

Regular Expression Types for XML

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We propose *regular expression types* as a foundation for statically typed XML processing languages. Regular expression types, like most schema languages for XML, introduce regular expression notations such as repetition (*), alternation (|), etc., to describe XML documents. The novelty of our type system is a semantic presentation of subtyping, as inclusion between the sets of documents denoted by two types. We give several examples illustrating the usefulness of this form of subtyping in XML processing.

The decision problem for the subtype relation reduces to the inclusion problem between tree automata, which is known to be EXPTIME-complete. To avoid this high complexity in typical cases, we develop a practical algorithm that, unlike classical algorithms based on determinization of tree automata, checks the inclusion relation by a top-down traversal of the original type expressions. The main advantage of this algorithm is that it can exploit the property that type expressions being compared often share portions of their representations. Our algorithm is a variant of Aiken and Murphy's set-inclusion constraint solver, to which are added several new implementation techniques, correctness proofs, and preliminary performance measurements on some small programs in the domain of typed XML processing.

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1. INTRODUCTION

XML [Bray et al. 2000] is an emerging standard format for tree-structured data. One of the reasons for its popularity is the existence of a number of schema languages, including DTDs [Bray et al. 2000], XML-Schema [Fallside 2001], DSD [Klarlund et al. 2000], RELAX [Murata 2001], and RELAX NG [Clark and Murata 2001], that can be used to define “types” (or “schemas”) describing structural constraints on data and thereby improve the safety of data processing and exchange.

However, the use of types in mainstream XML processing technology is often limited to checking only *data*, not *programs*. Typically, an XML processing program first reads an XML document and checks that it conforms to a given type using a *validating parser*. The program then uses either a generic tree manipulation library such as DOM [DOM 2001] or a dedicated XML language such as XSLT [Clark 1999] or XML-QL [Deutsch et al. 1998]. Since these tools make no systematic connection between the program and the types of the documents it manipulates, they provide no compile-time guarantee that the documents produced by the program will always conform to an intended type.

In this article, we propose *regular expression types* as a foundation for statically typed processing of XML documents. Regular expression types capture (and generalize) the regular expression notations (*, ?, |, etc.) commonly found in schema languages for XML, and support a natural semantic notion of subtyping.

We have used regular expression types in the design of a domain-specific language called XDuce (“transduce”) for XML processing [Hosoya and Pierce 2002, 2003]. In the present article, however, our focus is on the structure of the types themselves, their role in describing transformations on XML documents, and the algorithmic problems they pose. Interested readers are invited to visit the XDuce home page

<http://xduce.sourceforge.net>

for more information on the language as a whole.

As a simple example of regular expression types, consider the definitions

```
type Addrbook = addrbook[Person*]
type Person   = person[Name,Email*,Tel?]
type Name     = name[String]
type Email    = email[String]
type Tel      = tel[String]
```

corresponding to the following set of DTD declarations:

```
<!ELEMENT addrbook person*>
<!ELEMENT person   (name,email*,tel?)>
<!ELEMENT name     #PCDATA>
<!ELEMENT email    #PCDATA>
<!ELEMENT tel      #PCDATA>
```

Type constructors of the form `label[...]` classify tree nodes with the tag `label` (i.e., XML structures of the form `<label>...</label>`). Thus, the inhabitants of the types `Name`, `Email`, and `Tel` are all strings with an appropriate identifying label. Types may also involve the regular expression operators `*` (repetition) and `?` (optional occurrence), as well as `|` (alternation). Thus, the type `Addrbook` describes a label `addrbook` whose content is zero or more repetitions of subtrees of type `Person`. Likewise, the type `Person` describes a label `person` whose content is a `Name` subtree, zero or more `Email` subtrees, and an optional `Tel` subtree. An instance of the type `Addrbook` is the following XML document:

```
<addrbook>
  <person> <name> Haruo Hosoya </name>
           <email> hahosoya@upenn </email>
           <email> haruo@u-tokyo </email> </person>
  <person> <name> Jerome Vouillon </name>
           <email> vouillon@upenn </email>
           <tel> 123-456-789 </tel> </person>
</addrbook>
```

We define subtyping between regular expression types in a semantic fashion. A type, in general, denotes a set of documents; subtyping is simply inclusion between the sets denoted by two types. For instance, consider again the `Person` type definition from above

```
type Person = person[Name,Email*,Tel?]
```

and the following variant:

```
type Person2 = person[(Name | Email | Tel)*]
```

Elements of the `Person` type can have one name, zero or more emails, and zero or one tels in this order, while the `Person2` type allows any number of such nodes in any order. Therefore, `Person2` describes strictly more documents, which implies that `Person` is a subtype of `Person2`. Such subtype inclusions can be quite useful in programming. For example, suppose that we originally have a value of type `Person`. The above inclusion allows us to process this value using code that does not care about the ordering among the name, email, and tel nodes. (Such a situation might arise, for example, if we want to display the child nodes in a linear format, where we would naturally write a single loop over the sequence of child nodes with a case branch for each tag.)

Note that if we replaced `Person` with a more conventional type

```
person(name(string) × email(string) list × tel(string) option)
```

(using sum, product, and ML-like list and option types, plus unary covariant constructors `person`, `name`, `email`, and `tel`) and `Person2`, analogously, with

```
person((name(string) + email(string) + tel(string)) list)
```

the conventional subtyping of sum and product types would *not* yield the inclusion above. In general, the subtype relation obtained from our definition is

quite a bit more permissive than conventional subtyping. We give some further examples in Section 2.

Regular expression types exactly correspond to *tree automata* [Comon et al. 1999]—a finite-state machine model for accepting trees. It is easy to construct, from a given type, a tree automaton that accepts just the set of trees denoted by the type (Appendix A). Therefore, the subtyping problem can be reduced to the inclusion problem between tree automata, which is known to be decidable. Unfortunately, in the worst case, this problem is EXPTIME-complete [Seidl 1990].

To address this high complexity, we develop an algorithm that runs efficiently for the cases that we expect to encounter most often in programming with regular expression types. Instead of beginning from classical algorithms for testing tree automata inclusion [Comon et al. 1999], which work by (expensively) transforming their input automata to other completely separate automata before testing their properties, we take Aiken and Murphy’s algorithm [Aiken and Murphy 1991] as the starting point. Like standard subtyping algorithms for other type systems, this algorithm works top-down—that is, a given pair of types, we check matching of the top-most type constructors, proceed to the subcomponents of the types, and repeat the same check recursively. The main advantage of this top-down algorithm is that it enables many simple optimizations. In particular, we can exploit reflexivity ($T \leq T$) in order to decide subtype relations by looking at only a part of the whole input type expressions.

Our additions to the work of Aiken and Murphy are fourfold. First, we give proofs of soundness, completeness, and termination for (a reformulation of) their algorithm. Second, we give a detailed, intuitive explanation of the key step in the algorithm involving “untagged” union types, which was not presented well in their original paper. Third, we incorporate several useful implementation techniques, including (1) sharing type expressions as much as possible by using a representation that exploits equivalence between types, and (2) caching intermediate results of the subtyping algorithm by using a *functional* data structure (for which the operations do not destructively update the data). We describe these techniques in detail in Section 5. Fourth, we describe some preliminary experiments with our algorithm (as implemented in the prototype XDuce interpreter), confirming its practicality on several small examples. (Aiken and Murphy’s algorithm also addresses a more general problem: set-constraint solving. See Section 7 for further discussion of the relationship between their algorithm and ours.)

The contributions of this article can be summarized as follows:

- We motivate the use of regular expression types and set-inclusion-based subtyping for the domain of XML processing.
- We formalize the connection of regular expression types to tree automata.
- We develop a practical subtyping algorithm, present it in detail (along with an intuitive explanation of its key ideas), and give soundness, completeness, and termination proofs.
- We describe several implementation techniques for the subtyping algorithm and present preliminary measurements of their practical effects.

The article is organized as follows: In the next section, we give some examples of programming with regular expression types. In Section 3, we describe the connection of regular expression types to tree automata and give a precise definition of subtyping. In Section 4, we present our subtyping algorithm and prove its correctness. Section 5 describes our implementation techniques, and Section 6 presents the results of our preliminary experiments. We survey related work in Section 7 and conclude in Section 8. Appendix A presents an algorithm for translating regular expression types to tree automata, with its proof of correctness.

2. REGULAR EXPRESSION TYPES

We begin with a series of examples illustrating the application of regular expression types and subtyping to XML processing.

2.1 Values

Each type in our language denotes a set of sequences. Types like `String` and `tel[String]` denote singleton sequences; the type `Tel*` denotes sequences formed by repeating the singleton sequence `Tel` any finite number of times. So each element of the type `person[Tel*]` is a singleton sequence labeled with `person`, containing an arbitrary-length sequence of `Tels`. If `S` and `T` are types, then the type `S, T` denotes all the sequences formed by concatenating a sequence from `S` and a sequence from `T`. The comma operator is associative: the types `(Name, Tel*), Addr` and `Name, (Tel*, Addr)` have exactly the same elements. As the unit element for the comma operator, we have the *empty* sequence type, written `()`. Thus, `Name, ()` and `() , Name` are equivalent to `Name`.

2.2 Subtyping

The *subtype* relation between two types is simply inclusion between the sets of sequences that they denote. (See Section 3.1 for the formal definition.)

To illustrate the subtype relation, let us show the sequence of steps involved in verifying that the XML document given in the introduction actually has type `Addrbook`. First, from the intuition that `?` means “optional,” we expect the following inclusions:

$$\begin{aligned} \text{Name, Addr} &<: \text{Name, Addr, Tel?} \\ \text{Name, Addr, Tel} &<: \text{Name, Addr, Tel?} \end{aligned}$$

Notice that each right-hand side describes a larger set of sequences than the left-hand side. Similarly, `*` means “zero or more,” so in particular it can be three:

$$T, T, T <: T^*$$

Wrapping both sides of the first two inclusions with the label `person` and combining these with the third, we obtain:

$$\begin{aligned} &\text{person[Name, Addr]}, \\ &\text{person[Name, Addr, Tel]}, \\ &\text{person[Name, Addr]} \\ &<: (\text{person[Name, Addr, Tel?]})^* \end{aligned}$$

Finally, enclosing both sides with the `addrbook` constructor, we obtain

```
addrbook[
  person[Name,Addr] ,
  person[Name,Addr,Tel] ,
  person[Name,Addr]]
<: addrbook[(person[Name,Addr,Tel?])*]
= Addrbook
```

Since the XML document given in the introduction trivially has the type on the left hand side, it has also the type on the right hand side.

2.3 Recursion

As in many type systems, we support recursive types for describing arbitrarily nested structures. Consider the following definitions.

```
type Fld = Rcd*
type Rcd = name[String], folder[Fld]
         | name[String], url[String], (good[] | broken[])
```

The mutually recursive types `Fld` (“folder”) and `Rcd` (“record”) define a simple template for storing structured lists of bookmarks, such as might be found in a web browser: a folder is a list of records, while a record is either a named folder or a named URL plus either a good or a broken tag indicating whether or not the link is broken.

We can write another pair of types

```
type GoodFld = GoodRcd*
type GoodRcd = name[String], folder[GoodFld]
              | name[String], url[String], good[]
```

which are identical to `Fld` and `Rcd` except that links are all good. Intuitively, we expect that `GoodFld` should be a subtype of `Fld` because `GoodFld` allows fewer possibilities than `Fld`. Our type system validates this inclusion.

2.4 Regular Expression Types as Derived Forms

We have seen a rich variety of type constructors, but some of them can actually be derived as combinations of a smaller set of other constructors—concatenation, labeling, alternation, empty sequence, and recursive definition. For example, the optional type `T?` can be rewritten as `T | ()`, using an alternation type (a.k.a. *union type*) and the empty sequence.

Other regular expression operators are also definable. `T+`, standing for one or more repetitions of `T`, can be rewritten as `(T, T*)`. Furthermore, `T*` itself can also be derived using recursion. That is, `T*` is equal to a type name `X` defined by the following equation:

```
type X = T, X | ()
```

Note the similarity to the definition of `list` as a datatype in ML.

2.5 Subtagging

In XML processing, we sometimes encounter situations where we have to handle a large number of labels and it is convenient to organize them in a hierarchy, in the style of object-oriented languages. This leads us to support a notion of “subtagging” in our type system, allowing subtyping between types with different labels. This feature goes beyond the expressive power of DTDs, but a similar mechanism called “substitution groups” can be found in XML-Schema [Fallside 2001, Sect. 4.6].

The subtagging relation is a reflexive and transitive relation on labels. We declare subtagging explicitly with a set of global subtag forms. For example, the following declares that the tags *i* (italic) and **b** (bold) are subtags of `fontstyle`:

```
subtag i  <: fontstyle
subtag b  <: fontstyle
```

In the presence of these declarations, we have the subtype relations

```
i[T] <: fontstyle[T]
b[T] <: fontstyle[T]
```

for all T . These relations allow us to collapse two case branches for *i* and *b* into one for `fontstyle`, when both branches behave the same. This use of subtagging is similar to the common technique in object-oriented programming of defining an abstract class `fontstyle` with subclasses *i* and *b*.

Subtagging is also useful for other purposes. Our type system provides the special label `~`, denoting “any label.” That is, for every label *l*, we have the following built-in subtagging relation:

```
subtag l <: ~
```

Thus, the type `~[T]` describes *any* labeled structure whose contents belong to type T . The label `~`, in turn, can be used to define a completely generic type `Any` as follows (assuming the base types in our language are `String`, `Int`, and `Float`):

```
type Any = (~[Any] | String | Int | Float)*
```

That is, the values described by `Any` consist of zero or more repetitions of arbitrary labels (containing `Any`) and base types.

2.6 Pattern Matching

Regular expression types can enhance the pattern matching mechanisms found in mainstream functional languages. For example, in XDuce, we can write the following pattern match:

```
person[Name,Addr]*, (val x as person[Name,Addr,Tel]), Person*
-> (* do some stuff *)
| val y as person[Name,Addr]*
-> (* do other stuff *)
```

A pattern of the form `(val x as T)` matches any value of type `T` and binds the variable `x` to this value. In this example, the first case matches values containing at least one person with `tel`. In this case, the variable `x` is bound to the first person with a `tel`. The second case matches values containing no person with `tel`.

Notice that the first pattern—`person[Name,Addr]*`—in the first case matches a *variable length* sequence, something that is beyond the power of ML pattern matching. This makes standard techniques for exhaustiveness checking somewhat difficult to apply to such patterns. Fortunately, we can use subtyping for checking exhaustiveness of pattern matches. We assume that the type for the input value to the pattern match is available from the context. (In XDuce, this is ensured by type annotations on function headers.) In the example, suppose that the type of the input value is `Person*`. In order to show exhaustiveness of the pattern match with respect to this type, it is sufficient to show that every value of type `Person*` is accepted by the pattern. This leads to the following subtyping check

```
Person* <: (person[Name,Addr]*,
           person[Name,Addr,Tel],
           Person*)
          | person[Name,Addr]*
```

where the right-hand side is calculated by taking the union of the types of the two pattern clauses. A more thorough study on regular expression pattern matching and various static checking methods for it (including redundancy checks, type inference for pattern variables, and ambiguity checks) can be found in our separate papers [Hosoya and Pierce 2002; Hosoya 2003].

2.7 Semistructured Data

One of the main application domains for XML is storing and transmitting databases. Since this flexible representation allows smooth evolution of database schemas and integration of databases with different schemas, it is often called a *semistructured* format in the database community. This view is especially useful for wrapper/mediator systems for the Web that integrate multiple independent data sources, which may themselves occasionally evolve. In this section, we present some different scenarios of database evolution and integration and show how our regular expression types ensure static safety in a flexible and robust way.

Suppose that we begin with the following XML database *A*

```
<addrbook>
  <person>
    <name>Haruo Hosoya</name>
    <addr>Tokyo</addr>
  </person>
  <person>
    <name>Jerome Vouillon</name>
    <addr>Paris</addr>
```



```

    <tel>123-456-789</tel>
  </person>
</addrbook>

```

belonging to the type Addrbook, defined as follows:

```

type Addrbook = addrbook[Person*]
type Person = person[Name,Addr,Tel?]

```

Now, suppose we upgrade this database so that some person records can contain arbitrarily many tels. This process involves changes to types, databases, and programs. We change the types as follows:

```

type Person = person[Name,Addr,Tel*]

```

Notice that the new content type (Name,Addr,Tel*) of person is a supertype of the old type (Name,Addr,Tel?) and therefore the type Addrbook of the whole database becomes bigger as well. This means that our database, which had the old type, still conforms to the new type, without the need of restructuring. After adding some tel fields to our database, we arrive at the following database *B*:

```

<addrbook>
  <person>
    <name>Haruo Hosoya</name>
    <addr>Tokyo</addr>
    <tel>111-222-333</tel>
  </person>
  <person>
    <name>Jerome Vouillon</name>
    <addr>Paris</addr>
    <tel>123-456-789</tel>
    <tel>999-888-777</tel>
  </person>
</addrbook>

```

At each step in this process, the type of the database is Addrbook. The database can therefore smoothly evolve while preserving the robustness provided by type safety.

Upgrading the programs that operate on our database can be slightly trickier. Since the new type is a *supertype* of the old type rather than a subtype, all functions that *output* the old type can be treated as yielding the new type instead. On the other hand, functions that *input* the old type have to be modified so as to handle the additional cases. Some programmers may be happy with this, since the type system helps in isolating the part of the program requiring updates. Other programmers may feel that the types are preventing “forward compatibility” of old programs. For example, if we are interested in extracting specifically the Name field, then our program should work for the new database just as well as the old. But this sort of forward compatibility can easily be achieved, at the cost of writing the original program in a slightly more refined way: we maintain the convention that functions on persons should actually

be able to handle inputs of type $(\text{Name}, \text{Addr}, \text{Tel}?, \text{Any})$, simply ignoring the additional fields at the end. Now if, when the database's type is evolved, new fields are always added at the end, these old programs will work and typecheck without change.

XML also makes database integration easier than more structured formats such as relational databases. Again, regular expression types help ensure the type safety of integration steps. For example, consider integrating the previous database B with another database C with a slightly different type from B 's:

```
type Addrbook2 = addrbook[Person2*]
type Person2 = person[Name,Addr,Email*]
type Email = email[String]
```

Data integration again involves changes to types, databases, and programs. We integrate databases by constructing a tree whose root has the label `addrbook` and whose content is the concatenation of the contents of the two databases. The natural type of this merged database is:

```
type Addrbook = addrbook[Person*,Person2*]
```

Suppose that we want to write a program to scan the whole sequence and extract the names of all the persons. For writing such a program, the above type of the database is rather inconvenient, since it involves two occurrences of repetition, naturally leading to two separate loops for scanning the whole sequence. Obviously, it is better to roll these two loops into one. To do this, we can use a subtype inclusion that forgets the fact that all the Persons come before the Person2s:

```
Person*,Person2* <: (Person|Person2)*
```

Now we have *one* repetition, each of whose elements has either type `Person` or else type `Person2`, leading naturally to a one-loop scan. However, we can do better: each step of the natural scan over this type involves two very similar cases, both of which just extract the `Name` field. We can use one more subtype inclusion (actually an equivalence, hence written $=$, using the fact that alternation distributes over labels and concatenations) to rewrite the type so that the common structure is exposed:

```
person[Name,Addr,Tel*] | person[Name,Addr,Email*]
= person[(Name,Addr,Tel*) | (Name,Addr,Email*)]
= person[Name,Addr,(Tel*|Email*)]
```

After all this rewriting, the type `AddrBook` is now expressed in a form that leads naturally to scanning the Name fields with a single compact loop:

```
addrbook[ person[Name,Addr,(Tel*|Email*)]* ]
```

These distributive laws illustrate the flexibility of regular expression types. Note that the validity of these laws depends on the fact that our alternation operation is interpreted as an *untagged*, set-theoretic union—in contrast to the “tagged” sum types found in languages such as ML and Haskell.

3. DEFINITIONS

It will be useful to distinguish two forms of types: *external* and *internal*. The external form is one that the user actually reads and writes; all the examples in the previous sections are in this form. Internally, however, our subtyping algorithm uses the simpler internal representation to streamline both the implementation and its accompanying correctness proofs.

In this section, we define the syntax and semantics of each form as well as the subtype relations for each. For brevity, we omit base values (strings, numbers, etc.) and their types from our formalization; they are easily added.

3.1 External Form

We assume a countably infinite set of labels, ranged over by l . Values are defined as follows:

$$v ::= l_1[v], \dots, l_n[v] \quad (n \geq 0).$$

We write $()$ for the empty sequence and v, w for the concatenation of sequences v and w .

We assume a countably infinite set of type names, ranged over by X . Type expressions are then defined as follows:

$T ::= ()$	empty sequence
X	type variable
$l[T]$	label
T, T	concatenation
$T T$	union
\emptyset	empty set.

The bindings of type variables are given by a single, global set E of type definitions of the following form:

type $X = T$.

The body of each definition may mention any of the defined variables (in particular, definitions may be recursive). We regard E as a mapping from type variables to their bodies. The regular expression operators $*$, $+$, and $?$ are derived forms as described in Section 2.4. We write $dom(E)$ for the set of defined variables.

We assume a global subtagging relation, a reflexive and transitive relation on labels, written $<$.

As we have defined them so far, types correspond to arbitrary context-free grammars—for example, we can write definitions like:

type $X = a[], X, b[] \mid ()$

Since the decision problem for inclusion between context free languages is undecidable [Hopcroft and Ullman 1979, Theorem 8.12], we need to impose an additional restriction to reduce the power of the system so that types correspond to regular tree languages. Deciding whether an arbitrary context-free grammar is regular is also undecidable [Hopcroft and Ullman 1979, Theorem 8.12],

so we adopt a simple syntactic condition, called *well-formedness*, that ensures regularity. Intuitively, well-formedness allows unguarded (i.e., not enclosed by a label) recursive uses of variables, but restricts them to tail positions. For example, we allow the following type definitions:

```
type X = a[], Y
type Y = b[], X | ()
```

Formally, we define well-formedness in terms of a set of “nontail variables” and an auxiliary set of “top-level variables.” The set $\text{toplevel}(T)$ of top-level variables of a type T is the smallest set satisfying the following equations:

$$\begin{aligned} \text{toplevel}(X) &= \{X\} \cup \text{toplevel}(E(X)) \\ \text{toplevel}(T) &= \emptyset && \text{if } T = \emptyset, (), \text{ or } 1[T'] \\ \text{toplevel}(T|U) &= \text{toplevel}(T) \cup \text{toplevel}(U) \\ \text{toplevel}(T, U) &= \text{toplevel}(T) \cup \text{toplevel}(U). \end{aligned}$$

Likewise, the set $\text{nontail}(T)$ of non-tail variables of a type T is the smallest set satisfying the following equations:

$$\begin{aligned} \text{nontail}(X) &= \text{nontail}(E(X)) \\ \text{nontail}(T) &= \emptyset && \text{if } T = \emptyset, (), \text{ or } 1[T'] \\ \text{nontail}(T|U) &= \text{nontail}(T) \cup \text{nontail}(U) \\ \text{nontail}(T, U) &= \text{toplevel}(T) \cup \text{nontail}(U). \end{aligned}$$

Now, the set E of type definitions is said to be *well formed* if

$$X \notin \text{nontail}(E(X)) \text{ for all } X \in \text{dom}(E).$$

From now on, we assume that a well-formed set E of type definitions is given once and for all.

The semantics of external types is given by the denotation function $\llbracket T \rrbracket$, which is defined as the least solution of the following set of equations¹:

$$\begin{aligned} \llbracket \emptyset \rrbracket &= \emptyset \\ \llbracket () \rrbracket &= \{()\} \\ \llbracket X \rrbracket &= \llbracket E(X) \rrbracket \\ \llbracket 1[T] \rrbracket &= \{1'[v] \mid 1' < 1 \wedge v \in \llbracket T \rrbracket\} \\ \llbracket T, U \rrbracket &= \{v, w \mid v \in \llbracket T \rrbracket \wedge w \in \llbracket U \rrbracket\} \\ \llbracket T | U \rrbracket &= \llbracket T \rrbracket \cup \llbracket U \rrbracket. \end{aligned}$$

These rules are straightforward except that a labeled type with label 1 denotes not only values with label 1 but also those with labels $1'$ smaller than 1 in the sense of subtagging.

Subtyping is defined semantically: two internal types are in the subtype relation iff their denotations are in the subset relation:

$$S <: T \text{ iff } \llbracket S \rrbracket \subseteq \llbracket T \rrbracket.$$

¹A somewhat peculiar property of this type system is that one can define a type that is not syntactically \emptyset but denotes the empty set, by using a “nonterminating” recursive type:

```
type X = a[X]
```

3.2 Internal Form

In the external form, values are arbitrary-arity trees (i.e., each node can have an arbitrary number of children); in the internal form, we consider only binary trees.²

The labels l in the internal form are the same as labels in the external form. Internal (binary) *tree values* (or just *trees*) are defined as follows:

$$t ::= \epsilon \quad \text{leaf} \\ l(t, t) \quad \text{label.}$$

There is a straightforward isomorphism between binary trees and sequences of arbitrary-arity trees. The leaf value ϵ corresponds to the empty sequence, while $l(t, t')$ corresponds to a sequence whose head is labeled 1, with t corresponding to the content of 1 and t' to the remainder of the sequence. For example, from the arbitrary-arity tree

person[name[] , addr[]]

we can read off the binary tree

person(name(ϵ , addr(ϵ , ϵ)), ϵ)

and vice-versa.

To define the internal form of types, we begin as before by assuming a countably infinite set of (*internal*) *type states*, ranged over by X . A (*binary*) *tree automaton* M is a finite mapping from type states to (*internal*) *type expressions*, where a type expression T is a set of either leaves ϵ or labels $l(X, X)$. For convenience, we use the following syntax for internal types:

$$T ::= \emptyset \quad \text{empty set} \\ \epsilon \quad \text{leaf} \\ T \mid T \quad \text{union} \\ l(X, X) \quad \text{label.}$$

(Note that, in the internal form, type states can be used only under labels and labels cannot be nested.) We write $St(T)$ as the set of states appearing in T and $St(M)$ as $\bigcup_{X \in dom(M)} St(M(X))$. We write $card(T)$ for the cardinality of the set T .

There is an exact correspondence between external and internal types (i.e., for any external type, there is an equivalent internal type, and vice-versa) following the same intuition as for values. For example, the external type `person[name[], addr[*]]` corresponds to the internal type $person(X_1, X_0)$ where the states X_1 and X_0 are defined by the automaton M as follows:

$$M(X_0) = \epsilon \\ M(X_1) = \text{name}(X_0, X_2) \\ M(X_2) = \text{addr}(X_0, X_2) \mid \epsilon.$$

²Here, we consider automata accepting binary trees as the internal form of types. An alternative would be to use a more direct formalism, called hedge automata [Murata 2000], which accept arbitrary-arity trees.

The formal translation from external types to internal types is presented in full in Appendix A.

An internal type denotes a set of tree values. When $St(M) \subseteq dom(M)$, the denotation function $\llbracket T \rrbracket_M$ is defined as the least solution of the following set of equations:

$$\begin{aligned} \llbracket \epsilon \rrbracket_M &= \{\epsilon\} \\ \llbracket l(X, X') \rrbracket_M &= \{l'(t, t') \mid l' \prec l \wedge t \in \llbracket M(X) \rrbracket_M \wedge t' \in \llbracket M(X') \rrbracket_M\} \\ \llbracket T \mid T' \rrbracket_M &= \llbracket T \rrbracket_M \cup \llbracket T' \rrbracket_M \\ \llbracket \emptyset \rrbracket_M &= \emptyset. \end{aligned}$$

In the remainder of the article, we assume that a single tree automaton M is given once and for all and shared among all types. From now on, we write simply $\llbracket T \rrbracket$ to mean $\llbracket T \rrbracket_M$.

Just as we did for the external form, we define subtyping between internal types as inclusion between the sets of trees denoted by the types.

$$T <: T' \text{ iff } \llbracket T \rrbracket \subseteq \llbracket T' \rrbracket.$$

Since types are now represented as tree automata, the subtyping problem can be reduced to the inclusion problem of regular tree languages.

THEOREM 1 (COMPLEXITY OF SUBTYPING). *The decision problem for the (internal) subtype relation is EXPTIME-complete.*

PROOF. We use Seidl's result of EXPTIME-completeness of tree automata inclusion [Seidl 1990]. Our formulation is different from his only in that we have subtagging. EXPTIME-hardness of our problem is trivial since any instance of Seidl's problem is an instance of ours (with no subtagging). For the other direction, we can transform our form of tree automata to his by a polynomial-time reduction. Let L be the set of labels appearing in the two given tree automata and n be the number of labeled types appearing in them. We transform each labeled type $l(X, X')$ to $\{l'(X, X') \mid l' \in L \wedge l' \prec l\}$. Comparing the resulting automata with no subtagging is equivalent to comparing the original automata with subtagging. The transformation increases the number of labeled types by $O(|L|n)$. \square

Although the worst-case complexity is quite high, the subtyping algorithm presented below appears to behave satisfactorily on practical examples.

4. SUBTYPING ALGORITHM

This section develops an algorithm for deciding the subtyping relation that is tuned to the domain of typed XML processing, and gives proofs of soundness, completeness, and termination.

4.1 Highlights

There is a classical algorithm for checking inclusion between tree automata. Given two automata M and M' , it works as follows:

- (1) Take the complement $\overline{M'}$ of M' , by constructing a deterministic automaton from (the nondeterministic automaton) M' using a *subset* construction (where each state of the new automaton corresponds to a subset of the states of the original automaton), and then exchanging final and nonfinal states.
- (2) Take the intersection of M and $\overline{M'}$ using a *product* construction.
- (3) Test the emptiness of the result.

(See Comon et al. [1999] for the details of each technique.) In an early implementation of XDuce, we actually used this algorithm. The problem quickly became apparent: the tree automata constructions that it uses will *always* realize any potential exponential blowup in the types involved. This is wasteful, since it is seldom necessary to explore all the states of the automata—in practice, most of the inputs to the subtype checker involve types with a large degree of sharing. For example, consider the inclusion from Section 2.7.

$$\text{Person}^*, \text{Person2}^* <: (\text{Person} | \text{Person2})^*.$$

Since this inclusion holds whatever (and however big) the types *Person* and *Person2* may be, we should be able to check it without looking into their definitions.

In order to exploit this observation, we adopt, as a starting point, Aiken and Murphy's algorithm [Aiken and Murphy 1991]. Their algorithm works in a top-down manner—starting with a pair of types and, at each step, comparing the top-most type constructors and continuing recursively with corresponding sub-components until we reach leaves that require only trivial checks. The main advantage of this top-down algorithm is that it enables numerous optimizations. In particular, we can use reflexivity (e.g., $\text{Person} <: \text{Person}$ in the above example) to cut off large parts of the search. Also, we can cache intermediate results and use them for later comparisons.

This sort of top-down checking is quite standard in subtyping algorithms for conventional type systems. The differences from those algorithms arise from the presence of “untagged” union types in our setting, where two components of a union may have the same outermost label (unlike “tagged” or “disjoint” sum types, as in ML, where all components are required to have distinct tags). For example, we allow types like $l(R_1, S_1) \mid l(R_2, S_2)$.³ To see the difficulties that this raises, suppose we want to check the following inclusion:

$$l(T, U) <: l(R_1, S_1) \mid l(R_2, S_2).$$

What subgoals should we generate? The first rule we might try is the following:

$$\frac{\text{(WEAK-REC)} \quad l(T, U) <: l(R_1, S_1) \quad \text{or} \quad l(T, U) <: l(R_2, S_2)}{l(T, U) <: l(R_1, S_1) \mid l(R_2, S_2)}.$$

However, this rule is too weak. For example, if $T = R_1 \mid R_2$ and $U = S$, then neither premise holds, although we do have $l((R_1 \mid R_2), S) <: l(R_1, S) \mid l(R_2, S)$.

³We use the internal form of types from now on. Also, we use in examples an informal notation that allows nested labels.

To avoid this counterexample, we might try first distributing all unions over labels. For example, to verify $l((R_1 \mid R_2), S) <: l(R_1, S) \mid l(R_2, S)$, we could transform the left-hand side to $l(R_1, S) \mid l(R_2, S)$ and then check whether each clause on the left appears on the right. However, this approach does not work for recursive types, where we would apply distributivity infinitely.

Fortunately, we can overcome the difficulty by reorganizing the clauses, using a simple set-theoretic observation. Let us consider a slightly more general case

$$l(T, U) <: l(R_1, S_1) \mid l(R_2, S_2) \mid l(R_3, S_3)$$

and consider a series of transformations of this relation. (This discussion can be further generalized to cases where the subtype relation to check has an arbitrary number of clauses on the right-hand side.) First, in general, two arguments to a labeled type $l(R, S)$ can be seen as a cross product $R \times S$, which in turn is equal to $(R \times T) \cap (T \times S)$, where the maximal type T denotes the set of all trees. Therefore, the right-hand side of the subtype relation can be rewritten as follows:

$$(l(R_1, T) \cap l(T, S_1)) \mid (l(R_2, T) \cap l(T, S_2)) \mid (l(R_3, T) \cap l(T, S_3)).$$

Using distributivity of intersections over unions, we can turn this disjunctive form to the following conjunctive form.

$$\begin{aligned} & (l(R_1, T) \mid l(R_2, T) \mid l(R_3, T)) \cap \\ & (l(T, S_1) \mid l(R_2, T) \mid l(R_3, T)) \cap \\ & (l(R_1, T) \mid l(T, S_2) \mid l(R_3, T)) \cap \\ & (l(T, S_1) \mid l(T, S_2) \mid l(R_3, T)) \cap \\ & (l(R_1, T) \mid l(R_2, T) \mid l(T, S_3)) \cap \\ & \dots \end{aligned}$$

In each clause of the conjunctive form, if one argument R_i to l appears, then the other argument S_i does not appear, and vice-versa. Therefore, each clause can be rewritten as

$$\left(\bigcap_{i \in I} l(R_i, T) \right) \mid \left(\bigcap_{i \in \bar{I}} l(T, S_i) \right),$$

where I is a subset of $\{1, 2, 3\}$ and \bar{I} is $\{1, 2, 3\} \setminus I$. Since the whole conjunctive form above is the intersection of such formulas for all subsets I of $\{1, 2, 3\}$, the original subtype relation reduces to checking, for each I , that

$$l(T, U) <: \left(\bigcap_{i \in I} l(R_i, T) \right) \mid \left(\bigcap_{i \in \bar{I}} l(T, S_i) \right),$$

or equivalently that

$$l(T, U) <: l\left(\bigcap_{i \in I} R_i, T\right) \mid l\left(T, \bigcap_{i \in \bar{I}} S_i\right).$$

Write R_I for $\bigcap_{i \in I} R_i$ and $S_{\bar{I}}$ for $\bigcap_{i \in \bar{I}} S_i$.

Now, since each labeled type on the right has type T as one of its arguments, the situation becomes easier than the beginning: it suffices to test

$$T <: R_I \text{ or } U <: S_I.$$

To see why, suppose $l(T, U) <: l(R_I, T) \mid l(T, S_I)$ but $T \not<: R_I$ and $U \not<: S_I$. We can find trees $t \in T \setminus R_I$ and $u \in U \setminus S_I$. This means that $l(t, u) \in l(T, U)$ but neither $l(t, u) \in l(R_I, T)$ nor $l(t, u) \in l(T, S_I)$, which contradicts the assumption. (The other direction—if $T <: R_I$ or $U <: S_I$, then $l(T, U) <: l(R_I, T) \mid l(T, S_I)$ —is obvious.)

Our subtyping algorithm is based on these intuitions. In addition, we incorporate several implementation techniques to further improve the efficiency. First, to make effective use of reflexivity checks, we recognize semantically equal types as often as possible and ensure that their representations are physically shared (i.e., we use a form of “hash consing” for types). Second, the subtyping algorithm may backtrack when checking the above formula “ $T <: R_I$ or $U <: S_I$ ”; since, as we will see in Section 5.1, this requires reverting the cache that records intermediate results of subtype checking, we use functional data structures to represent the cache so as to make the reverting quick. Other implementation techniques capture some common special cases for untagged union types to avoid exploring an exponential number of subgoals (which, in general, is incurred by considering all the subsets I). In the following section, we describe the core of the algorithm, postponing the optimizations to Section 5.

4.2 Algorithm

The subtyping algorithm is expressed by two judgments, $\Pi \vdash T <: U \Rightarrow \Pi'$ and $\Pi \vdash^\dagger T <: U \Rightarrow \Pi'$, where Π is a set of pairs of types of the form $R <: S$, called “assumptions.” Both judgments should be read: “assuming that all relations $R <: S$ in Π hold, the algorithm verifies $T <: U$ and yields, in the output set Π' , both the old pairs in Π and the new pairs $T' <: U'$ that have been checked in the process.” At each step of the subtyping algorithm, we add the given pair of types to the assumption set. Later on, when we encounter the same pair, we immediately return “yes,” thus ensuring termination. Of course, we have to be careful not to check the assumption set immediately after storing the given pair, which would incorrectly establish subtyping between any pair of types. This is why we have two different judgments: we switch from one to the other immediately after adding a pair and switch back whenever it is safe. The accumulated assumptions are eventually returned in the output set Π' , which is propagated as the input to other subtype checks, avoiding repeated checks of the same pair. Furthermore, these pairs are reused not only in the process of checking a single inclusion, but all the way through the typechecking of the whole program, thus serving as a cache of all verified subtype relations. (Related discussions of assumptions can be found in Amadio and Cardelli [1993], Brandt and Henglein [1998], and Gapeyev et al. [2000].)

We now give the rules to define the above two judgments. If the pair $T <: U$ of input types is already in the set Π of assumptions, we immediately succeed

(rule HYP). Otherwise, we add the pair to this set and switch judgments (rule ASSUM).

$$\begin{array}{c}
 \text{(HYP)} \\
 \frac{(T <: U) \in \Pi}{\Pi \vdash T <: U \Rightarrow \Pi} \\
 \\
 \text{(ASSUM)} \\
 \frac{\begin{array}{c} (T <: U) \notin \Pi \\ \Pi; T <: U \vdash^\dagger T <: U \Rightarrow \Pi' \end{array}}{\Pi \vdash T <: U \Rightarrow \Pi'}
 \end{array}$$

These two rules are for ensuring termination as well as avoidance of repeated checks of the same pair. In ASSUM, we switch from the judgment of the form $\Pi \vdash T <: U \Rightarrow \Pi'$ to $\Pi \vdash^\dagger T <: U \Rightarrow \Pi'$, preventing the incorrect application of the rule HYP immediately following ASSUM. We keep using the judgment $\Pi \vdash^\dagger T <: U \Rightarrow \Pi'$ in the subsequent rules, and switch back to the judgment $\Pi \vdash T <: U \Rightarrow \Pi'$ in the last rule REC below.

The remaining rules depend on the shapes of the input types. The first three handle the cases where the left-hand side is either an empty set type, a union type, or a leaf type.

$$\begin{array}{c}
 \text{(EMPTY)} \\
 \frac{}{\Pi \vdash^\dagger \emptyset <: T \Rightarrow \Pi} \\
 \\
 \text{(SPLIT)} \\
 \frac{\begin{array}{cc} \text{card}(T) < \text{card}(T \mid T') & \text{card}(T') < \text{card}(T \mid T') \\ \Pi \vdash^\dagger T <: U \Rightarrow \Pi' & \Pi' \vdash^\dagger T' <: U \Rightarrow \Pi'' \end{array}}{\Pi \vdash^\dagger T \mid T' <: U \Rightarrow \Pi''} \\
 \\
 \text{(LEAF)} \\
 \frac{}{\Pi \vdash^\dagger \epsilon <: \epsilon \mid T \Rightarrow \Pi}
 \end{array}$$

If the left-hand side is an empty set type, we simply return since the relation clearly holds (rule EMPTY). If the left-hand side is the union of two types T and T' , we generate two subgoals for these types (rule SPLIT). The intuition behind this rule is the set-theoretic fact that $T \cup T' \subseteq U$ iff $T \subseteq U$ and $T' \subseteq U$. The side conditions $\text{card}(T) < \text{card}(T \mid T')$ and $\text{card}(T') < \text{card}(T \mid T')$ ensure that the algorithm makes progress.

If the left-hand side is a leaf type, we check that the right-hand side also contains a leaf type (rule LEAF). The algorithm can fail only at this rule.

In the remaining cases, the left-hand side is a singleton labeled type.

$$\begin{array}{c}
 \text{(PRUNE)} \\
 \frac{l'(U, U') \notin R \quad l \neq l' \quad \Pi \vdash^\dagger l(T, T') <: R \Rightarrow \Pi'}{\Pi \vdash^\dagger l(T, T') <: l'(U, U') \mid R \Rightarrow \Pi'} \\
 \\
 \text{(PRUNE-LEAF)} \\
 \frac{\epsilon \notin R \quad \Pi \vdash^\dagger l(T, T') <: R \Rightarrow \Pi'}{\Pi \vdash^\dagger l(T, T') <: \epsilon \mid R \Rightarrow \Pi'}
 \end{array}$$

$$\begin{array}{c}
\frac{\text{LEAF}}{\Pi \vdash^\dagger \epsilon <: \epsilon \mid l(Y, \epsilon) \Rightarrow \Pi} \quad \frac{\left(\frac{\text{HYP}}{\Pi \vdash X <: Y \Rightarrow \Pi} \right) \quad \text{and} \quad \left(\frac{\text{fail}}{\vdots} \right) \quad \frac{\text{LEAF}}{\Pi \vdash \epsilon <: \epsilon \Rightarrow \Pi}}{\Pi \vdash^\dagger l(X, \epsilon) <: l(Y, \epsilon) \Rightarrow \Pi} \\
\frac{\Pi \vdash^\dagger \epsilon <: \epsilon \mid l(Y, \epsilon) \Rightarrow \Pi \quad \Pi \vdash^\dagger l(X, \epsilon) <: \epsilon \mid l(Y, \epsilon) \Rightarrow \Pi}{\Pi \vdash^\dagger \epsilon \mid l(X, \epsilon) <: \epsilon \mid l(Y, \epsilon) \Rightarrow \Pi} \\
\frac{}{\emptyset \vdash X <: Y \Rightarrow \Pi}
\end{array}$$

Fig. 1. Example of the subtyping algorithm.

(REC)

$$\frac{\begin{array}{l} \text{for all } 1 \leq j \leq n, l < l_j, \\ \text{for all } 1 \leq i \leq 2^n, \text{ either} \\ \Pi_{i-1} \vdash M(X) <: \bigcup_{j \in I_i^n} M(Y_j) \Rightarrow \Pi_i \quad \text{or} \quad \Pi_{i-1} \vdash M(X') <: \bigcup_{j \in \bar{I}_i^n} M(Y'_j) \Rightarrow \Pi_i \end{array}}{\Pi_0 \vdash^\dagger l(X, X') <: l_1(Y_1, Y'_1) \mid \dots \mid l_n(Y_n, Y'_n) \Rightarrow \Pi_{2^n}}$$

The rules PRUNE and PRUNE-LEAF remove from the right-hand side a leaf type ϵ and all types with labels l' that are not greater than l . The side conditions $l'(U, U') \notin R$ and $\epsilon \notin R$ guarantees that the algorithm proceeds with smaller types. The rule REC handles the other cases—that is, where the right-hand side is the (possibly empty) union of types $l_j(Y_j, Y'_j)$, with all labels l_j greater than the label l —using the intuition explained in Section 4.1. We index the subsets of $\{1, \dots, n\}$ in some arbitrary order from I_1^n to $I_{2^n}^n$. We write \bar{I}_i^n for the complement $\{1, \dots, n\} \setminus I_i^n$. For each index i , we prove that either $M(X)$ is a subtype of the union $\bigcup_{j \in I_i^n} M(Y_j)$, or $M(X')$ is a subtype of the union $\bigcup_{j \in \bar{I}_i^n} M(Y'_j)$.

Figure 1 shows a proof-tree-like diagram that arises when we use the algorithm to check the subtype relation $X <: Y$, where we assume X and Y are defined in the global tree automaton as follows:

$$\begin{aligned}
M(X) &= \epsilon \mid l(X, \epsilon) \\
M(Y) &= \epsilon \mid l(Y, \epsilon).
\end{aligned}$$

In the figure, we write Π for $\{X <: Y\}$. Notice that when $X <: Y$ is checked for the second time, HYP is used; thus the check terminates. Also, checking $\Pi \vdash^\dagger l(X, \epsilon) <: l(Y, \epsilon)$ generates a complex form of premises, some of which fail; thus the check involves backtracking.

Intuitively, we can check that the algorithm terminates by reasoning as follows. The process from EMPTY to PRUNE-LEAF obviously terminates because the cardinalities of the types always decrease. In REC, observe that $X, X', Y_1, \dots, Y_n, Y'_1, \dots, Y'_n$ are always states of the tree automata of the input types. Since the set of such states is finite, the set of unions of such states, for which REC generates subgoals, is also finite. ASSUMP keeps track of all such unions

of states that the algorithm encounters, and HYP ensures termination. This argument is formalized in the next section.

4.3 Proofs

Before arguing the correctness of the algorithm, we need to clarify one subtlety in its definition. The rules in Section 4 can be viewed in two ways: as a “success” relation $\Pi \vdash T <: U \Rightarrow \Pi'$ and as a subtyping algorithm (which we call *Sub* in this section). The former is the relation that is defined by induction on the structure of the internal types using the given rules, whereas the latter is the algorithm that takes an assumption set Π and two internal types T and U as inputs, successively applies the rules in a bottom-up way, and returns another assumption set Π' as an output. Making this distinction is critical, particularly for discussing the algorithm’s termination. When the relation $\Pi \vdash T <: U \Rightarrow \Pi'$ holds, then there is a proof tree validating this relation, and this tree corresponds to *one* possible execution path of the algorithm on the inputs Π , T , and U . However, since the algorithm is nondeterministic, there could be other execution paths for the same inputs. Therefore, the relation $\Pi \vdash T <: U \Rightarrow \Pi'$ does not by itself imply termination.

In addition to the above two concepts, we will need to define a “failure” relation $\Pi \vdash T <: U \Rightarrow \perp$. The proof trees for this relation correspond to execution paths that lead the algorithm to a failure. We will use induction on these proof trees in the completeness proof below. We cannot just use the negation $\Pi \not\vdash T <: U \Rightarrow \Pi'$ of the success relation, since this statement just means that there is no (finite) proof tree validating the success relation, leaving open the possibility that there are only “infinite proof trees”—that is, that the algorithm diverges no matter what set of nondeterministic choices it makes.

Formally, the structure of the definitions and proofs is as follows. We first define (a) the subtyping algorithm, (b) the success relation, and (c) the failure relation. (The first two have already been defined and the last is coming soon.) Then, we prove that

- (1) the algorithm always terminates, and its execution path corresponds to a proof tree in either the success relation or the failure relation (termination),
- (2) the success relation implies the subtype relation (soundness), and
- (3) the failure relation implies the negation of the subtype relation (completeness).

We start with some definitions used in the soundness proof. We first define the height of tree values as follows:

$$\begin{aligned} h(\epsilon) &= 1 \\ h(l(t, t')) &= 1 + \max(h(t), h(t')). \end{aligned}$$

We write $\llbracket T \rrbracket_n$ for $\{t \in \llbracket T \rrbracket \mid h(t) \leq n\}$. The sets $\llbracket T \rrbracket_n$ can be characterized by the following equations:

$$\begin{aligned} \llbracket T \rrbracket_0 &= \emptyset \\ \llbracket \emptyset \rrbracket_{n+1} &= \emptyset \end{aligned}$$

$$\begin{aligned}
\llbracket \epsilon \rrbracket_{n+1} &= \{\epsilon\} \\
\llbracket l(X, X') \rrbracket_{n+1} &= \{l'(t, t') \mid l' \prec l \wedge t \in \llbracket M(X) \rrbracket_n \wedge t' \in \llbracket M(X') \rrbracket_n\} \\
\llbracket T \mid T' \rrbracket_{n+1} &= \llbracket T \rrbracket_{n+1} \cup \llbracket T' \rrbracket_{n+1}.
\end{aligned}$$

We also define a family of subtype relations:

$$T <:_n T' \text{ iff } \llbracket T \rrbracket_n \subseteq \llbracket T' \rrbracket_n.$$

The following properties clearly hold:

- If $T <:_n T'$ for all n , then $T <: T'$.
- If $T <: T'$, then $T <:_n T'$ for all n .
- $T <:_0 T'$ for all types T and T' .

We define Π to be *n-consistent* if $T <:_n U$ for all $(T <: U) \in \Pi$ and *consistent* if Π is *n-consistent* for all n (that is, $T <: U$ for all $(T <: U) \in \Pi$).

The proof of soundness uses the following lemma in the case of **REC**.

LEMMA 2. *Let m be a natural number and A_i, B_i be sets ($1 \leq i \leq m$). Define*

$$\begin{aligned}
I^m &= \{1, \dots, m\} \\
C_m &= \bigcup_{i \in I^m} A_i \times B_i \\
D_m &= \bigcap_{I \subseteq I^m} \left(\bigcup_{j \in I} A_j \times T \right) \cup \left(\bigcup_{j \in I^m \setminus I} T \times B_j \right).
\end{aligned}$$

Then, we have $C_m = D_m$ for all m .

PROOF. By mathematical induction on m .

Case: $m = 0$

$$C_m = D_m = \emptyset.$$

Case: $m \geq 1$

First, D_m can be transformed as follows:

$$\begin{aligned}
D_m &= \bigcap_{I \subseteq I^{m-1}} \left(\bigcup_{j \in I} A_j \times T \right) \cup \left(\bigcup_{j \in I^{m-1} \setminus I} T \times B_j \right) \cup (T \times B_m) \\
&\cap \bigcap_{I \subseteq I^{m-1}} \left(\bigcup_{j \in I} A_j \times T \right) \cup (A_m \times T) \cup \left(\bigcup_{j \in I^{m-1} \setminus I} T \times B_j \right) \\
&= \left(\bigcap_{I \subseteq I^{m-1}} \left(\bigcup_{j \in I} A_j \times T \right) \cup \left(\bigcup_{j \in I^{m-1} \setminus I} T \times B_j \right) \right) \cup ((T \times B_m) \cap (A_m \times T)) \\
&= \left(\bigcap_{I \subseteq I^{m-1}} \left(\bigcup_{j \in I} A_j \times T \right) \cup \left(\bigcup_{j \in I^{m-1} \setminus I} T \times B_j \right) \right) \cup (A_m \times B_m).
\end{aligned}$$

By using the induction hypothesis, the above formula equals to $C_{m-1} \cup (A_m \times B_m) = C_m$. \square

THEOREM 3 (SOUNDNESS). *If $\Pi \vdash T <: U \Rightarrow \Pi'$ and Π is consistent, then $T <: U$ and Π' is consistent.*

PROOF. We prove stronger statements:

- Suppose $\Pi \vdash T <: U \Rightarrow \Pi'$. For all n , if Π is n -consistent, then $T <:_n U$ and Π' is n -consistent.
- Suppose $\Pi \vdash^\dagger T <: U \Rightarrow \Pi'$. For all n , if Π is n -consistent, then $T <:_{n+1} U$ and Π' is n -consistent.

Then the assumption $\vdash T <: U \Rightarrow \Pi'$ entails that $T <:_n U$ and Π' is n -consistent for all n , which imply the result.

The proof proceeds by simultaneous induction on derivations of $\Pi \vdash T <: U \Rightarrow \Pi'$ and $\Pi \vdash^\dagger T <: U \Rightarrow \Pi'$. (We elide the side conditions involving sizes in some of the rules, since they are irrelevant here.)

Case HYP:

$$\frac{(T <: U) \in \Pi}{\Pi \vdash T <: U \Rightarrow \Pi}$$

Immediate.

Case ASSUM:

$$\frac{\begin{array}{c} (T <: U) \notin \Pi \\ \Pi; T <: U \vdash^\dagger T <: U \Rightarrow \Pi' \end{array}}{\Pi \vdash T <: U \Rightarrow \Pi'}$$

Suppose that Π is n -consistent. We must show that $(\Pi; T <: U)$ is also n -consistent. From this, $T <:_n U$ will follow immediately and we can obtain the n -consistency of Π' using the induction hypothesis.

We trivially have $T <:_0 U$. Also, since Π is n -consistent, we can easily show that Π is also i -consistent for all $0 \leq i \leq n$. Furthermore, by the induction hypothesis, for all $0 \leq i \leq n$, if $(\Pi; T <: U)$ is i -consistent, then $T <:_{i+1} U$. By an inner induction on i , it follows that $(\Pi; T <: U)$ is i -consistent for all $0 \leq i \leq n$, and in particular for $i = n$.

Case SPLIT:

$$\frac{\begin{array}{c} \text{card}(T) < \text{card}(T \mid T') \quad \text{card}(T') < \text{card}(T \mid T') \\ \Pi \vdash^\dagger T <: U \Rightarrow \Pi' \quad \Pi' \vdash^\dagger T' <: U \Rightarrow \Pi'' \end{array}}{\Pi \vdash^\dagger T \mid T' <: U \Rightarrow \Pi''}.$$

Suppose Π is n -consistent. Using the induction hypothesis on the first premise, we find that $T <:_{n+1} U$ and Π' is n -consistent. Further using the induction hypothesis on the second premise, we obtain $T' <:_{n+1} U$, together with the n -consistency of Π'' .

It remains to prove that $T \mid T' <_{n+1} U$, that is, $\llbracket T \rrbracket_{n+1} \cup \llbracket T' \rrbracket_{n+1} \subseteq \llbracket U \rrbracket_{n+1}$. This holds because, from $T <_{n+1} U$ and $T' <_{n+1} U$, we have $\llbracket T \rrbracket_{n+1} \subseteq \llbracket U \rrbracket_{n+1}$ and $\llbracket T' \rrbracket_{n+1} \subseteq \llbracket U \rrbracket_{n+1}$.

Case EMPTY:

$$\overline{\Pi \vdash^\dagger \emptyset <: T \Rightarrow \Pi}$$

$\llbracket \emptyset \rrbracket_{n+1} = \emptyset \subseteq \llbracket T \rrbracket_{n+1}$ for any T and n .

Case LEAF:

$$\overline{\Pi \vdash^\dagger \epsilon <: \epsilon \mid T \Rightarrow \Pi}$$

$\llbracket \epsilon \rrbracket_{n+1} = \{\epsilon\} \subseteq \{\epsilon\} \cup \llbracket T \rrbracket_{n+1} = \llbracket \epsilon \mid T \rrbracket_{n+1}$, for any T and n .

Case PRUNE:

$$\frac{l'(U, U') \notin R \quad l \not\prec l' \quad \Pi \vdash^\dagger l(T, T') <: R \Rightarrow \Pi'}{\Pi \vdash^\dagger l(T, T') <: l'(U, U') \mid R \Rightarrow \Pi'}$$

Suppose that Π is n -consistent. By the induction hypothesis, $l(T, T') <_{n+1} R$ and Π' is n -consistent. Therefore, $\llbracket l(T, T') \rrbracket_{n+1} \subseteq \llbracket R \rrbracket_{n+1}$. The result follows from this and $\llbracket R \rrbracket_{n+1} \subseteq \llbracket R \rrbracket_{n+1} \cup \llbracket l'(U, U') \rrbracket_{n+1} = \llbracket l'(U, U') \mid R \rrbracket_{n+1}$.

Case PRUNE-LEAF:

$$\frac{\epsilon \notin R \quad \Pi \vdash^\dagger l(T, T') <: R \Rightarrow \Pi'}{\Pi \vdash^\dagger l(T, T') <: \epsilon \mid R \Rightarrow \Pi'}$$

Similar.

Case REC:

$$\frac{\begin{array}{c} \text{For all } 1 \leq j \leq m, l \prec l_j \\ \text{for all } 1 \leq i \leq 2^n, \text{ either} \\ \Pi_{i-1} \vdash M(X) <: \left|_{j \in I_i^n} M(Y_j) \Rightarrow \Pi_i \quad \text{or} \quad \Pi_{i-1} \vdash M(X') <: \left|_{j \in \overline{I_i^n}} M(Y'_j) \Rightarrow \Pi_i \right. \end{array}}{\Pi_0 \vdash^\dagger l(X, X') <: l_1(Y_1, Y'_1) \mid \dots \mid l_m(Y_m, Y'_m) \Rightarrow \Pi_{2^m}}$$

Let $T = M(X)$, $T' = M(X')$, $U_j = M(Y_j)$, and $U'_j = M(Y'_j)$. Suppose that Π is n -consistent. By assumption, Π_0 is n -consistent. By successive uses of the induction hypotheses, for all $1 \leq i \leq 2^m$, we have either $T <_{n+1} \big|_{j \in I_i^m} U_j$ or $T' <_{n+1} \big|_{j \in \overline{I_i^m}} U'_j$, where Π_i is n -consistent. We must show that

$$l(X, X') <_{n+1} l_1(Y_1, Y'_1) \mid \dots \mid l_m(Y_m, Y'_m)$$

that is,

$$\llbracket l(X, X') \rrbracket_{n+1} \subseteq \llbracket l_1(Y_1, Y'_1) \rrbracket_{n+1} \cup \dots \cup \llbracket l_m(Y_m, Y'_m) \rrbracket_{n+1}.$$

For each j , since $l \prec l_j$, we have

$$\begin{aligned} \llbracket l(Y_j, Y'_j) \rrbracket_{n+1} &= \{l'(t, t') \mid l' \prec l \wedge t \in \llbracket U_j \rrbracket_n \wedge t' \in \llbracket U'_j \rrbracket_n\} \\ &\subseteq \{l'(t, t') \mid l' \prec l_j \wedge t \in \llbracket U_j \rrbracket_n \wedge t' \in \llbracket U'_j \rrbracket_n\} \\ &\subseteq \llbracket l_j(Y_j, Y'_j) \rrbracket_{n+1}. \end{aligned}$$

It is therefore sufficient to prove that

$$\llbracket \mathcal{L}(X, X') \rrbracket_{n+1} \subseteq \llbracket \mathcal{L}(Y_1, Y'_1) \rrbracket_{n+1} \cup \dots \cup \llbracket \mathcal{L}(Y_m, Y'_m) \rrbracket_{n+1}.$$

Notice that, by definition, we have

$$\llbracket \mathcal{L}(X, X') \rrbracket_{n+1} = \{l'(t, t') \mid l' \prec l \wedge t \in \llbracket T \rrbracket_n \wedge t' \in \llbracket T' \rrbracket_n\}$$

and this set is isomorphic to $\{l' \mid l' \prec l\} \times \llbracket T \rrbracket_n \times \llbracket T' \rrbracket_n$; similarly, $\llbracket \mathcal{L}(Y_i, Y'_i) \rrbracket_{n+1}$ is isomorphic to $\{l' \mid l' \prec l\} \times \llbracket U_i \rrbracket_n \times \llbracket U'_i \rrbracket_n$. Thus, the inclusion relation that we are working on proving can be rewritten as follows:

$$\{l' \mid l' \prec l\} \times \llbracket T \rrbracket_n \times \llbracket T' \rrbracket_n \subseteq \bigcup_{1 \leq j \leq m} (\{l' \mid l' \prec l\} \times \llbracket U_j \rrbracket_n \times \llbracket U'_j \rrbracket_n).$$

Since the label part $\{l' \mid l' \prec l\}$ of the products can be canceled on both sides, we only consider the other parts. Hence,

$$\llbracket T \rrbracket_n \times \llbracket T' \rrbracket_n \subseteq \bigcup_{1 \leq j \leq m} (\llbracket U_j \rrbracket_n \times \llbracket U'_j \rrbracket_n).$$

Since $\llbracket U_j \rrbracket_n \times \llbracket U'_j \rrbracket_n = (\llbracket U_j \rrbracket_m \times \mathcal{T}) \cap (\mathcal{T} \times \llbracket U'_j \rrbracket_n)$, where \mathcal{T} is the set of all trees, this inclusion is equivalent to

$$\llbracket T \rrbracket_n \times \llbracket T' \rrbracket_n \subseteq \bigcup_{1 \leq j \leq m} ((\llbracket U_j \rrbracket_n \times \mathcal{T}) \cap (\mathcal{T} \times \llbracket U'_j \rrbracket_n)),$$

which is in turn equivalent to

$$\llbracket T \rrbracket_n \times \llbracket T' \rrbracket_n \subseteq \bigcap_{1 \leq i \leq 2^m} (\bigcup_{j \in I_i^m} (\llbracket U_j \rrbracket_n \times \mathcal{T}) \cup \bigcup_{j \in \overline{I_i^m}} (\mathcal{T} \times \llbracket U'_j \rrbracket_n))$$

using Lemma 2. What we need to show is now that for all $1 \leq i \leq 2^n$,

$$\llbracket T \rrbracket_n \times \llbracket T' \rrbracket_n \subseteq \bigcup_{j \in I_i^m} (\llbracket U_j \rrbracket_n \times \mathcal{T}) \cup \bigcup_{j \in \overline{I_i^m}} (\mathcal{T} \times \llbracket U'_j \rrbracket_n).$$

This follows, for each i , from one of the following inclusions:

$$\begin{aligned} \llbracket T \rrbracket_n &\subseteq \bigcup_{j \in I_i^m} \llbracket U_j \rrbracket_n \\ \text{or } \llbracket T' \rrbracket_n &\subseteq \bigcup_{j \in \overline{I_i^m}} \llbracket U'_j \rrbracket_n. \end{aligned}$$

Each of these, in turn, follows from the definition of one of the inclusions

$$\begin{aligned} T &\prec_n \bigg|_{j \in I_i^m} U_j \\ \text{or } T' &\prec_n \bigg|_{j \in \overline{I_i^m}} U'_j, \end{aligned}$$

which we derived above using the induction hypotheses. \square

We next proceed to the proof of termination, where we show that the algorithm always terminates and yields a successful derivation of subtype judgment or fails. We first define the failure relation $\Pi \vdash T \prec: U \Rightarrow \perp$, read “under assumption Π , the algorithm fails to prove $T \prec: U$.” The relation is defined by

the following set of rules:

$$\begin{array}{c}
\text{(N-ASSUM)} \\
\frac{(T <: U) \notin \Pi \quad \Pi; T <: U \vdash^\dagger T <: U \Rightarrow \perp}{\Pi \vdash T <: U \Rightarrow \perp} \\
\\
\text{(N-SPLIT1)} \\
\frac{\text{card}(T) < \text{card}(T \mid T') \quad \Pi \vdash^\dagger T <: U \Rightarrow \perp}{\Pi \vdash^\dagger T \mid T' <: U \Rightarrow \perp} \\
\\
\text{(N-SPLIT2)} \\
\frac{\text{card}(T) < \text{card}(T \mid T') \quad \text{card}(T') < \text{card}(T \mid T') \quad \Pi \vdash^\dagger T <: U \Rightarrow \Pi' \quad \Pi' \vdash^\dagger T' <: U \Rightarrow \perp}{\Pi \vdash^\dagger T \mid T' <: U \Rightarrow \perp} \\
\\
\text{(N-LEAF)} \\
\frac{\epsilon \notin T}{\Pi \vdash^\dagger \epsilon <: T \Rightarrow \perp} \\
\\
\text{(N-PRUNE)} \\
\frac{l'(U, U') \notin R \quad l \not\prec l' \quad \Pi \vdash^\dagger l(T, T') <: R \Rightarrow \perp}{\Pi \vdash^\dagger l(T, T') <: l'(U, U') \mid R \Rightarrow \perp} \\
\\
\text{(N-PRUNE-LEAF)} \\
\frac{\epsilon \notin R \quad \Pi \vdash^\dagger l(T, T') <: R \Rightarrow \perp}{\Pi \vdash^\dagger l(T, T') <: \epsilon \mid R \Rightarrow \perp} \\
\\
\text{(N-REC)} \\
\begin{array}{c}
\text{For all } 1 \leq j \leq n, l \prec l_j \\
\text{For some } 1 \leq k \leq 2^m, \text{ for all } 1 \leq i \leq k-1, \\
\Pi_{i-1} \vdash M(X) <: \left|_{j \in I_i^m} M(Y_j) \Rightarrow \Pi_i \text{ or } \Pi_{i-1} \vdash M(X') <: \left|_{j \in I_i^m} M(Y'_j) \Rightarrow \Pi_i \right. \\
\text{and both} \\
\Pi_{k-1} \vdash M(X) <: \left|_{j \in I_k^m} M(Y_j) \Rightarrow \perp \text{ and } \Pi_{k-1} \vdash M(X') <: \left|_{j \in I_k^m} M(Y'_j) \Rightarrow \perp \right. \\
\hline
\Pi_0 \vdash^\dagger l(X, X') <: l_1(Y_1, Y'_1) \mid \dots \mid l_n(Y_n, Y'_n) \Rightarrow \perp
\end{array}
\end{array}$$

Note that each rule corresponds to the “negation” of a rule in the success relation $\Pi \vdash T <: U \Rightarrow \Pi'$. N-SPLIT1 corresponds to the case where the first premise of SPLIT fails and N-SPLIT2 corresponds to the case where the first premise of SPLIT succeeds but the second fails. (The asymmetry here arises because the result set of the first premise is used in the second premise.) Similarly, N-REC corresponds to the case where either premise of REC succeeds up to the $(k-1)$ -th iteration and both premises fail at the k th iteration.

Now, the termination property can be summarized and proved as follows:

THEOREM 4 (TERMINATION). *For all Π , T , and U , the algorithm *Sub* terminates and derives either $\Pi \vdash T <: U \Rightarrow \Pi'$ for some Π' or else $\Pi \vdash T <: U \Rightarrow \perp$.*

PROOF. We prove instead the following statement (from which the one above follows directly): for all Π , T , and U , the algorithm *Sub* terminates and derives either $\Pi \vdash^\dagger T <: U \Rightarrow \Pi'$ for some $\Pi' \supseteq \Pi$ or else $\Pi \vdash^\dagger T <: U \Rightarrow \perp$.

Let \mathcal{X} be the set of states in the global automaton. The statement can be proved by induction on the lexicographic order of $(|X| \cdot 2^{|\mathcal{X}|} - |\Pi|, \text{card}(T) + \text{card}(U))$. The crucial observations are that the rules **SPLIT**, **PRUNE**, and **PRUNE-LEAF** always decrease $\text{card}(T) + \text{card}(U)$, and that the rule **REC** is always followed either by **HYP**, which immediately terminates, or by **ASSUM**, which increases Π . Since an element of Π is always a pair of the body of a state and the union of the bodies of some states, the size of Π is bounded by $|X| \cdot 2^{|\mathcal{X}|}$. \square

Finally, we prove the completeness property.

THEOREM 5 (COMPLETENESS). *If $T <: U$, then, for all Π , there is some Π' such that $\Pi \vdash T <: U \Rightarrow \Pi'$.*

PROOF. We prove a stronger statement: If $T <: U$, then $\Pi \vdash T <: U \Rightarrow \Pi'$ and $\Pi \vdash^\dagger T <: U \Rightarrow \Pi'$. We show this by contradiction, that is, if $T <: U$ but $\Pi \not\vdash T <: U \Rightarrow \Pi'$ or $\Pi \not\vdash^\dagger T <: U \Rightarrow \Pi'$, then there is a tree $t \in \llbracket T \rrbracket \setminus \llbracket U \rrbracket$, contradicting $T <: U$.

Since the termination property (Theorem 4) ensures that $\Pi \not\vdash T <: U \Rightarrow \Pi'$ implies $\Pi \vdash T <: U \Rightarrow \perp$, and $\Pi \not\vdash^\dagger T <: U \Rightarrow \Pi'$ implies $\Pi \vdash^\dagger T <: U \Rightarrow \perp$, then we prove the statement in the last paragraph by induction on the height of this “failure” proof tree.

Case N-ASSUM:

$$\frac{(T <: U) \notin \Pi \quad \Pi; T <: U \vdash^\dagger T <: U \Rightarrow \perp}{\Pi \vdash T <: U \Rightarrow \perp}$$

The result follows by the induction hypothesis.

Case N-SPLIT1:

$$\frac{\text{card}(T) < \text{card}(T \mid T') \quad \Pi \vdash^\dagger T <: U \Rightarrow \perp}{\Pi \vdash^\dagger T \mid T' <: U \Rightarrow \perp}$$

By the induction hypothesis, there is $t \in \llbracket T \rrbracket \setminus \llbracket U \rrbracket$, from which the result follows:

Case N-SPLIT2:

$$\frac{\text{card}(T) < \text{card}(T \mid T') \quad \text{card}(T') < \text{card}(T \mid T') \quad \Pi \vdash^\dagger T <: U \Rightarrow \Pi' \quad \Pi' \vdash^\dagger T' <: U \Rightarrow \perp}{\Pi \vdash^\dagger T \mid T' <: U \Rightarrow \perp}$$

By the induction hypothesis, there is $t \in \llbracket T' \rrbracket \setminus \llbracket U \rrbracket$, from which the result follows:

Case N-LEAF:

$$\frac{\epsilon \notin T}{\Pi \vdash^\dagger \epsilon <: T \Rightarrow \perp}.$$

The result follows from $\epsilon \notin \llbracket T \rrbracket$ and therefore $\epsilon \in \llbracket \epsilon \rrbracket \setminus \llbracket T \rrbracket$.

Case N-PRUNE:

$$\frac{l'(U, U') \notin R \quad l \not\prec l' \quad \Pi \vdash^\dagger l(T, T') <: R \Rightarrow \perp}{\Pi \vdash^\dagger l(T, T') <: l'(U, U') \mid R \Rightarrow \perp}$$

By the induction hypothesis, there is $t \in \llbracket l(T, T') \rrbracket$ such that $t \notin \llbracket R \rrbracket$. By definition, $t = k(t_1, t_2)$ for some $k \prec l$ and t_1 and t_2 with $t_1 \in \llbracket T \rrbracket$ and $t_2 \in \llbracket T' \rrbracket$. This implies that $t' = l(t_1, t_2) \in \llbracket l(T, T') \rrbracket$ and $t' \notin \llbracket R \rrbracket$. Since $l \not\prec l'$, we also have $t' \notin \llbracket l'(U, U') \rrbracket$. Therefore, $t' \notin \llbracket l'(U, U') \mid R \rrbracket$.

Case N-PRUNE-LEAF:

$$\frac{\epsilon \notin R \quad \Pi \vdash^\dagger l(T, T') <: R \Rightarrow \perp}{\Pi \vdash^\dagger l(T, T') <: \epsilon \mid R \Rightarrow \perp}$$

The result can be proved similarly to the previous case.

Case N-REC:

$$\frac{\begin{array}{c} \text{For all } 1 \leq j \leq n, l \prec l_j \\ \text{For some } 1 \leq k \leq 2^m, \text{ for all } 1 \leq i \leq k-1, \\ \Pi_{i-1} \vdash M(X) <: \left|_{j \in I_i^m} M(Y_j) \Rightarrow \Pi_i \text{ or } \Pi_{i-1} \vdash M(X') <: \left|_{j \in \overline{I}_i^m} M(Y'_j) \Rightarrow \Pi_i \right. \\ \text{and both} \\ \Pi_{k-1} \vdash M(X) <: \left|_{j \in I_k^m} M(Y_j) \Rightarrow \perp \text{ and } \Pi_{k-1} \vdash M(X') <: \left|_{j \in \overline{I}_k^m} M(Y'_j) \Rightarrow \perp \right. \end{array}}{\Pi_0 \vdash^\dagger l(X, X') <: l_1(Y_1, Y'_1) \mid \dots \mid l_n(Y_n, Y'_n) \Rightarrow \perp}$$

Let $T = M(X)$, $T' = M(X')$, $U_j = M(Y_j)$, and $U'_j = M(Y'_j)$. By the induction hypothesis, there are

$$\begin{array}{l} -t \in \llbracket T \rrbracket \text{ such that } t \notin \left|_{j \in I_i^n} U_j \right| = \bigcup_{j \in I_i^n} \llbracket U_j \rrbracket, \text{ and} \\ -t' \in \llbracket T' \rrbracket \text{ such that } t' \notin \left|_{j \in \overline{I}_i^n} U'_j \right| = \bigcup_{j \in \overline{I}_i^n} \llbracket U'_j \rrbracket. \end{array}$$

Then, we have $l(t, t') \in \llbracket l(X, X') \rrbracket$. Suppose we had $l(t, t') \in \llbracket l_1(Y_1, Y'_1) \mid \dots \mid l_n(Y_n, Y'_n) \rrbracket$. Then there would be some j such that $l(t, t') \in \llbracket l_j(Y_j, Y'_j) \rrbracket$, which implies $t \in \llbracket U_j \rrbracket$ and $t' \in \llbracket U'_j \rrbracket$. This contradicts either $t \notin \bigcup_{j \in I_i^n} \llbracket U_j \rrbracket$ or $t' \notin \bigcup_{j \in \overline{I}_i^n} \llbracket U'_j \rrbracket$. \square

5. IMPLEMENTATION

Our subtype checker implementation embodies the rules in the previous section, plus a number of optimizations specialized to the subtyping problems that arise in practice in the domain of typed XML processing. This section describes several of these techniques. They are categorized into low-level,

representational techniques and higher-level heuristics inspired by set-theoretic observations.

5.1 Low-Level Techniques

In order for the subtyping algorithm to be able to exploit fast reflexivity checks (as mentioned in 4.1), it is crucial to share the physical representations of internal types as much as possible. To this end, we recognize some semantically equal external types and translate them to the same internal type. The equalities that we exploit are commutativity, associativity, and idempotency of the union operator, and associativity of the concatenation operator. In order to make the recognition of equal types quick, we represent external types in such a way that equal types are *structurally* equal. We represent the union of types as a set of types, with nested unions flattened; thus, $((R|S) | (T|U))$ and $(U | (S | (T|R)))$ can be recognized as equal, for example. Since we use only union and equality for the operations on such sets, a suitable representation is a sorted list, which allows us to perform these two operations in linear time. (We sort the types in the list by their hash values, which are computed from the structures of the types.) We represent the concatenation of types simply by a list of types, with nested concatenations flattened and the empty sequence removed; thus, $((R, ()), S), (T, U))$ and $(R, (S, (T, U)))$ can be recognized as equal, for example. To further improve the speed of equality tests, we use *hash consing*, which associates each type expression with its integer hash value, so that equality can be quickly checked in most cases by comparing their hash values. (An alternative approach to hash consing might be to use *multiset discrimination* [Cai and Paige 1995], but we have not tried this yet.)

In the internal form of types, we use cyclic structures to represent recursion. Compared to explicit use of recursion variables and table lookups, this representation not only makes the dereferencing slightly faster, but simplifies the implementation.

We need to be a little careful about the representation of the sets of assumptions Π described in Section 4.2. A hash table might initially appear to be a suitable representation, but it is not. To see why, recall that in REC, we have the premises “ $T <: \dots$ or $T' <: \dots$.” Suppose that the first premise fails. Then we try the second premise, to which we need to pass the original set of assumptions. However, if we represent the set of assumptions by a hash table, then the hash table available after the first premise may contain some assumptions that have been added during the processing of the first premise. Since those assumptions may be wrong, we have to remove them from the hash table. Or, we could copy the whole table before trying the first premise. Either would be very expensive. Instead, we use a functional representation of sets. Since the required operations here are insertion and membership testing (unlike the unions of types described above), we use *balanced binary trees*. Operations for balanced binary trees take logarithmic time in the size of the set, as opposed to constant time for hash tables. We have not examined this factor, but we believe that this is negligible in practice since the number of assumptions stored is usually not large (less than 500 in our test cases, as shown in Section 6).

5.2 High-Level Techniques

5.2.1 Literal Equality. In the implementation of a subtype checker for any type system, the most trivial optimization is, before going deeply into the structures, checking if the given types are literally equal (we use a physical equality check in the implementation). In the presence of union types, this can be slightly generalized, using the fact that $T <: T \mid U$. In our implementation, we use the following rule

$$\frac{(\text{TRIV}) \quad \Pi \vdash U <: T \mid R \Rightarrow \Pi'}{\Pi \vdash T \mid U <: T \mid R \Rightarrow \Pi'}$$

which can be seen as a combination of SPLIT and $T <: T \mid U$. (We try this rule before HYP.) (Notice that this rule simply takes the difference between two sets of types. Since we represent the union of types as a sorted list, the difference can be calculated by a linear scan.)

5.2.2 Empty Type Elimination. In this optimization (which we call EMP), before starting the subtyping algorithm, we perform a preprocessing step that eliminates all the types denoting the empty set. Since the subtyping algorithm can now assume that any type it encounters is not empty, some tests can be short cut. For example, two subgoals of the form $T <: \emptyset$ in the rule REC need not be generated. In particular, in the special case of REC where the right-hand side is just a single labeled type— $l(T, U) <: l'(R, S)$ —we only need to check $T <: R$ and $U <: S$ with $l < l'$, which involves no backtracking at all.

The preprocessing of empty type elimination can be highly tuned. In theory, identifying and eliminating empty states can be performed in linear time [Comon et al. 1999]. However, we use a simpler but potentially quadratic algorithm, which seems to perform well enough in practice. (See Section 6.) We have not compared these two algorithms yet.

5.2.3 Merging Labeled Types. In order to make the previous optimization more effective, we merge types on the right-hand side when either the first or second arguments are the same:

$$\frac{(\text{MERGE1}) \quad \Pi \vdash^\dagger l(T, U) <: l'(R, (S \mid E)) \Rightarrow \Pi' \quad l < l'}{\Pi \vdash^\dagger l(T, U) <: l'(R, S) \mid l'(R, E) \Rightarrow \Pi'}$$

$$\frac{(\text{MERGE2}) \quad \Pi \vdash^\dagger l(T, U) <: l'((R \mid S), E) \Rightarrow \Pi' \quad l < l'}{\Pi \vdash^\dagger l(T, U) <: l'(R, E) \mid l'(S, E) \Rightarrow \Pi'}$$

(We try these rules before SINGLE.) In our experience, the first case is more common than the second case. This is because, in the external form of types, labels of the same name often have the same content type. (When we “import” existing DTDs and interpret them as regular expression types, labels of the same name are even *required* to have the same content type.) Therefore, we check MERGE1 first and then MERGE2.

5.2.4 Default Case. If the type on the right-hand side in REC has the form $U \mid R_1 \mid \dots \mid R_n$ and U is larger than any R_i , then we only need to compare the left hand side with this largest type U :

$$\begin{array}{c} \text{(SUPER)} \\ \frac{1 \leq i \leq n \quad \Pi_i \vdash^\dagger R_i <: U \Rightarrow \Pi_{i+1} \quad \Pi_{n+1} \vdash^\dagger T <: U \Rightarrow \Pi'}{\Pi_1 \vdash^\dagger T <: U \mid R_1 \mid \dots \mid R_n \Rightarrow \Pi'} \end{array}$$

(We try this rule before REC after MERGE1/2.) This typically happens when the programmer writes a “default” case in a pattern match, which is given a type (e.g., Any) that covers all the other cases.

In principle, this optimization can generate, in the course of searching for the largest type, so many subgoals that the cost surpasses the gain. However, we have not found such a situation in practice so far (see Section 6). This is probably because most of the cases are handled by the previous optimizations, and because the general rule REC (which we would have to use if not using SUPER) is often more expensive.

5.2.5 Type Any. Although the type Any can be encoded using the special label \sim , as described in Section 2.5, it is useful to represent Any as a special state in the internal form, so that we can use the following rule:

$$\begin{array}{c} \text{(ANY)} \\ \hline \Pi \vdash^\dagger T <: \text{Any} \Rightarrow \Pi \end{array}$$

(We try this rule before SPLIT.) Any is used quite heavily in our programs, especially when we do not want to specify precise types in patterns. Without this optimization, we would have to traverse the whole structure of the left-hand side type T , where every step just compares a subphrase of T with Any.

Using these heuristics together, almost all uses of the rule REC are eliminated in our example programs (see the following section).

6. PRELIMINARY EXPERIMENTS

We have incorporated the subtyping algorithm described above in a prototype implementation of the XDuce language. XDuce is a simple first-order functional language; a typical program consists of a collection of type declarations and recursive functions that use pattern matching to analyze input values. XDuce can parse external DTDs, interpreting them as regular expression type declarations. Some of our applications use this feature to incorporate fairly large DTDs from real-world XML applications.

In this section, we present the results of some preliminary performance measurements of our implementation. In the experiments, we are interested in (1) the wall-clock time that our algorithm takes to typecheck various application programs (subtype checks consume most of this time), and (2) the separate effects of each high-level optimization. The platform for our experiment was a Sun Enterprise 3000 (250MHz UltraSPARC) running SunOS 5.7.

Table I. Applications and Measurement Results

Application	# of Lines		Subtyping Time (sec)		Subtype Alg. Internals	
	XDuce	DTD	Total	Main	States	Assumptions
Bookmarks	310	1197	0.48	0.018	756	133
Diff	355	—	0.039	0.014	276	165
Html2Latex (strict)	307	989	0.58	0.22	783	345
Html2Latex (transitional)	312	1197	0.88	0.36	946	433
Html2Latex (frameset)	323	1226	0.87	0.34	975	454

Our test suite consists of three small applications written in XDuce:

Bookmarks. Takes as input a Netscape bookmarks file of type `Bookmarks`, which is a small subset of the type `HTML`. It extracts a particular folder named “Public,” adds a table of contents at the front, and inserts links between the contents and the body. The type of the result is the full `HTML` type.

Html2Latex. Takes an `HTML` file (of type `HTML`) and converts it into `LaTeX` (a value of type `String`) by interpreting some of the markup commands.

Diff. Implements Chawathe’s “tree diff” algorithm [Chawathe 1999]. It takes a pair of XML files of type `Xml`, which is the type of all XML documents, and returns a tree with annotations indicating whether each subtree has been retained, inserted, deleted, or changed between the two inputs.

The `HTML` type (more precisely, `XHTML`, which is an XML implementation of `HTML`) is currently one of the largest schemas that are in real use. This makes it an excellent benchmark case for our implementation. There are actually three versions of `XHTML`: `XHTML-strict`, `XHTML-transitional`, and `XHTML-frameset`; accordingly, our `Html2Latex` application comes in three versions. The first is smallest and the third is slightly larger than the second.

The first group of columns in Table I shows the number of lines in XDuce code (counting functions and types written in XDuce syntax, but not external DTDs), and the number of lines in external DTDs (if used). The difference between the three versions of `Html2Latex` is mainly in the number of lines of DTDs. The column “total” in the table shows the total time spent by the subtyping algorithm during the type checking of the whole program. It includes conversion from the external form to the internal form (INT), empty type elimination (EMP), and the main subtyping algorithm (SUB). The optimizations are all turned on for this table. The column “main” shows the time spent by the main algorithm SUB. The table gives two more columns: “states” and “assumptions.” The “states” column indicates the number of states of the internal form stored in the system, and the “assumps” indicates the number of pairs stored in the set of assumptions.

As the table indicates, the speed of type checking is acceptable for these applications. In particular, it takes less than one second to type check programs involving the full `HTML` type. Also, the space consumed by the subtyping algorithm indicated by the number of assumptions is relatively small, despite its potential of blow-up.

Notice that for the applications that we use, the ratio of the “main” column to the “total” column is relatively small, even though subtyping takes exponential time in the worst case. This ratio would not be so small if we did not use any of our high-level optimizations, as we will discuss next.

We ran a further set of measurements to evaluate the effects of each high-level optimization. The measurement strategy is as follows: We start with a subtype checker that implements the set of rules in Section 4.2 and the low-level techniques described in Section 5.1 (NONE). We then turn on the high-level optimizations one by one and measure the performance. We add high-level optimizations in this order: EMP, TRIV, MERGE1, MERGE2, SUPER and ANY.

The graphs in Figure 2 show how the behavior of our algorithm varies as we add each optimization. We only consider the “transitional” version of Html2Latex since the graphs look alike for other versions. Each graph provides two kinds of information. The first is given by bars indicating the transition of the time spent by the algorithm. Each bar in the graphs has three components corresponding to each phase of the subtyping procedure. We are mainly interested in SUB. The others are only presented so as to show the ratio of SUB in the total time. The times, except for SUB, should not depend on the optimization switches but one can see noises in the graphs (especially in Diff). We suspect that this is due to garbage collection since this occurs consistently over several runs (we have not confirmed this by any further experiment). The second piece of information is given by a line indicating the number of times the rule REC is used (NDET). (For NONE and EMP, these numbers are sometimes too big to fit on the graph.)

We first notice that EMP and TRIV are always effective. (For Diff, unfortunately, the total time becomes bigger with MERGE1 because of the “GC noise.”) ANY is effective in Bookmarks. The effects of MERGE1 and MERGE2 are unclear in these graphs.

The number of times the rule REC is used almost always decreases each time an optimization is added; by using all the optimizations, this rule can almost be avoided. However, even though REC rule is usually avoided, the savings in wall-clock time is less. This indicates that, in some cases, the cost of the optimization and that of blindly using REC rule may be comparable.

Also, notice that the cost of empty type elimination (EMP) is very low.

The experimental results indicate that the efficiency of our subtyping algorithm is quite acceptable in our small applications. In particular, it is quite encouraging that it works well even in the applications using XHTML. This is a big step forward from our earliest version of XDuce, which used a naive text-book algorithm for tree automata inclusion (which is mentioned in the beginning of Section 4.1) and which performed terribly on such programs. However, in order to conclude that the present algorithm is adequate for a wide range of programs, benchmarks using much bigger applications would be required.

7. RELATED WORK

Static typing of programs for XML processing has been approached from several different angles. One popular idea is to embed a type system for XML in an

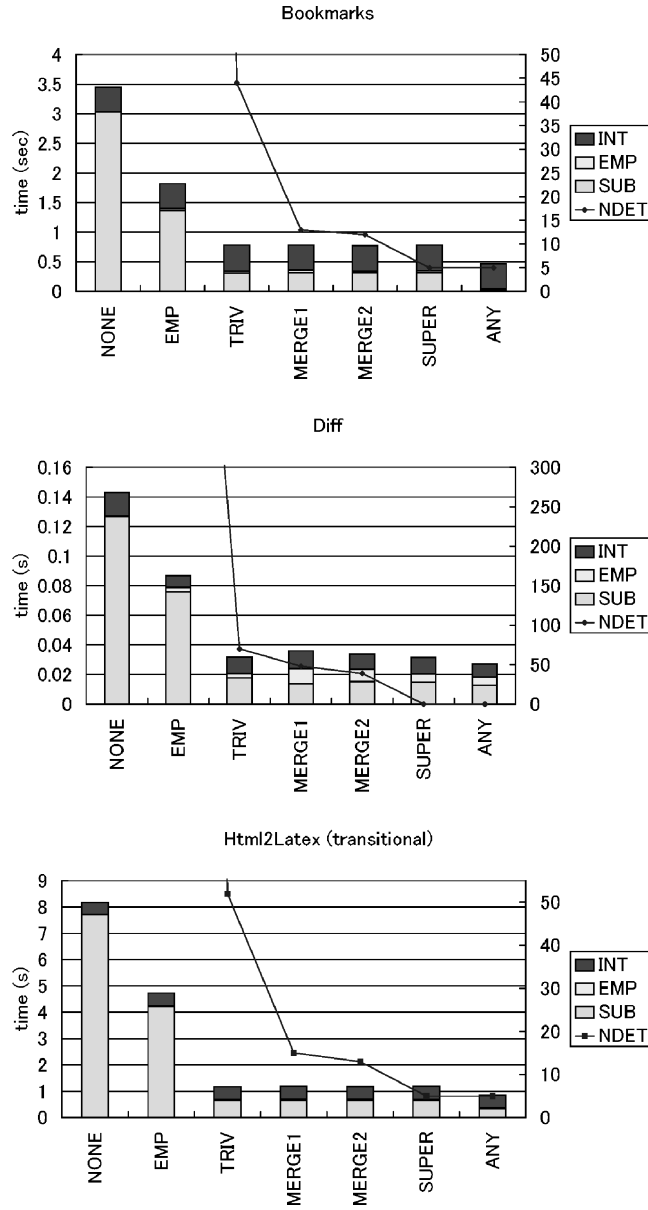


Fig. 2. The effects of high-level optimizations.

existing typed language. The advantage is that we can enjoy not only static type safety, but also all the other features provided by the host language. The cost is that XML values and their corresponding schemas must somehow be “injected” into the value and type spaces of the host language; this usually involves adding more layers of tagging than were present in the original XML documents, which inhibits subtyping. The lack of subtyping (or availability of only restricted forms

of subtyping) is not a serious problem for simple traversal of tree structures; it becomes a stumbling block, though, in tasks like the “database integration” that we discussed in Section 2.7, where ordering-forgetting subtyping and distributivity were critically needed.

A recent example of the embedding approach is Wallace and Runciman’s [1999] proposal to use Haskell as a host language for XML processing. The only thing they add to Haskell is a mapping from DTDs into Haskell datatypes. This allows their programs to make use of other mechanisms standard in functional programming languages, such as higher-order functions, parametric polymorphism, and pattern matching. However, they do not have any notion of subtyping. A difference in the other direction is that our type system does not currently support higher-order functions or parametric polymorphism. (We are working on both of these extensions, but they are nontrivial.)

Meijer and Shields propose the typed functional language $\text{XM}\lambda$ for XML processing [Meijer and Shields 1999; Shields and Meijer 2001]. Although their type system stems from Haskell’s, they attain additional flexibility required in XML processing by incorporating, instead of subtyping, extensible records and variants based on row polymorphism.

The query language YAT [Cluet and Siméon 1998] has a type system similar to regular expression types. They also provide a notion of subtyping that can make programs flexible against changes of types. However, their subtyping is somewhat weaker than ours: they can handle the first example (“database evolution”) in Section 2.7, but since their subtyping lacks distributive laws, they cannot treat the second example (“database integration”) given in the same section.

Since its initial publication, our work on regular expression types has influenced proposals by other researchers. In particular, Fernández et al. [2001] propose XML Query Algebra for the basis of XML query processing and optimization, and they use our regular expression types in their type system and our subtyping algorithm in their implementation.

Milo et al. [2000] have studied a typechecking problem for their general framework called *k-pebble Tree transducers*, which can capture a wide range of query languages for XML. The types there are based on tree automata and conceptually identical to regular expression types. Papakonstantinou and Vianu [2000] present a typechecking algorithm for their query language *loto-ql*, where the algorithm uses extensions to DTDs. One of their extensions is equivalent to tree automata.

Although schema languages for XML do not treat static verification of programs, the type structures in these languages and regular expression types are worth discussing. Since schema languages are too many to enumerate, we consider the following representative ones: DTD [Bray et al. 2000], XML-Schema [Fallside 2001], DSD (Document Structure Description) [Klarlund et al. 2000], RELAX [Murata 2001], TREX [Clark 2001], RELAX NG [Clark and Murata 2001], and SOX (Schema for Object-Oriented XML) [Davidson et al. 1999]. As our regular expression types do, all of these use regular expressions or similar mechanisms for describing sequences. Both our regular expression types and RELAX (and its relatives TREX and RELAX NG) are equivalent to tree

automata (more precisely, nondeterministic tree automata; see Comon et al. [1999] for a detailed classification of tree automata), and therefore these have the same expressiveness. The other schema languages are more restrictive. Some of the above schema languages consider subtyping. XML-Schema has a “restriction” subtyping (which yields a subtype by decreasing the number of choices in an existing type), and both XML-Schema and SOX have an “extension” subtyping (which yields a subtype by adding new fields to the tail, similarly to inheritance in object-oriented languages). Both forms of subtyping are subsumed by ours. Kuper and Siméon [2001] define a schema language similar to regular expression types; they provide a subtype relation that is weaker than ours but allows efficient storage layout for XML databases.

Several researchers have developed type systems for query languages for tree-like data that are similar to XML except that they have no ordering among children nodes (some of the formalisms also allow cycles). Their unorderedness makes these type systems quite different from those for XML. In the type system studied by Buneman et al. [1997], types are graph structures and their conformance and subtype relations are defined in terms of graph simulation (which is weaker than the inclusion relation). The query language Lorel uses a somewhat similar type system called DataGuide [Goldman and Widom 1997].

Our investigation of regular expression types was originally motivated by an observation by Buneman and Pierce [1998] that untagged union types correspond naturally to forms of variation found in semistructured databases. The difference from the present work is that they study unordered record types instead of ordered sequences and do not treat recursive types.

Our subtyping algorithm can be seen as an extension to subtyping algorithms for conventional type systems. In particular, the technique used in our algorithm for keeping track of a set of “assumptions” to detect termination (Section 4.2) can be found in many subtyping algorithms for simple recursive types [Amadio and Cardelli 1993; Brandt and Henglein 1998; Gapeyev et al. 2000]. The ingredient that is rarely found in classical algorithms is the complex rule shown Section 4.2 for the “untagged” union type operator.

As we discussed above, our subtyping algorithm can be viewed as a variant of Aiken and Murphy’s algorithm [Aiken and Murphy 1991]. They actually treat a more general problem called set-constraint solving. In their setting, types can contain free variables and the goal is to obtain a substitution for the variables that satisfies the given set-constraints, if they are satisfiable. In our setting, on the other hand, types do not contain free variables and the goal is to know only whether the constraints are satisfiable. Thus, we can solve our subtyping problem by using their algorithm and removing the rules related to free variables. They do not, however, prove correctness in their paper—in particular, they have no completeness proof since their algorithm is indeed incomplete. (Incompleteness means that the algorithm may answer “no” even if the constraints are solvable.) What we have proved is that their algorithm is actually complete if we restrict types to contain no free variables.

In their framework for *refinement types* for ML, Freeman and Pfenning [1991] adopt regular tree languages to express more precise types than ML datatypes and use tests for inclusion between tree automata in type

checking. Davies [2000] addresses the efficiency issue of the inclusion test in this context.

Damm [1994] considers a type system with union, intersection, and recursive types where the subtype relation is defined in a semantic way. He presents a decision procedure for subtype checking that first encodes types into regular tree expressions and then uses existing set-constraint solving algorithms. Furthermore, he treats higher-order functions, which we have not handled yet. On the other hand, he does not address the efficiency of the subtyping algorithm.

8. CONCLUSIONS

We have proposed regular expression types for XML processing, arguing that set-inclusion-based subtyping and subtagging yield useful expressive power in this domain. We developed an algorithm for subtyping that can exploit sharing among type expressions, giving soundness, completeness, and termination proofs. By incorporating several optimization techniques, our algorithm runs at acceptable speeds on several applications involving fairly large types, such as the complete DTD for HTML documents.

Our work on type systems for XML processing has just begun. In the future, we hope to incorporate other standard features from functional programming, such as higher-order functions and parametric polymorphism. The combination of these features with regular expression types raises some difficult problems. For function types, our current approach—define subtyping by inclusion of the semantics of types and reduce it to the decidability of tree automata inclusion—does not easily extend simply because functions are not trees. (Some achievements in this direction have recently been reported by the CDuce team [Frisch et al. 2002].) Also for polymorphism, our current scheme needs to be substantially extended since usual tree automata do not have any concept corresponding to “type variables.” A promising direction might be to incorporate ideas from *tree set automata* [Gilleron et al. 1999], though we have not gone far.

APPENDIX

A. TRANSLATION OF TYPES

This section presents an algorithm for translating from the external to the internal form of types and proves its soundness, completeness, and termination. (A related algorithm can be found in Hornung [1996]).

A.1 Algorithm

We first show the translation of values, which is straightforward.

$$\begin{aligned} ts(()) &= \epsilon \\ ts(1[v_1], v_2) &= 1(ts(v_1), ts(v_2)) \end{aligned}$$

We now consider the translation of types. Let us first illustrate it by an example. Consider the following external type:

$$a[]*, d[]$$

By expanding the abbreviation $a[]*$, this type is equivalent to $(X, d[])$ where:

$$\text{type } X = (a[], X) \mid ()$$

Now, we want to compute the internal type corresponding to $(X, d[])$. For this, we transform it in such a way that all head labels are revealed. We first expand X to its definition and then use distributivity of the union operator, associativity of the concatenation operator, and neutrality of the empty sequence⁴:

$$\begin{aligned} & (X, d[]) \\ \Rightarrow & ((a[], X) \mid ()), d[]) \\ \Rightarrow & ((a[], X), d[]) \mid (((), d[])) \\ \Rightarrow & ((a[], (X, d[])) \mid d[]) \end{aligned}$$

Since the head labels a and d are revealed now, this type can be translated to an internal type:

$$a(X_0, X_1) \mid d(X_0, X_0)$$

Since the content of a and both the content and the remainder of d are all the empty sequence, we can share the internal types corresponding to these by translating them all as the single state X_0 and associating it with ϵ :

$$M(X_0) = \epsilon$$

The remainder of a , that is, the type $(X, d[])$, is translated as the state X_1 . To associate with this state, we need an internal type corresponding to $(X, d[])$. By transforming it exactly in the same way as above, we obtain $((a[], (X, d[])) \mid d[])$. This type is then translated to the internal type $a(X_0, X_1) \mid d(X_0, X_0)$. Here, note that we can reuse the states X_0 and X_1 since we have already translated the empty sequence to X_0 and the type $(X, d[])$ to X_1 . Thus, we obtain

$$M(X_1) = a(X_0, X_1) \mid d(X_0, X_0).$$

Before presenting the general translation algorithm, it is convenient to make a small restriction on the syntax of types. In the given type definitions, we make sure that all recursive uses of type names must be preceded by nonnullable type, where a nullable type is one whose denotation contains the empty sequence. For example, both of the type definitions of X and Y in the following are rejected.

```
type X = X
type Y = a[]*, Y | ()
```

Semantically, these type definitions make sense since we can interpret X to be the empty set and Y to be the set of sequences of zero or more $a[]$ s. Nonetheless, we reject such type definitions, since handling these would make our formulation complicated, while these have no practical interest. (There are more straightforward ways of writing both the empty set type and the type of sequences of zero or more $a[]$ s.)

⁴This technique of revealing head labels is similar to *derivatives* [Brzozowski 1964], which are often used in construction of string automata from regular expressions.

Formally, we define a function *nohead* that takes a type and a set of type names and answers true if and only if all recursive uses of type names in the type are preceded by non-nullable types:

$$\begin{aligned}
nohead(\emptyset)\sigma &= \text{true} \\
nohead(\cdot)\sigma &= \text{true} \\
nohead(1[T])\sigma &= \text{true} \\
nohead(T_1 | T_2)\sigma &= nohead(T_1)\sigma \wedge nohead(T_2)\sigma \\
nohead(X)\sigma &= \begin{cases} nohead(E(X))(\sigma \cup \{X\}) & (X \notin \sigma) \\ \text{false} & \text{otherwise} \end{cases} \\
nohead(\emptyset, T)\sigma &= \text{true} \\
nohead(\cdot, T)\sigma &= nohead(T)\sigma \\
nohead(1[T_1], T_2)\sigma &= \text{true} \\
nohead((T_1, T_2), T_3)\sigma &= nohead(T_1, (T_2, T_3))\sigma \\
nohead((T_1 | T_2), T_3)\sigma &= nohead(T_1, T_2)\sigma \wedge nohead(T_1, T_3)\sigma \\
nohead(X, T)\sigma &= \begin{cases} nohead(E(X), T)(\sigma \cup \{X\}) & (X \notin \sigma) \\ \text{false} & \text{otherwise} \end{cases}
\end{aligned}$$

Then, we require:

$$nohead(E(X))\emptyset = \text{true} \quad \forall X \in \text{dom}(E)$$

The main translation procedure takes an external type T_0 and a set E of type definitions, and computes an internal type T_0 and a tree automaton M . The procedure works by double loops where the outer loop constructs the automaton and the inner loop computes an internal type expression, which will be associated with a state in the automaton. Each state corresponds to an external type expression from which the internal type associated with the state is computed; to make this correspondance explicit, we write each state in the form X_T .

The translation function *ts* (“the inner loop”) takes an external type and returns an internal type. The rules below define the function *ts*:

$$\begin{array}{ll}
ts(\emptyset) = \emptyset & \text{TR-EMP1} \\
ts(\cdot) = \epsilon & \text{TR-EPS1} \\
ts(1[T]) = 1(X_T, X_{\cdot}) & \text{TR-LAB1} \\
ts(T_1 | T_2) = ts(T_1) | ts(T_2) & \text{TR-OR1} \\
ts(X) = ts(E(X)) & \text{TR-NM1} \\
ts(\emptyset, T) = \emptyset & \text{TR-EMP2} \\
ts(\cdot, T) = ts(T) & \text{TR-EPS2} \\
ts(1[T_1], T_2) = 1(X_{T_1}, X_{T_2}) & \text{TR-LAB2} \\
ts((T_1, T_2), T_3) = ts(T_1, (T_2, T_3)) & \text{TR-ASSOC} \\
ts((T_1 | T_2), T_3) = ts(T_1, T_3) | ts(T_2, T_3) & \text{TR-OR2} \\
ts(X, T) = ts(E(X), T) & \text{TR-NM2}
\end{array}$$

These rules simply generalize the operations that we have seen in the example. Rule TR-ASSOC uses associativity and rule TR-OR2 uses distributivity. Also, rule TR-EPS2 uses the neutrality of \cdot . Rules TR-NM1 and TR-NM2 unfold the type name X to their definitions. (Since the syntactic restriction described above

ensures that any type name never recurs without non-nullable types in front, rules TR-NM1 and TR-NM2 never unfold the same type name more than once. This ensures that the function ts terminates.) By using all these rules, the head of a sequence is eventually revealed, and turned (by rules TR-LAB1 and TR-LAB2) into an internal label type whose “car” state corresponds to the content type of the label and whose “cdr” state corresponds to the remainder of the sequence. The empty sequence is turned into the leaf transition (by rules TR-EPS1 and TR-LAB2).

The outer loop proceeds as follows. The translation begins with setting M to the empty automaton and computing T_0 from T_0 by using the above function ts . The resulting internal type T_0 may contain some states that are not yet in the domain of M . For each such state X_T , we calculate an internal type T from the external type T by the ts function and add the mapping $X_T \mapsto T$ to M . We repeat this process until the states appearing in T_0 and M are all in the domain of M . Thus, the result of the whole translation satisfies the following:

$$\begin{aligned} T_0 &= ts(T_0) \\ M(X_T) &= ts(T) \quad \forall X_T \in dom(M) \\ dom(M) &\supseteq St(T_0) \cup St(M). \end{aligned}$$

THEOREM A.1 (SOUNDNESS AND COMPLETENESS). *Suppose that a given tree automaton M satisfies $M(X_T) = ts(T)$ for all $X_T \in dom(M)$ and $dom(M) \supseteq St(M)$. If $ts(v) = t$ and $ts(T) = T$, then $v \in \llbracket T \rrbracket$ implies $t \in \llbracket T \rrbracket$ and vice-versa.*

PROOF. By lexicographic induction on the pair of the height $h(t)$ of the tree t and the size of the derivation of $ts(T) = T$, with the case analysis on the rule used in the derivation. \square

A.2 Termination

In a classical algorithm dealing with recursive types, the type expressions encountered by the algorithm are always subphrases of the original type expression. By using the fact that such subphrases are finite in number, termination of such an algorithm can easily be proved. However, the types encountered by our translation algorithm are not necessarily subphrases of the original type, since the algorithm may restructure types from their original form. For example, consider the following type definitions:

```
type X = a[], X | ()
type Y = X, b[]
```

Let us see the steps involved in the translation of the type Y . We first unfold Y , which yields $(X, b[])$. The next step unfolds X , and yields $(a[], X \mid ()), b[]$. Note that this type does not appear in the above type definitions. (But note that this type is the *concatenation* of some subphrases appearing there.) Nonetheless, the number of types encountered by the algorithm is finite. Let us continue the translation. We expand the type $(a[], X \mid ()), b[]$ by distributivity, and then proceed to each of the clauses $((a[], X), b[])$ and $((()), b[])$. The second clause ends up with $b[]$. On the other hand, the first clause transforms

to $(a[], (X, b[]))$ by associativity and finally leads to $(X, b[])$. Since we have already seen this type, we can stop here. (Notice that the type name X was once unfolded before, but X was followed by the same type $b[]$ at that unfolding as it is now. This is ensured by the well-formedness of the type definitions, which restricts unguarded, recursive uses of type names to be in tail positions.)

For our proof of termination, we devise a special transition system where each transition represents the fact that, if the algorithm encounters the first type expression in one step, then it will encounter the second in a later step. We concentrate on what the algorithm encounters and abstract away what it returns. The reason is that outputs of the translation are never used by the translation itself and therefore termination only depends on inputs to the translation.

The terms over which the transition function is defined are type expressions augmented with several kinds of information that were hidden in the description of the algorithm. First, we separate concatenations introduced by the algorithm from those already in the original type definitions, writing the former with semicolons. Second, we associate each type with a list of type names that records a history of which type names have been unfolded before the algorithm gets to this type. Formally, a *type sequence* K (i.e., a term to transit) has the form $T_1^{\tau_1}; \dots; T_n^{\tau_n}$, where each type is superscripted with a *name list* of the form $X_1; \dots; X_m$. We write \cdot for the empty type sequence and \emptyset for the empty name list. Intuitively, a type sequence of the form $T_1^{\tau_1}; \dots; T_n^{\tau_n}$ corresponds to the type $(T_1, (\dots, T_n))$. In each name list τ_i , the first name was unfolded most recently. The empty type sequence \cdot corresponds to the empty sequence type. We write $\tau \supseteq \tau'$ when $\tau = \tau''; \tau'$ for some τ'' . Also, we write $X \in \tau$ when X is an element of τ .

We now define the transition relation of the form $K_1 \Rightarrow K_2$, read “if the algorithm encounters the type corresponding to K_1 , then it also encounters the type corresponding to K_2 .” The transition relation is the smallest relation including the following rules:

$$\begin{aligned}
& \emptyset; K \Rightarrow \cdot \\
& ()^\tau; K \Rightarrow K \\
& Y^\tau; K \Rightarrow E(Y)^{(Y; \tau)}; K \\
& 1[T]^\tau; K \Rightarrow T^\emptyset \\
& 1[T]^\tau; K \Rightarrow K \\
& (T_1, T_2)^\tau; K \Rightarrow T_1^\tau; T_2^\tau; K \\
& (T_1 | T_2)^\tau; K \Rightarrow T_1^\tau; K \\
& (T_1 | T_2)^\tau; K \Rightarrow T_2^\tau; K.
\end{aligned}$$

In the first rule (corresponding to the algorithm rules TR-EMP1 and TR-EMP2), if we encounter a sequence beginning with the empty sequence, then we also encounter the sequence obtained by removing the empty sequence. Likewise in the second rule (corresponding to the algorithm rules TR-NM1 and TR-NM2), if we encounter a sequence beginning with a type name, then we also encounter a sequence obtained by unfolding the type name. (Note that we record the

unfolded type name in the type name list.) Further, in the combination of the third and the fourth rules (corresponding to TR-LAB1 and TR-LAB2), if we encounter a sequence beginning with a label, then we also encounter both its content and its remainder. (For the content, we reset the type name list to the empty, since, when we get to the content T , we freshly consider the transitions from this.) The other rules are analogous.

Our goal is then to show that the set of type sequences reachable from the starting type sequence is finite. The key observation in the proof is that the transition function defined as above preserves the invariant that all T_i s in a type sequence $T_1^{r_1}; \dots; T_n^{r_n}$ appear in *different places* in the original type definitions; the type sequences satisfying this invariant are finite in number. To see why this is so, let us consider the above example again. We start with the type sequence Y^θ and transitions go as follows:

$$Y^\theta \Rightarrow (X, b[])^Y \Rightarrow X^Y; b[]^Y \Rightarrow ((a[], X) | ())^{X;Y}; b[]^Y.$$

The last type sequence then branches into two chains of transitions: first

$$\begin{aligned} &\Rightarrow (a[], X)^{X;Y}; b[]^Y \Rightarrow a[]^{X;Y}; X^{X;Y}; b[]^Y \Rightarrow X^{X;Y}; b[]^Y \\ &\Rightarrow ((a[], X) | ())^{X;X;Y}; b[]^Y \end{aligned}$$

(since the last type sequence here is the same as the beginning of the chain, continuing the transition will only yield the same type sequences, modulo differences in type name lists.), and second

$$\Rightarrow ()^{X;Y}; b[]^Y \Rightarrow b[]^Y.$$

Notice that each type sequence appearing in all the above transitions is a sequence of types from different places in the original type definitions.

In order to prove the above-mentioned invariant, we will also need several other invariants. First, a type appearing earlier in any type sequence has an equal or longer type name list than any other type appearing later (since unfolding occurs only in the first type in the sequence). Second, in any type sequence, only the right-most type that has X in its type name list may contain X in tail positions; any other type with X in its type name list cannot contain X at all. Third, in order for induction to work, the condition “appearing different places” has to be slightly strengthened: each type in the type sequence must not be a subphrase of any other type in the sequence “in the same level.”

To formalize these invariants, we make a few notational preparations. First, we assume that a unique id is implicitly added to each type expression appearing in the given type definitions or the type T_0 that we want to translate, and that type expressions referred to by the same metavariable name always have the same id. We define the *same-level subphrase* relation, written \sqsubseteq , as the smallest reflexive and transitive relation including the following:

$$\begin{aligned} T_1 &\sqsubseteq (T_1, T_2) \\ T_2 &\sqsubseteq (T_1, T_2) \\ T_1 &\sqsubseteq (T_1 | T_2) \\ T_2 &\sqsubseteq (T_1 | T_2). \end{aligned}$$

We write Σ_X for the all unguarded (not enclosed by a label) subphrases in the body of X , that is, $\{T \mid T \subseteq E(X)\}$. Also, we write Σ for *all* the subphrases (including guarded ones) appearing in the type definitions E and the type T_0 ; we write Σ_\emptyset for all the guarded subphrases in the type definitions, that is, $\Sigma \setminus \bigcup_{X \in \text{dom}(E)} \Sigma_X$.

All the properties of type sequences that we have discussed can be summarized by the following definition of invariants.

Definition A.2 (Invariants). Let $K = T_1^{\tau_1}; \dots; T_n^{\tau_n}$. We call the following the *invariants* for K .

- (1) $T_i \in \Sigma_\emptyset$ if $\tau_i = \emptyset$, and $T_i \in \Sigma_Y$ if $\tau_i = Y; \tau'_i$.
- (2) $\tau_1 \supseteq \dots \supseteq \tau_n$
- (3) For all X , if $X \in \tau_m$ and $X \notin \tau_{m+1}$ (or $m = n$), then $X \notin \text{nontail}(T_m)$ and $X \notin \text{toplevel}(T_i)$ for all $1 \leq i < m$.
- (4) $T_i \not\sqsubseteq T_j$ for all $i \neq j$.

Here, $\text{nontail}(T)$ is the set of non-tail variables of a type T and $\text{toplevel}(T)$ is the set of top-level variables of a type T . The definitions of these are given in Section 3.1.

The main theorem is that transition preserves the invariants for type sequences.

THEOREM A.3. *If K satisfies the invariants and $K \Rightarrow K'$, then K' satisfies the invariants.*

Since Σ is finite, condition (4) ensures that the set of type sequences that satisfy the invariants is finite. Furthermore, we begin the translation with a singleton sequence T_0^\emptyset , which trivially satisfies the invariants. Therefore the theorem implies that the translation terminates.

PROOF OF THEOREM A.3. Conditions (1) and (2) can easily be checked; the others require a little more work. Let us begin with condition (3).

Case: $()^\tau; K \Rightarrow K$
or $1[T]^\tau; K \Rightarrow K$

Condition (3) for the lhs immediately implies condition (3) for the rhs.

Case: $1[T]^\tau; K \Rightarrow T^\emptyset$

The result holds trivially, since there is no X that satisfies the condition.

Case: $Y^\tau; K \Rightarrow E(Y)^{(Y;\tau)}; K$

Let $\tau_0 = Y; \tau$ and $K = U_1^{\tau_1}; \dots; U_n^{\tau_n}$. From condition (2), we have two cases on X .

Subcase: $X \in \tau_0 \quad X \notin \tau_1$
or $n = 0$

If $X \neq Y$, then we have $X \notin \text{nontail}(Y)$ from condition (3) for the left-hand side. Therefore, $X \notin \text{nontail}(E(Y))$ by the definition of *nontail*. If $X = Y$, then, from well-formedness of E , we immediately obtain $Y \notin \text{nontail}(E(Y))$.

Subcase: $X \in \tau_m \quad X \notin \tau_{m+1} \quad 1 \leq m \leq n$
or $n = m$

If $X \neq Y$, then we have $X \notin \text{toplevel}(Y)$ from condition (3) for the left-hand side. Therefore, $X \notin \text{toplevel}(E(Y))$ by the definition of *toplevel*. If $X = Y$, then condition (3) requires $Y \notin \text{toplevel}(Y)$. But this is impossible from the definition of *toplevel*.

Case: $(T_1, T_2)^\tau; K \Rightarrow T_1^\tau; T_2^\tau; K$

Let $K = U_1^{\tau_1}; \dots; U_n^{\tau_n}$. From condition (2), we have two cases on X .

Subcase: $X \in \tau_0 \quad X \notin \tau_1$
or $n = 0$

Since $X \notin \text{nontail}((T_1, T_2))$ from condition (3) for the left-hand side, $X \notin \text{toplevel}(T_1)$ and $X \notin \text{nontail}(T_2)$ follow by the definitions of *nontail* and *toplevel*.

Subcase: $X \in \tau_m \quad X \notin \tau_{m+1} \quad 1 < m < n$
or $n = m$

Conditions $X \notin \text{nontail}(U_m)$ and $X \notin \text{toplevel}(U_i)$ for $1 < i < m$ immediately hold. Since $X \notin \text{toplevel}((T_1, T_2))$ from condition (3) for the left-hand side, conditions $X \notin \text{toplevel}(T_1)$ and $X \notin \text{toplevel}(T_2)$ follow from the definition of *toplevel*.

Case: $(T_1 | T_2)^\tau; K \Rightarrow T_1^\tau; K$
or $(T_1 | T_2)^\tau; K \Rightarrow T_2^\tau; K$

Similar to the previous case.

Finally, we prove condition (4). All the cases are obvious except for

$$Y^\tau; K \Rightarrow E(Y)^{(Y;\tau)}; K.$$

Let $K = U_1^{\tau_1}; \dots; U_n^{\tau_n}$. The case $Y \notin \tau_i$ with $i \geq 1$ never happens, because otherwise, condition (3) would require $Y \notin \text{toplevel}(Y)$, which is impossible from the definition of *toplevel*. Therefore, the result follows from condition (1). \square

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