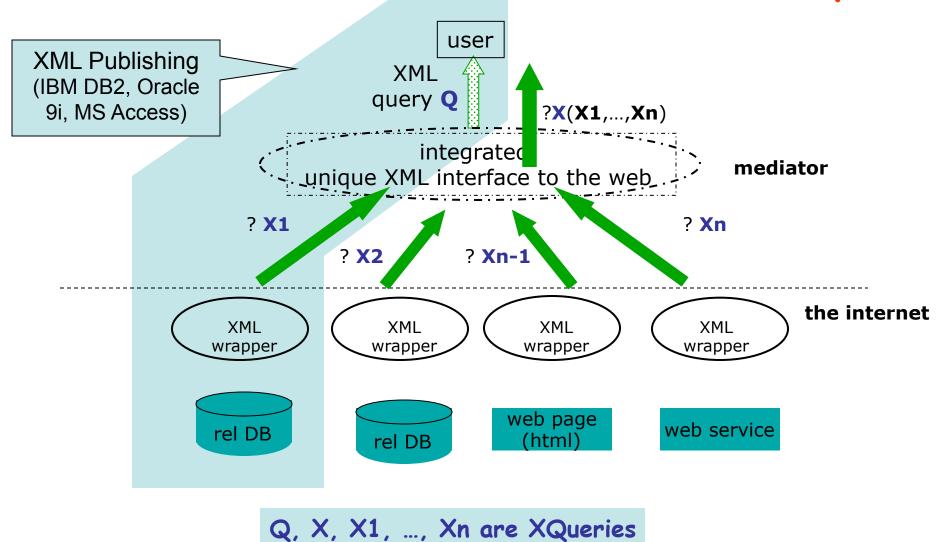
# XQuery Advanced Topics

Alin Deutsch

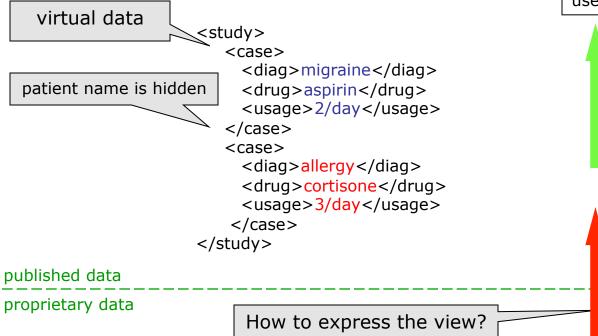
### Roadmap

- Use of XQuery for Web Data Integration
- XQuery Evaluation Models
- Optimization
- Flavor of Standardization Issues
  - Equality in XQuery
- More on Optimization

#### The Web as Database Queried in XQuery



A Simple Publishing Scenario



#### prescription

usage	drug	name
2/day	aspirin	John
3/day	cortisone	Jane

#### patient

name	diagnosis	
John	migraine	
Jane	allergy	

How to "compose" the user query with the view, obtaining the reformulation?

reformulation

(SQL)

user query

(XQuery)

carrespondence

is called **view** 

## Encoding relational data as XML

Want to specify view from proprietary  $\rightarrow$  published data as XML  $\rightarrow$  XML view expressed in XQuery

#### prescription

usage	drug	name
2/day	aspirin	John
3/day	cortisone	Jane

#### patient

name	diagnosis	
John	migraine	
Jane	allergy	

# Proprietary > Published View: XML > XML

public.xml

```
<study>
          <case><diag>migraine</diag><drug>aspirin</drug>
                 <usage>2/day</usage>
          </case>
          <case><diag>allergy</diag><drug>cortisone</drug>
                 <usage>3/day</usage>
           </case>
       </study>
                                                  expressible as XQuery
published data
proprietary data
        cription>
           <tuple><usage>2/day</usage>
                  <drug>aspirin</drug><name>John</name>
           </tuple>
           <tuple><usage>3/day</usage>
                  <drug>cortisone</drug><name>Jane</name>
            </tuple>
        </prescription>
```

encoding.xml

#### The View

```
<study>
         $t1 in document("encoding.xml")//patient/tuple,
 for
             $n1 in $t1/name/text(),
             $di in $t1/diagnosis/text(),
         $t2 in document("encoding.xml")//prescription/tuple,
             $n2 in $t2/name/text(),
             $dr in $t2/drug/text(),
             $u in $t2/usage/text(),
 where
         $n1=$n2
 return
         <case><diag>$di</diag>
                <drug>$dr</drug>
                <usage>$u</usage>
         <case>
</study>
```

### A Client Query

Find high-maintenance illnesses (require drug usage thrice a day):

Not directly executable, public.xml does not exist

## The Reformulated Query

Directly executable, expressed in SQL against the proprietary database:

**Select** pr.drug

**From** patient pa, prescription pr

**Where** pa.name = pr.name and

pr.usage = "3/day"

#### prescription

usage	drug	name
2/day	aspirin	John
3/day	cortisone	Jane

#### patient

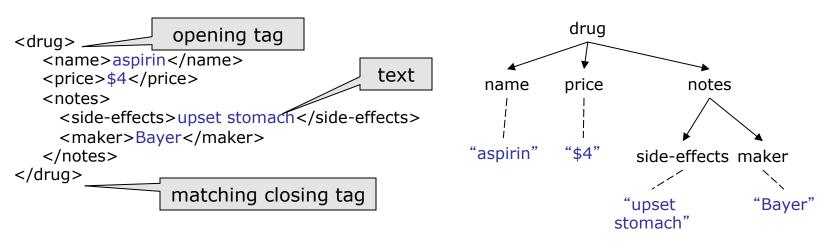
name	diagnosis
John	migraine
Jane	allergy

### Roadmap

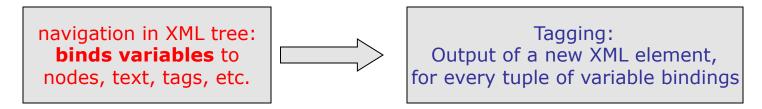
- Use of XQuery for Web Data Integration
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## XQuery Semantics: Navigation & Tagging

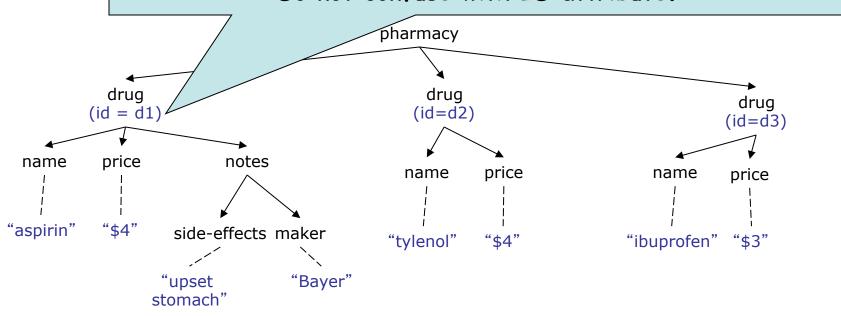
XML data model is a tagged tree



#### XQueries compute in two stages:

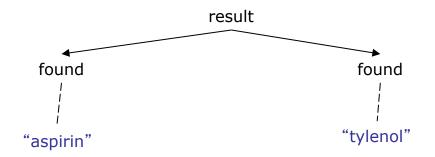


# Node identity, for example java reference of DOM node. Do not confuse with ID attribute.



<b>\$</b> x	\$n	\$p
d1	"aspirin"	<b>"</b> \$4"
d2	"tylenol"	<b>"\$4</b> "
d3	ibu"	<b>-"\$3</b> "

# XQuery Semantics: Tagging

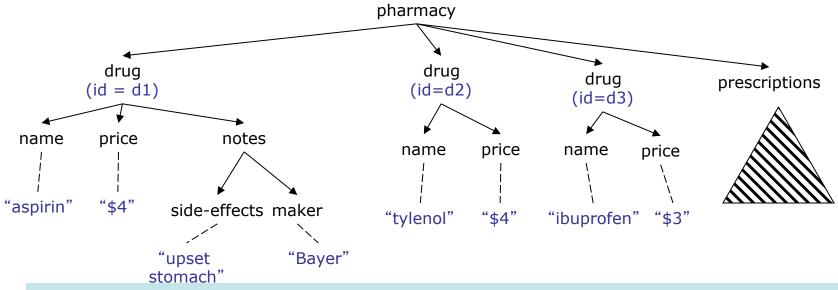


## Descendant Navigation

Direct implementation of descendant navigation is wasteful:

for \$x in \$d//drug

Go to all descendants of the root (all elements), keep <drug>-tagged ones

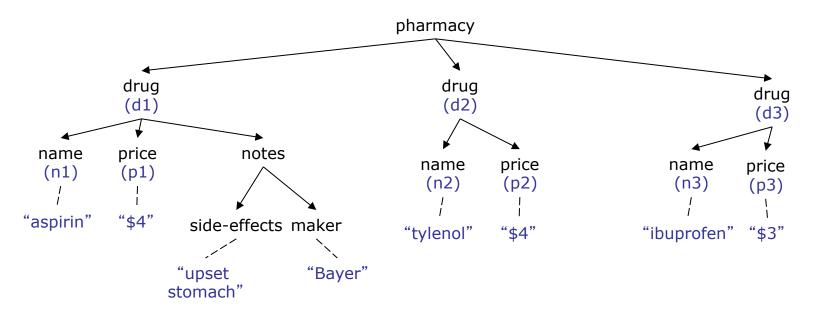


To find the 3 drug elements, a direct implementation visits all elements in the document (e.g. drug). The full query does so repeatedly. In general, a query with n descendant steps may visit |doc size|^n elements!

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  - Stream-based
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#### Index-based Evaluation



Idea 1: keep an index (associative array, hash table) associating tags with lists of node ids. Allows random access into XML tree.

idx: tag	node ids	lookup operation: idx[price] = [p1,p2,p3]
drug	d1,d2,d3	
name	n1,n2,n3	
price	p1,p2,p3	

#### Index-based Evaluation (2)

```
lookup operation: idx[price] = [p1,p2,p3]
idx: tag node ids
    drug
           d1,d2,d3
           n1,n2,n3
    name
           p1,p2,p3
    price
foreach $p in idx[price]
                                                  // p1, p2, p3
    if $p/text() = "$4"
                                                   // p1, p2
      foreach $x in idx[drug]
                                                  // d1, d2, d3
          if $p descendant_of $x
                                                  // p1 of d1, p2 of d2
                                                  // n1, n2, n3
            foreach $n in idx[name]
                if $n descendant_of $x
                                                  // n1 of d1, n2 of d2
                return <found>$n</found>
```

Only 9 elements visited, regardless of size of irrelevant XML subtrees.

But doesn't the implementation of descendant\_of require more visiting?

### Ancestor-Descendant Testing in O(1)

Idea 2: identify each node n by a pair of integers pre(n), post(n), with

pre(n) = the rank of n in the preorder traversal of the tree
post(n) = the rank of n in the postorder traversal

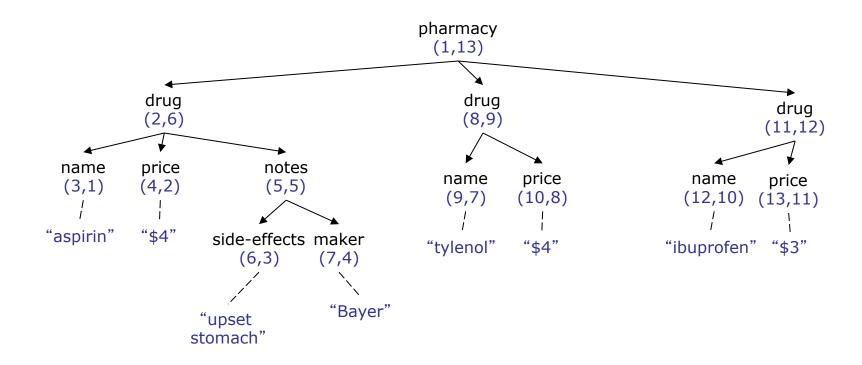
Then

d is descendant of a



pre(d) >= pre(a) and post(d) <= post(a)</pre>

### Example post-preorder node ids



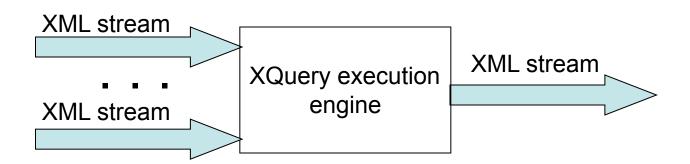
Additional advantage: node identity independent of particular in-memory representation of DOM objects.

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### Stream-based XQuery Execution

- So far, we assumed construction of DOM tree in memory.
- XML documents can be XML representations of databases. The DOM approach does not scale to typical database sizes.
- We want an execution model that minimizes the memory footprint of the XQuery engine.



#### Applications of Stream-based Execution

- Besides scaling to database sizes. There are applications where the data is inherently received in streamed form:
- Sensor networks (attend faculty candidate Sam Madden's talk)
- Network monitoring/XML packet routing
- XML document publish/subscribe systems

### Stream-based XML Parsing

- A parser generates a stream of predefined events (according to the standard SAX API)
- Applications consume these events.
- Each event triggers a handler. The application is coded by providing the code for the handlers.

```
XML input to parser
                                  stream of events output by parser
                                  open("a")
<a>
                                  open("b")
 <b>
                                  open("c")
  <c>
   someText
                                  text("someText")
                                  close("c")
  </c>
                                  close("b")
 </b>
                                  open("d")
 <b
                                  text("moreText")
 moreText
                                  close("d")
 </d>
                                  close("a")
</a>
```

A free SAX parser: http://xml.apache.org/xerces-j/

### Stream-Based XQuery Navigation

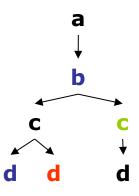
Idea: turn path expressions into Finite Automata over alphabet containing the set of element tags

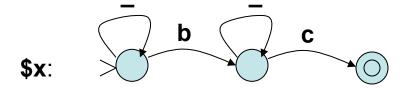
**E.g.** for \$x in //b//c, \$y in \$x/d

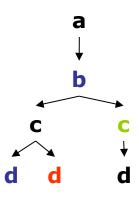
compiles to

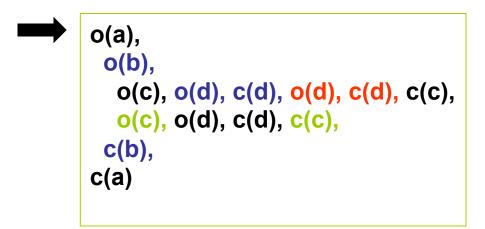


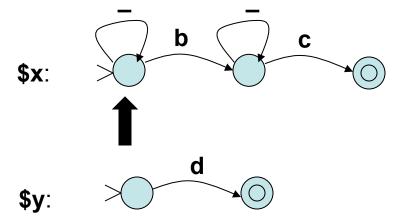
Only one automaton active at any moment. Automaton of y is active only as long as that of x is in final state

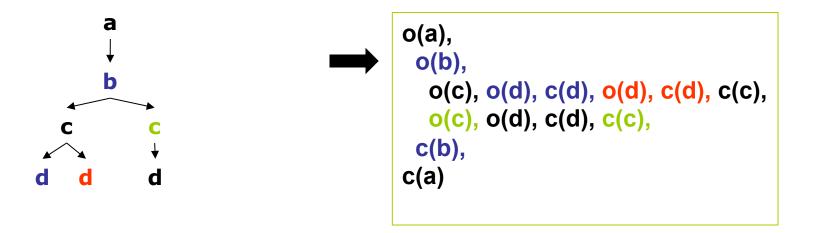


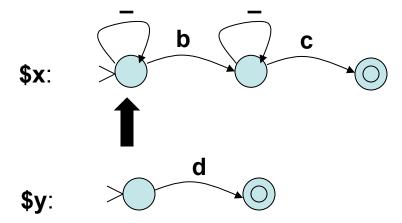


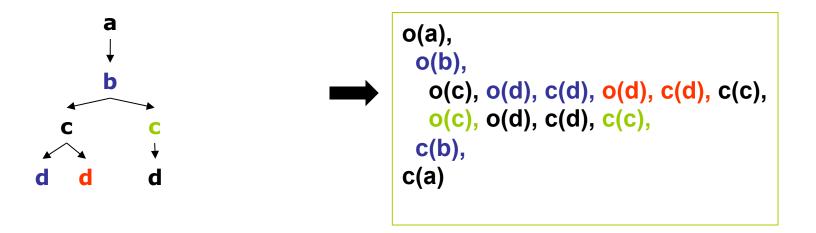


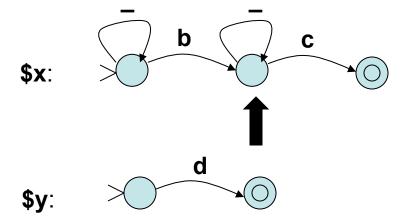


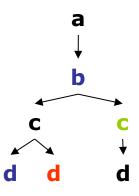


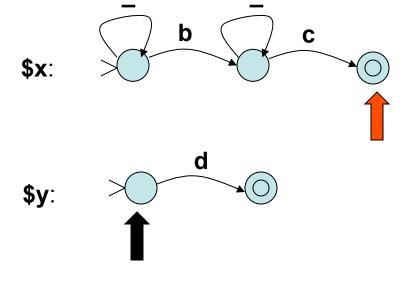


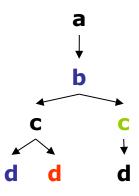




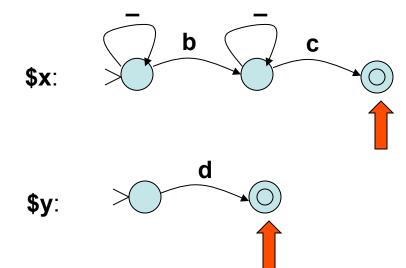




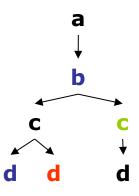


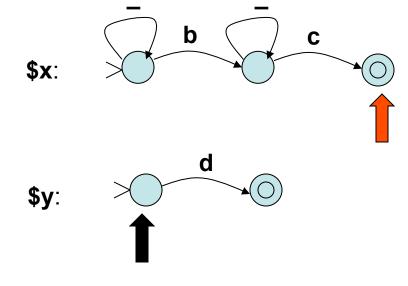


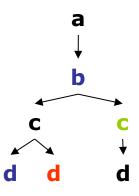
for \$x in //b//c, \$y in \$x/d

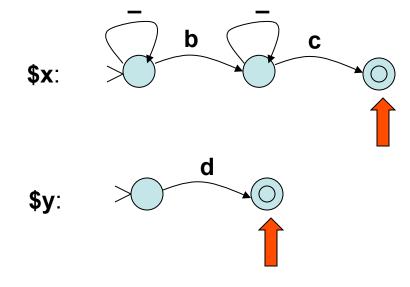


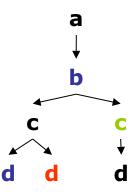
Need to reset automaton for \$y





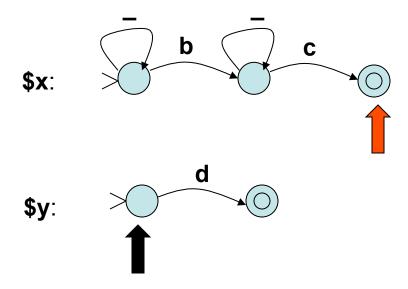


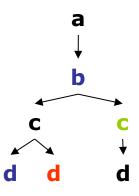


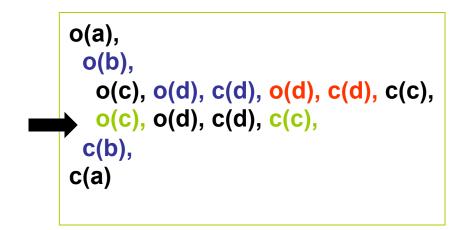


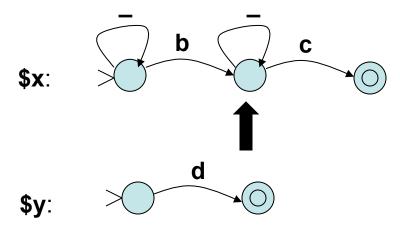
for \$x in //b//c, \$y in \$x/d

Need to reset automaton for \$x to state prior to reading black c element









#### Automaton Extended with Stack

Let d be the transition function of automaton A. The corresponding extension of A with a stack is defined as follows:

current state	current event in stream	stack action	next state
Q	open(tag)	push(Q)	d(Q)
Q	close(tag)	Q' =pop()	Q'

Convince yourselves that the run of this automaton on the stream in the example corresponds to the intended sequence of states.

An additional use of PDAs, aside from parsing.

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#### Semantic Optimization

- Sometimes, we can translate away descendant computation.
- Consider the following DTD describing the structure of drug.xml

```
<!ELEMENT pharmacy (drug*)>
<!ELEMENT drug (name,price,notes?)>
```

Then for all documents satisfying DTD:

```
for $x in $d//drug, $n in $x//name/text() is equivalent to for $x in $d/drug, $n in $x/name/text()
```

## Semantic Optimization As Typechecking

For all XML documents conforming to the DTD

```
<!ELEMENT pharmacy (drug*)>
```

<!ELEMENT drug (name,price,notes?)>

we can determine statically that

for \$x in \$d//drug, \$m in \$d/maker

returns the empty answer.

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## Element Equality in XQuery

- Two kinds of equality:
  - "==" id-based (an element node is equal only to itself)
  - "=" value-based
- Value-based equality underwent several drafts,
- Initially (about one year into standardization process): text-centric point of view. XML elements are value-equal iff their text values are equal after stripping away the XML annotations.
  - E.g. <a><b>f</b><c>oo</c></a> = <m>foo</m>
- · Currently: XML elements are equal iff their corresponding trees are isomorphic

### Id-based Element Equality

Let \$x be bound to an XML tree. Then

creates a new XML tree (fresh node ids) and it is short for

#### Always true:

$$(\$x)/a/* = $x$$

(value-based equality)

#### **Always false:**

(id-based equality)

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### More on XQuery Optimization

- There are many ways to write the same query (i.e. there are many distinct XQuery expressions with identical semantics)
- Some of these expressions lead to cheaper execution than their counterparts.
- Goal of query optimization: given a query Q, find the optimal query Q' with identical semantics (we say that Q and Q' are equivalent)
- · Basic test in query optimization: checking query equivalence
- · The more expressive a language, the harder it is to test equivalence
- · Various classes of XQueries have distinct complexity:
  - PTIME (1), NP-complete (1),  $\Pi_2^p$ -complete (4), PSPACE-complete (1), EXPTIME-complete, undecidable

#### The UCSD Database Lab

- · Main Focus: XML Query Optimization
- · Check out the weekly DB Research Meeting
- Faculty
  - Victor Vianu
  - Yannis Papakonstantinou
  - Alin Deutsch
- San Diego SuperComputer Resaerchers
  - Ilkay Altintas
  - Amarnath Gupta

www.db.ucsd.edu