the high key is the highest key in the subtree pointed by  $P_m$ . For a leaf node, the high key is the key is the same as  $K_m$ . The additional pointer is called *link* and it points to the right sibling. The node on every level is now linked together, hence the name B<sup>link</sup> tree. Note this change still satisfy the sorted property of a node, that is, all keys in the subtree pointed by  $P_m$  are greater than  $K_m$  and less than or equal to the high key  $K_{m+1}$ .

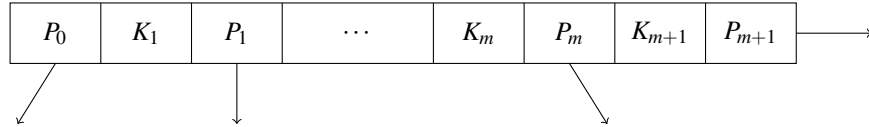


Figure 4.2: A B<sup>link</sup> tree node

The B<sup>link</sup> tree is essentially a fine-grained locking scheme. Each node is paired with a lock. When a thread is accessing the fields in a node, it has to acquire the lock first. The high key and the link then serve as a way to know if the current node is being split by another thread. When accessing a node, if the search key exceeds the high key, it means that the current node has been split and the thread should follow the link to the right sibling.

Figure 4.3 from Lehman and Yao shows an example of a B<sup>link</sup> tree. Our specification differs from Lehman and Yao in that the rightmost node at each level has  $+\infty$  as its high value, i.e., all the 99's in the diagram are changed to  $+\infty$ . Although the key  $+\infty$  never appears in the tree, this decision is consistent as long as in the beginning when the empty root is created,  $+\infty$  is set as the high value. We will see later that this will make the implementation easier in section 4.5.

The type definition of the B<sup>link</sup> tree in Flix is as follows:

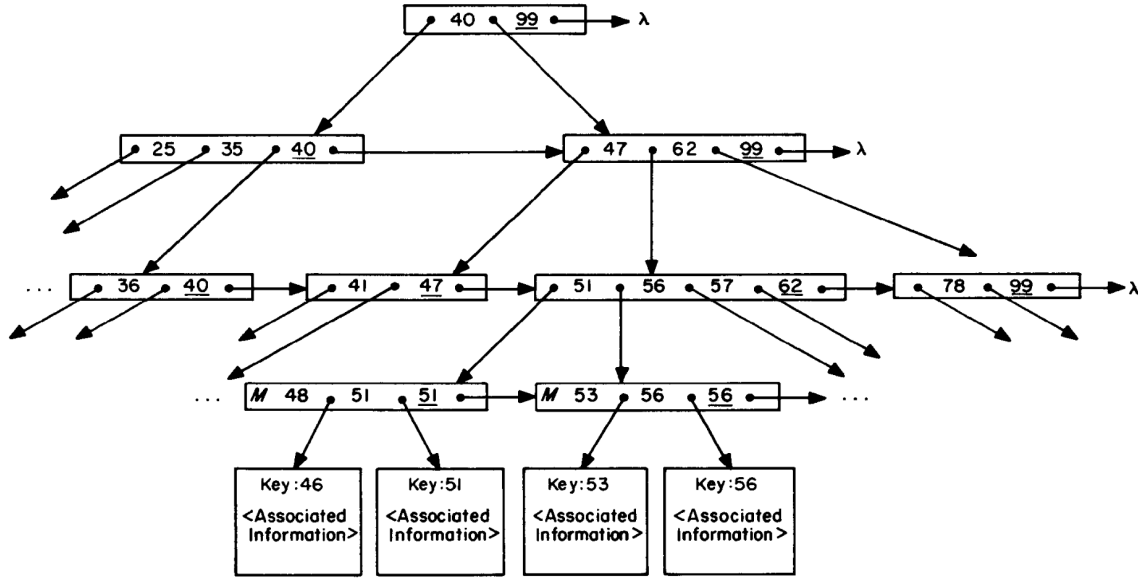


Figure 4.3: A  $B^{\text{link}}$  tree

```

enum BLinkTree[k: Type, v: Type, r: Region](Region[r], Ref[Node[k, v, r], r])
enum Node[k: Type, v: Type, r: Region] {
  case Inner(Int32, Ref[Int32, r], Array[k, r], Array[Node[k, v, r], r],
    Ref[Node[k, v, r], r], ReentrantReadWriteLock[r])
  case Leaf(Int32, Ref[Int32, r], Array[k, r], Array[v, r],
    Ref[Node[k, v, r], r], ReentrantReadWriteLock[r])
  case Empty
}

```

The node structure is similar to that of the B+ tree, with the addition of the high key, stored in the last element of the key array, the link, and the lock. The original paper uses a simple mutex lock because in their setting, a node is stored as a block in the disk. Reading a block is an atomic operation so there is no locking involved in the get method. It is only when the intention of reading the block is to update it that the lock is acquired. In our setting, we need to read-write locks because the nodes are stored in memory and when reading the fields in a node, we need to make sure that these fields are not being modified at the same time by another thread. The extra integer represents the level of the node in the tree. Leaf nodes are at level 0, and the level increases as we go up the tree. Once a node is created, the level is fixed and will not change. Level is needed here to deal with the case that Lehman and Yao did not discuss, that is, when the root node is concurrently split by other threads. We will discuss this in section 4.5.

## 4.4 Search in B<sup>link</sup> Trees

To search for a value  $k$  in the tree, the search process begins at the root and proceeds by comparing  $k$  with the keys in each node in a path down the tree. At each node, we first acquire the read lock and then examine the keys and high key. If  $k$  is greater than the high key, we follow the link to the right sibling, unlock the current node, and repeat searching in the sibling. Otherwise, we examine the key array to find the index of the first key that is greater than or equal to  $k$ . Then we use the index to get the child node to follow. We unlock the current node, push it the stack, and repeat the search in the child node until we reach a leaf node.

The details of the search function is shown in algorithm 4, following the same naming convention in 3. Note how the high key is used to determine if we need to follow the link to the right sibling. In the descending function, even if we have found the correct child to follow at some level  $i + 1$ , when we examine that child node at level  $i$ , it is still possible that it has been split by another thread, and the key we are looking for is moved to the right.

Locking is involved in the function `searchInner` and `searchLeaf`. The difference is that in `searchInner`, when we have finished searching the current level and have obtained the correct child node to follow, we release the lock as we will then proceed to the next level. In `searchLeaf`, when the function returns, the lock is still held, as the node needs to be examined further depending on if we are only reading or doing insertion.

## 4.5 Insertion in B<sup>link</sup> Trees

To insert a key-value pair, we first perform operations similar to that for searching. Beginning at the root, we scan down the tree to the leaf node that should contain the key, keeping track of the rightmost node that we examined at each level in a stack.

The insertion of the key-value pair into the leaf node may necessitate splitting the node. In this case, we split the node as what we have seen in the B+ tree, with the addition of changing the high key. The high key of the new node is set to the high key of the old node, and the high key of the old node is set to the last key that remains in the old node after splitting, which is then pushed up to the parent node recorded in the stack. The parent node, too, may need to be split. If so, we backtrack up the tree, splitting nodes and inserting new pointers into their parents. In all cases, we lock a node before modifying it. Just as what we discussed in the insertion algorithm of the B+ tree, the splitting of the inner node and the leaf node slightly differs. When splitting the leaf node, the high key is pushed up to the parent, while when splitting the inner node, it is the middle key which neither remains in the old node nor be put in the new node that is pushed up to the parent.

The details of the insertion algorithm is shown in algorithm 5, again following the same naming convention in 3, where *keys*, *vals*, and *level* are used to represent the key array, value array, and the level of the node that is currently examined in the context, i.e. *node'*.

We mentioned before that unlike the original paper, for the rightmost node at each level, the high key is set to  $+\infty$  instead of the actual largest key in this subtree. This makes splitting the rightmost



leaf node easier. When the rightmost leaf is split, we can always safely set the high key of the new node to that of the old node, which is  $+\infty$ , regardless of if we have inserted a key that is greater than all existing keys. This makes splitting the rightmost node consistent with splitting other nodes.

We also mentioned that Lehman and Yao fail to discuss what must happen when the root node becomes full and must be split. When an inserter recurses up the tree, it is possible it has recursed up to the root but the root level has been split by other threads and a new root has been created. In this case the stack it made while descending does not help for finding the correct parents anymore. We adopt the solution from PostgreSQL. PostgreSQL has adopted  $B^{\text{link}}$  trees as base for its index structure. Their solution is that when this happens, we re-descending the tree until we reach the level one above the old root. Typically the re-descending only needs to examine one level of nodes, because in most cases, if the old root level has been split, there will be enough space for insertion and re-descending is not needed. It is rare that the root level is split and is quickly filled up again by other threads. In this case, the re-descending is needed. But it is highly unlikely that even the new root level is split and filled up.

This locking scheme is efficient in that only a constant number of nodes are locked at any time. The original paper couples locks as part of moving right when relocating a child node's downlink during an ascent of the tree. This is the only point where it has to simultaneously hold three locks (a lock on the child, the original parent, and the original parent's right sibling). However, in our implementation, an inserter never couples any locks, neither between the child and the parent, nor between siblings. This means that only one lock is held at any time. We will not give a rigorous proof of the correctness here but we argue that as long as we always obtain the correct pointer to follow from the current node when the current node is locked, and we always lock the node the pointer points to before accessing its fields, the implementation is correct.

## 4.6 Range Query

The implementation of range query in B+ trees and  $B^{\text{link}}$  trees is similar. We find the first key that is greater than or equal to the left bound of the range in the inner node, with moving to the right involved in  $B^{\text{link}}$  tree, and then traverse down the tree to the leaf node. This is exactly the same as in searches and insertions. The `forAll` function simply follow the leftmost child in each node and traverse the leaf level all the way to the rightmost leaf node.

In  $B^{\text{link}}$  trees, we always lock the read lock when we are reading the fields in both inner nodes and leaf nodes. This enables threads doing queries or iterations to not block other threads from reading or writing the tree. Dead-locks could occur if in the query or iteration tries to modify the tree. Programmers using these two functions should be aware of this and avoid modifying the tree in the query or iteration.

---

**Algorithm 4** Search in  $B^{\text{link}}$  Tree

---

```
1: function GET( $k, t$ )
2:    $root \leftarrow \text{root}(t)$ 
3:    $leaf, stack \leftarrow \text{descendToLeaf}(k, root, [])$ 
4:    $leaf', i \leftarrow \text{searchLeaf}(k, leaf, \text{false})$ 
5:   if  $i = \text{num}(keys)$  or  $k \neq keys[i]$  then
6:     release read lock of  $leaf'$ 
7:     return None
8:   else
9:      $v \leftarrow vals[i]$ 
10:    release read lock of  $leaf'$ 
11:    return Some( $v$ )
12: function DESCENDTOLEAF( $k, node, stack$ )
13:   if  $node$  is leaf then
14:     return  $node, stack$ 
15:   else
16:      $child, stack' \leftarrow \text{searchInner}(k, node, stack)$ 
17:     return  $\text{descendToLeaf}(k, child, stack')$ 
18: function SEARCHINNER( $k, node, stack$ )
19:   acquire read lock of  $node$ 
20:   if  $\text{needToGoRight}(k, node)$  then
21:      $sibling \leftarrow \text{link}(node)$ 
22:     release read lock of  $node$ 
23:     return  $\text{searchInner}(k, sibling, stack)$ 
24:   else
25:      $i \leftarrow \text{findFirstGreaterOrEqual}(k, keys)$ 
26:      $child \leftarrow vals[i]$ 
27:     release read lock of  $node$ 
28:     return  $child, node :: stack$ 
29: function SEARCHLEAF( $k, node, isWriting$ )
30:   acquire read or write lock depending on  $isWriting$ 
31:   if  $\text{needToGoRight}(k, node)$  then
32:      $sibling \leftarrow \text{link}(node)$ 
33:     release read or write lock depending on  $isWriting$ 
34:     return  $\text{searchInner}(k, sibling, isWriting)$ 
35:   else
36:      $i \leftarrow \text{findFirstGreaterOrEqual}(k, keys)$ 
37:     return  $node, i$ 
38: function NEEDTOGORIGHT( $k, node$ )
39:   return  $\text{link}(node) \neq \text{null} \wedge k > \text{high key}(node)$ 
```

---

---

**Algorithm 5** Insert in  $B^{\text{link}}$  Tree

---

```
1: function PUT( $k, v, t$ )
2:    $root \leftarrow root(t)$ 
3:    $leaf, stack \leftarrow descendToLeaf(k, root, [])$ 
4:    $insertLeaf(k, v, leaf, stack, root)$ 
5: function INSERTLEAF( $k, v, node, stack, root$ )
6:    $node', i \leftarrow searchLeaf(k, node', true)$ 
7:   if  $i < num(keys)$  and  $k = keys[i]$  then
8:      $vals[i] \leftarrow v$ 
9:     release write lock of  $node'$ 
10:  else
11:     $k', v' \leftarrow$  insert  $k$  and  $v$  into the key and value array
12:    if  $k'$  and  $v'$  are null then
13:      increase the number of keys in the  $node'$  by 1
14:      release write lock of  $node'$ 
15:    else
16:       $newNode, upKey \leftarrow splitLeaf(node', k', v')$ 
17:      release write lock of  $node'$ 
18:      if  $stack$  is empty then
19:        if  $level(root) = level$  then
20:           $newRoot \leftarrow$  create a new inner node
21:          put  $upKey, node'$ , and  $newNode$  in  $newRoot$ 
22:           $root \leftarrow newRoot$ 
23:        else
24:           $redescend(upKey, newNode, level, root)$ 
25:      else
26:         $parent \leftarrow pop(stack)$ 
27:         $insertInner(upKey, newNode, parent, stack, root)$ 
28: function INSERTINNER( $k, n, node, stack, root$ )
29:   same as  $insertLeaf$ , with the function of  $splitLeaf$  replaced by  $splitInner$ 
30: function REDESCEND( $k, n, toLevel, root$ )
31:    $node, stack \leftarrow descendToLevel(k, toLevel, root, [])$ 
32:    $insertInner(k, n, node, stack, root)$ 
33: function DESCENDTOLEVEL( $k, toLevel, node, stack$ )
34:   if  $level(node) = toLevel + 1$  then
35:     return  $node, stack$ 
36:   else
37:      $child, stack' \leftarrow searchInner(k, node, stack)$ 
38:     return  $descendToLevel(k, toLevel, child, stack')$ 
```

---

## Chapter 5

# Evaluation

In this chapter, we answer the following five research questions:

**RQ0** Does our implementation of the trees satisfy the asymptotic bound?

**RQ1** Is insertion and range query in the B+ tree faster than in the built-in red-black trees?

**RQ2** Is insertion in the  $B^{\text{link}}$  tree faster than in the B+ tree and red-black tree?

**RQ3** Is Datalog evaluation using B+ trees faster than using red-black trees?

**RQ4** Is Datalog evaluation using  $B^{\text{link}}$  trees faster than using B+ trees and red-black trees?

The experiments are run on a machine with 64 GB of memory and an Intel Core i5-13500 that has 14 threads.

To answer the question 0 and 1, we do the following experiment. We create trees where the keys and values are integers. The value does not matter in the experiment so we set the value to be the same as the key. We then first propagate the tree with a certain number of keys that are generated randomly. We then measure the time of inserting another key-value pair into the tree. Because the time complexity of insertion in both B+ trees and red-black trees are  $O(\log_2 n)$  when binary search is used in B+ trees, if we divide the time by  $\log_2 n$ , we should expect a constant. Since measuring the time of inserting one key-value pair is not accurate, we insert 100 random key-value pairs and divide the time by 100. Because we propagate the tree to a size that is much greater than 100, dividing  $100 \log_2 n$  is acceptable even though  $n$  is changing.

If not mentioned otherwise, the random number is generated under a fixed seed so that the same set of keys are generated for different trees. And we run the experiment 5 times and take the average time for the benchmark on trees.

The result of insertion is shown in figure 5.1 where the size of the trees range from  $2^{17}, 2^{18}, \dots, 2^{21}$ . We can see that both B+ trees and red-black trees satisfy the bound since all lines are almost flat. We can clearly see that insertion in B+ trees is faster than in red-black trees because in B+ trees, when

comparing keys in a node, it is more likely that the key array is in the cache. The B+ tree with order 16 is obviously slower and there is no significant difference between the B+ tree with order 32, 64 and 128. We mention that the built-in red-black trees in the flx standard library is an optimized and benchmarked implementation while our B+ trees is a simple and quick implementation.

B+ trees are often used for database systems where the nodes are stored in a block in a disk so the size of the node is usually determined so that one node fits in one block, minimizing the number of I/O operations. In our case, the nodes are stored in memory and it is fine that a node does not fit in a cache line. Smaller nodes increase the tree height and traversal time but fit better in CPU caches, leading to potentially faster access times within each node. We expect that the B+ tree with a too small size of node will be slower since the L1 cache is usually large to hold a node so the height of the tree is more important. The result of insertion meets our expectation as the B+ tree with order 16 is slower than the other three.

The result of range query is shown in figure 5.2. We first propagate the tree with  $2^{20}$  keys starting from 1 to  $2^{20}$  but the order is randomly shuffled. We then measure the time of querying different ranges where the left bound is half of the size of tree, i.e.  $2^{19}$ , and the size of the range is  $2^6$ ,  $2^{10}$ ,  $2^{14}$ ,  $2^{18}$ . We expect that range query on a small range in B+ trees is faster than red-black tree which is true in the experiment. However when the range becomes bigger, e.g.  $1/4$  of the size of the tree, namely  $2^{18}$ , the B+ tree is slower than red-black tree. This is because, as we have seen in section 4.6, the implementation of range query tries to find the first key that is greater than or equal to the left bound in each node using binary search. When the range is large, we need to do plenty of comparisons in each node to find the child to follow. But the values are only stored in leafs, so we have to traverse down to the leaf before we actually do any work. However in red-black trees, when traversing the inner nodes, if the key falls in the range, we can immediately run the given function on the value in the node.

To answer the question 2, we propagate the trees to a certain size with random integers and measure the time. For  $B^{\text{link}}$  trees, we use 14 threads which is the number of cores in the machine and the input is partitioned evenly among the threads. The result in 5.3 shows that the best  $B^{\text{link}}$  tree is on average 3 times faster than the best B+ tree which is again 3 times faster than the red-black tree.  $B^{\text{link}}$  with order 32 is the best in the experiment and 64 also comes very close. If the nodes are too small, the height of tree is too large and nodes are split more often, leading to more locks and more time. If the nodes are too large, the time it takes to examine a node and do insertion on a node becomes longer and locks are more likely to compete with each other. We have also tested different combinations of the number of threads and the order of the tree and the result is consistent – using threads equal to the number of cores and the order of 32 and 64 are the best.

We did not test the range query of the  $B^{\text{link}}$  tree because it is implemented in the same way as the B+ tree with locks acquired if other threads are modifying the node and there is no parallelism available for a range query.

To answer the question 3 and 4, we run the benchmark with four different implementations of the engine. The baseline is the built-in engine in Flix release 0.46.0 that uses red-black trees. Because Flix is a functional-first language that adopts the concept of regions to separate pure and impure code as we have mentioned in section 4.1. The definition of Datalog programs, including the rules

(constraints) and the results (models) is defined as follow:

```
pub enum Datalog[v] {  
  case Datalog(Vector[Constraint[v]], Vector[Constraint[v]])  
  case Model(Map[RamSym[v], Map[Vector[v], v]])  
  case Join(Datalog[v], Datalog[v])  
}
```

If we use a mutable data structure in the engine, after the evaluation of the whole Datalog program, we need to copy the results back to into an immutable map. We can not simply change Map to BLinkTree because it will introduce a region variable *r* in the type, i.e. Datalog[v, *r*]. Therefore, for fair comparisons for the B+ tree and B<sup>link</sup> tree, we add another engine where the only difference between the built-in engine is that we add an extra copy of the results after evaluation. The results are shown in figure 5.4.

Compared to the baseline, the B+ tree engine is on average slightly slower than the baseline, partly due to the copying. If we compare it with the baseline with copying, the B+ tree engine is almost always faster, although just merely. The B<sup>link</sup> tree engine is significantly faster than the baseline and the B+ tree engine. On the four realistic programs in the Closure.flix and the first points-to analysis in the PointsTo.flix, the B<sup>link</sup> tree engine is 1.2x to 2.2x faster than the baseline, depending on the number of rules in the program and how rule looks like. Usually, we gain more from parallelism if there are more rules. For example, on the program propagate3, which we illustrated in section 3.4, the B<sup>link</sup> tree engine is 5x faster than the baseline.

The only exception where both B<sup>link</sup> tree engine and the B+ tree engine are slower than the baseline is the program points2. The difference between this points2 and points1 is that two atoms in the body of the load rule are swapped, causing a range query statement falls back to a search-and-if statement in the compiled RAM code. The reason why the B<sup>link</sup> tree engine is slower than the baseline is not clear.

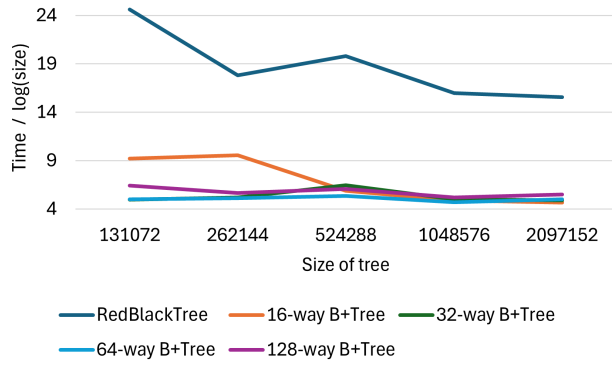


Figure 5.1: Time of inserting 1 value in Red-Black Tree and B+Trees

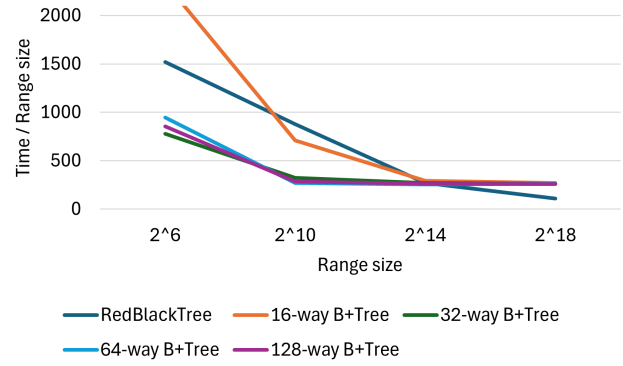


Figure 5.2: Time of range query in Red-Black Tree and B+Trees containing  $2^{20}$  keys

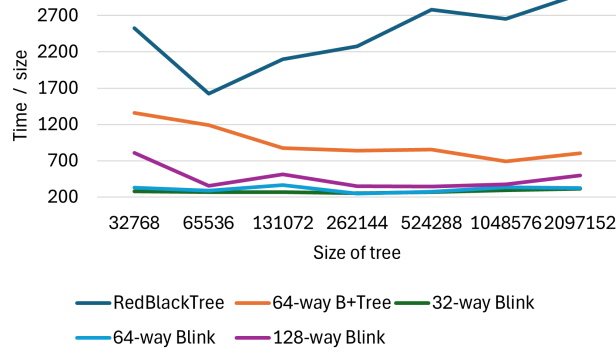


Figure 5.3: Time of propagating different trees

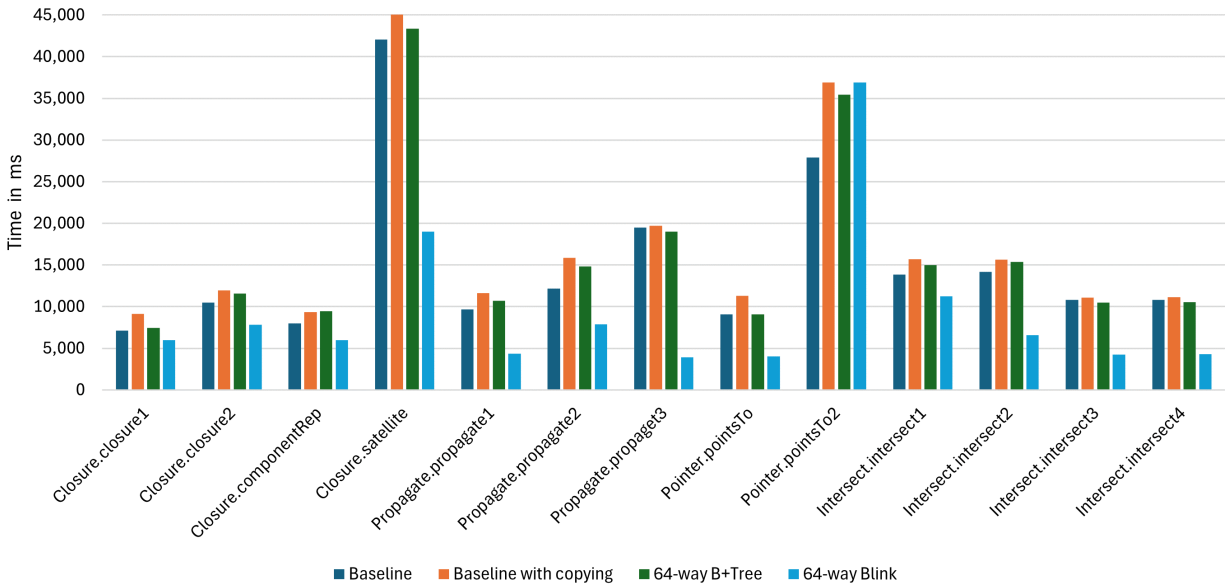


Figure 5.4: Benchmark of engines using different trees

## Chapter 6

# Future Work

### 6.1 Optimizing the $B^{\text{link}}$ Tree

The implementation of the  $B^{\text{link}}$  tree is a quick implementation. There are possibly many optimizations that can be done with respect to both the algorithm and the code. For example, there are different ways to implement the range query. The current implementation is to find the first key that is greater than or equal to the left bound in each level of the tree. This is not optimal because in the inner nodes, we will do plenty of comparisons of keys that are in the range which will be revisited again in the leaf level if the range is large. Since we have high keys in  $B^{\text{link}}$  trees, we can instead find the node whose high key is the first one that is greater than or equal to the left bound in each level. Only after we have reached the correct leaf node, we will start to examine the key array.

### 6.2 Index Selection and Rule Rewriting

We can also consider optimizations at the source level. For example, the benchmark `points1` contains the rules in example 3.1. We have seen that the load rule will produce two range queries. However, programmers are probably not aware the underlying implementation of the engine and they might write a load rule where the second and the third atom in the body are swapped. If then, the second atom will be compiled to a search statement as follows:

```
search 0$ ∈ Load do
  search 1$ ∈ ΔPointsTo do
    if((0$[0], 1$[1]) ∉ PointsTo) then
      query {2$ ∈ PointsTo | 2$[0] == 0$[1] ∧ 2$[1] == 1$[0]} do
        project (0$[0], 1$[1]) into ΔPointsTo'
      end
    end
  end
end;
search 0$ ∈ Load do
```



```

search 1$ ∈ PointsTo do
  if((0$[0], 1$[1]) ∉ PointsTo) then
    query {2$ ∈ ΔPointsTo | 2$[0] == 0$[1] ∧ 2$[1] == 1$[0]} do
      project (0$[0], 1$[1]) into ΔPointsTo'
    end
  end
end
end;

```

We can add an analyzing phase in the compiler to reorder the atoms in the rules so that range queries are used as much as possible.

We can also analyze the program to determine which index is the best for each relation, that is, what order of the elements in the tuple is the best when the tuple is stored in a sorted tree data structure. For example, the benchmark `intersect3` contains rules similar to the following:

```

A(1 , i) :- Check(a, b, c), In(i, a, b, c).
A(2 , i) :- Check(a, _, c), In(i, a, _, c).
A(3 , i) :- Check(a, _, _), In(i, a, _, _).
A(4 , i) :- Check(a, b, _), In(i, a, b, _).
A(5 , i) :- Check(a, b, c), In(i, a, b, c).

```

where the EDB is `Check` and `In` and the IDB is `A`. The current implementation will compile all five rules to be search statements. But if we analyze the program, we can actually store the tuples of the `In` relation as `(a, b, c, i)`. Then these rules can be translated into range queries.

Multiple indices can be created for the same relation. This means that for one relation, we can have multiple trees that store the tuples in different orders of the elements. Although performance can substantially improve, the memory footprint can become very large.

Index selection is studied and implemented by some other Datalog engines. For example, Souffle, A Datalog Synthesis Tool for Static Analysis, uses a strategy to automatically constructs an appropriate set of indices [5].

### 6.3 Parallelism through Data Partitioning

Another direction of parallelism is to partition the tuples of the relations and let different threads do the evaluation of the rules on different partitions. This parallelism is probably more promising in that the level of parallelism is not limited by the number of rules in the program or by the skewed workload of the rules. But data partitioning is also difficult because we want such a partition that the workload of each thread is balanced and the communication between threads is minimized. Communication here means that the threads probably need to exchange their newly found facts in the IDB with other thread based on the partitioning strategy. A rich line of research on partitioning strategies for Datalog exists since the 1990s. The monograph *Modern Datalog Engines* [2] has listed a good resource of literature and Datalog engines on this topic.

## Chapter 7

# Conclusion

► Conclude on the problem statement from the introduction ◀

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