# XQuery

CS 445

Fall 2008

# Querying XML Data

- XPath = simple navigation through the tree
- XQuery = the SQL of XML

# Query Language and Data Model

- A query language is "closed" w.r.t. its data model if input and output of a query conform to the model
- SQL
  - Set of tuples in, set of tuples out
- XPath 1.0
  - A tree of nodes (well-formed XML) in, a node set out.
- XQuery 1.0
  - Sequence of items in, sequence of items out
- Compositionality of a language
  - Output of Query 1 can be used as input to Query 2

### **XQuery Values**

- Item = node or atomic value.
- Value = ordered sequence of zero or more items.
- Examples:
  - () = empty sequence.
  - ("Hello", "World")
  - ("Hello", <PRICE>2.50</PRICE>, 10)

### Sample Data for Queries

```
<bib>
   <book> <publisher> Addison-Wesley </publisher>
           <author> Serge Abiteboul </author>
           <author> <first-name> Rick </first-name>
                     <|ast-name> Hull </|ast-name>
           </author>
           <author> Victor Vianu </author>
           <title> Foundations of Databases </title>
           <year> 1995 </year>
   </book>
   <book price="55">
           <publisher> Freeman </publisher>
           <author> Jeffrey D. Ullman </author>
           <title> Principles of Database and Knowledge Base Systems </title>
           <year> 1998 
   </book>
</bib>
```

### **Document Nodes**

- Form:
  - doc("<file name>").
- Establishes a document to which a query applies.
- Example:
  - doc("/courses/445/bib.xml")

### FLWOR expressions

- FLOWR is a high-level construct that
  - supports iteration and binding of variables to intermediate results
  - is useful for joins and restructuring data
- Syntax: For-Let-Where-Order by-Return

```
for $x in expression1  /* similar to FROM in SQL */

[let $y := expression2]  /* no analogy in SQL */

[where expression3]  /* similar to WHERE in SQL */

[order by expression4 (ascendingIdescending)?]

/* similar to ORDER-BY in SQL */

return expression4  /* similar to SELECT in SQL */
```

# Example FLOWR Expression

- The for clause iterates over all books in an input document, binding \$x to each book in turn.
- For each binding of \$x, the let clause binds \$y to all authors of this book.
- The result of for and let clauses is a tuple stream in which each tuple contains a pair of bindings for \$x and \$y, i.e. (\$x, \$y).
- The where clause filters each tuple (\$x, \$y) by checking predicates.
- The order by clause orders surviving tuples.
- The return clause returns the count of \$y for each surviving tuple.

Find all book titles published after 1995:

```
for $x in doc("bib.xml")/bib/book
```

where x/year/text() > 1995

return \$x/title

#### Result:

Equivalently (perhaps more geekish)

for \$x in doc("bib.xml")/bib/book[year/text() > 1995] /title
return \$x

And even shorter:

doc("bib.xml")/bib/book[year/text() > 1995] /title

• Find all book titles and the year when they were published:

We can construct whatever XML results we want!

#### Answer

```
<answer>
   <what> How to cook a Turkey </what>
   <when> 2003 </when>
</answer>
<answer>
   <what> Cooking While Watching TV </what>
   <when> 2004 </when>
</answer>
<answer>
   <what> Turkeys on TV</what>
   <when> 2002 </when>
</answer>
```

- Notice the use of "{" and "}"
- What is the result without them?

### More Examples of WHERE

#### Selections

### Aggregates

Find all books with more than 3 authors:

```
for $x in doc("bib.xml")/bib/book
where count($x/author)>3
return $x
```

```
count = a function that counts
avg = computes the average
sum = computes the sum
distinct-values = eliminates duplicates
```

### Aggregates

#### Same thing:

```
for $x in doc("bib.xml")/bib/book[count(author)>3]
RETURN $x
```

### FOR v.s. LET

#### **FOR**

• Binds *node variables*  $\rightarrow$  iteration

#### LET

• Binds *collection variables*  $\rightarrow$  one value

### FOR v.s. LET

```
for $x in /bib/book
return <result> { $x } </result>
```

```
let $x := /bib/book
return <result> { $x } </result>
```

#### Returns:

```
<result> <book>...</book></result> <result> <book>...</book></result> <result> <book>...</book></result> ...
```

#### Returns:

```
<result> <book>...</book> <book>...</book> <book>...</book> ...</book> ...</result>
```

• "Flatten" the authors, i.e. return a list of (author, title) pairs

```
Answer:

<answer>
  <title> abc </title>
  <author> efg </author>
  </answer>
  <answer>
  <title> abc </title>
  <author> hkj </author>
  </answer>
  </answer>
```

# XQuery: Nesting

For each author of a book by Morgan Kaufmann, list all books she published:

In the <u>RETURN</u> clause comma concatenates XML fragments

### XQuery

#### Result:

# Getting Distinct Values from FOR

- Distinct values: the fn:distinct-values function eliminates duplicates in a sequence by value
  - The <u>for expression</u> evaluates to a sequence of nodes
  - fn:distinct-values converts it to a sequence of atomic values and removes duplicates

### Value Comparison

```
<author>
<first>Jeffery</first>
<last>Ullman</last>
</author>
```

- Value comparison "eq": compares single values
- "eq" applies atomization (fn:data()) to each operand
  - Given a sequence of nodes, fn:data() returns an atomic value for each node which consists of:
    - a <u>string value</u>, i.e., the concatenation of the string values of *all*\_its *Text Node descendants* in *document order*
    - a type, e.g., xdt:untypedAtomic
  - For each operand, "eq" uses the fn:data() result if it evaluates to a singleton sequence, o.w. runtime error.
- for \$a in doc("bib.xml")/bib/book/author
  where \$a eq "JefferyUllman"
  return \$a/..
- for \$b in doc("bib.xml")/bib/book where \$b/author eq "JefferyUllman" return \$b/author

# General Comparison

- General comparison operators (=, !=, <, >, <=, >=):
   existentially quantified comparisons, applied to operand
   sequences of any length
- Atomization (fn:data()) is applied to each operand to get a sequence of atomic values
- Comparison is true if one value from a sequence satisfies the comparison

```
for $b in doc("bib.xml")/bib/book
where $b/author = "JefferyUllman"
return $b/author
```

# String Operations

Functions for string matching

```
fn:contains(xs:string, xs:string)
fn:starts(ends)-with(xs:string, xs:string
fn:substring-before(after)(xs:string, xs:string)
...
```

 Again, atomization (fn:data()) is applied to each function parameter to get an atomic value.

```
for $a in doc("bib.xml")//author
where contains($a, "Ullman")
return $a
```

```
<author>
<name>Jeffery Ullman</name>
</author>
```

```
<author>
<first>Jeffery</first>
<last>Ullman</last>
</author>
```

### **Element Construction**