



## Advanced databases and data models:

### Theme2: Query languages

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### Today's lecture

Different kinds of query languages

Languages

Lorel

XPath

XQuery

SPARQL

Transformation of data

XSL

FOR XML

Data guides – why and what



### Semi-structured data - properties

Data model/data guide could be supportive or a hinder while querying

Data model/guide changes commonly

Object can change type/class

The distinction between data and schema is blurred

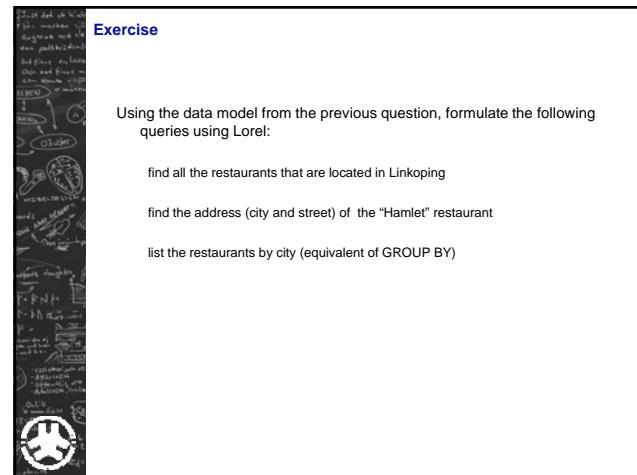
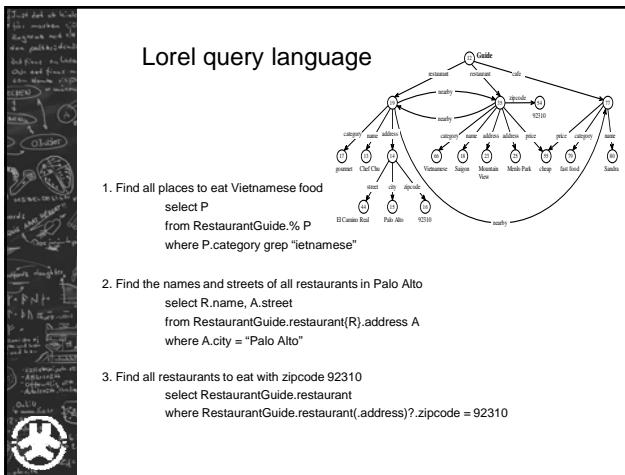
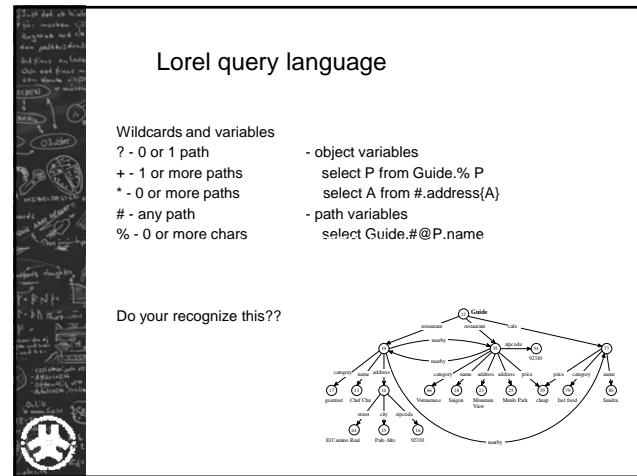
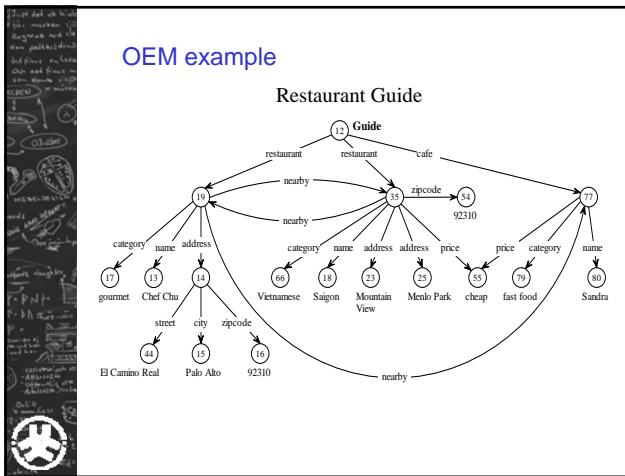
### Types of query languages

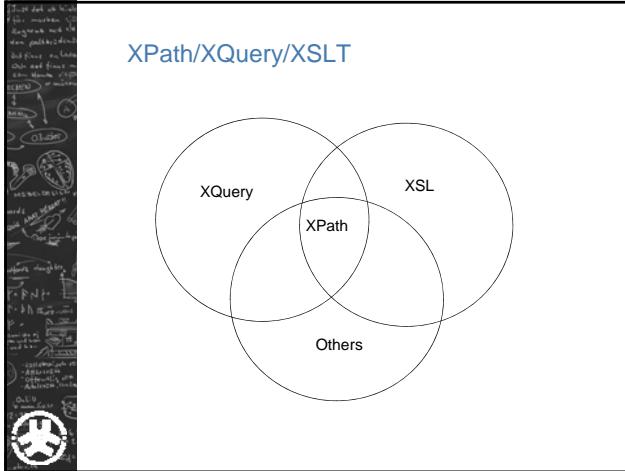
Typically:

Data	Unstructured	Semi-structured	Structured
Example	Text	XML, RDF, OEM	Relational database
Typical query language	Keywords	XPath, XQuery, Lorel ...	SQL, ....
Typical user	Every day		Expert

However, this is changing... (more in coming lectures)







**XPath**

**Objects:**

- Nodes
- Atomicvalues
- Items

**Relations:**

- Parent
- Children
- Ancestor
- Descendants

```

<minimodel name="sugartransport" >
  <listOfCompartments>
    <compartment id="blood" name="inblood" />
    <compartment id="cell" name="musclecell" />
  </listOfCompartments>
  <listOfSpecies>
    <species id="sug1" name="sugarinblood" compartment="blood" />
    <species compartment="ins" id="insulin" name="blood" />
    <species compartments="sug2" id="sugarcell" name="cell" />
    <species compartment="en" id="energy" name="cell" />
  </listOfSpecies>
  <listOfReactions>
    <reaction id="moveall" name="sugarcodil" >
      <listOfReactants>
        <speciesReference species="sug1"/>
        <speciesReference species="ins"/>
      </listOfReactants>
      <listOfProducts>
        <speciesReference species="sug2"/>
      </listOfProducts>
    </reaction>
    <reaction id="move" name="makemovement" >
      <listOfReactants>
        <speciesReference species="sug2"/>
      </listOfReactants>
      <listOfProducts>
        <speciesReference species="en"/>
      </listOfProducts>
    </reaction>
  </listOfReactions>
</minimodel>
  
```

**XPath**

**Node selection:**

```

<minimodel name="sugartransport" >
  <listOfCompartments>
    <compartment id="blood" name="inblood" />
    <compartment id="cell" name="musclecell" />
  </listOfCompartments>
  <listOfSpecies>
    <species id="sug1" name="sugarinblood" compartment="blood" />
    <species compartment="ins" id="insulin" name="blood" />
    <species compartments="sug2" id="sugarcell" name="cell" />
    <species compartment="en" id="energy" name="cell" />
  </listOfSpecies>
  <listOfReactions>
    <reaction id="moveall" name="sugarcodil" >
      <listOfReactants>
        <speciesReference species="sug1"/>
        <speciesReference species="ins"/>
      </listOfReactants>
      <listOfProducts>
        <speciesReference species="sug2"/>
      </listOfProducts>
    </reaction>
    <reaction id="move" name="makemovement" >
      <listOfReactants>
        <speciesReference species="sug2"/>
      </listOfReactants>
      <listOfProducts>
        <speciesReference species="en"/>
      </listOfProducts>
    </reaction>
  </listOfReactions>
</minimodel>
  
```

**Predicates:**

```

</species[@id='ins']>
  
```

**Wildcards:**

```

/* 
  /reaction/@*
  
```

**Several paths:**

```

</species>/<compartment>
  
```

And a lot of functions for selecting ancestors etc.

```

data(doc('minimodel.xml')/compartment[@id='blood']/@name)
  
```

**XQuery**

**XML Query language**

```

for $x in doc('minimodel.xml')//species
  where data($x/@compartment)="blood"
  return $x
  
```



## XPath vs. XQuery

When do you need XQuery?



## XQuery allows many variations

```
for $x in doc('minimodel.xml')//species,  
      $y in doc('minimodel.xml')//compartment  
    where data($x/@compartment)=data($y/@id)  
return  
  <pair> {data($x/@name)} is inside {data($y/@name)} </pair>  
  
for $x in doc('minimodel.xml')//species,  
      $y in doc('minimodel.xml')//compartment  
    [@id=data($x/@compartment)]  
return  
  <pair> {data($x/@name)} is inside {data($y/@name)} </pair>
```



## XQuery functions

Declare local functions  
Can be recursive!

```
declare function local:fak($n)  
{if ($n=1)  
  then 1  
  else ($n * local:fak($n - 1))  
};  
<res>{local:fak(3)}</res>
```



## XQuery functions

```
declare function local:next($r)  
{  
  <next>  
  {data($r/listOfProducts/speciesReference/@species)}  
  for $x in $r/listOfProducts/speciesReference,  
        $y in doc('minimodel.xml')  
          //listOfReactants/speciesReference  
          [@species=data($x/@species)]  
  return local:next($y/..)}  
  </next>}  
for $x in doc('minimodel.xml')//species,  
      $y in doc('minimodel.xml')  
        //listOfReactants/speciesReference  
        [@species=data($x/@id)]  
return <answer> {data($x/@id)} leads to  
  {(local:next($y/..))} </answer>
```



## XQuery - tips

- The Xpath expressions can return many answers.
- Attribute values – use data()
- Start from something simple – build on it.



## SPARQL – Query language for RDF

```
PREFIX books: <http://example.org/book/>
PREFIX dc: <http://purl.org/dc/elements/1.1/>
SELECT ?book ?title
WHERE { ?book dc:title ?title }
```

```
PREFIX dc: <http://purl.org/dc/elements/1.1/>
CONSTRUCT { $book dc:title $title }
WHERE { $book dc:title $title }
```



## Transformation: XSL

```
<catalog>
<cd>
<title>Empire Burlesque</title>
<artist>Bob Dylan</artist>
<country>USA</country>
<company>Columbia</company>
<price>10.90</price>
<year>1985</year>
</cd>
...
<xsl:stylesheet version="1.0" encoding="ISO-8859-1">
<xsl:template match="/">
<html>
<body>
<h2>My CD Collection</h2>
<table border="1">
<tr bgcolor="#9acd32">
<th>Title</th>
<th>Artist</th>
</tr>
<xsl:for-each select="catalog/cd">
<xsl:for-each>
<td><xsl:value-of select="title" /></td>
<td><xsl:value-of select="artist" /></td>
</xsl:for-each>
</xsl:for-each>
</table>
</body>
</html>
</xsl:template>
</xsl:stylesheet>
```

Transform XML to  
another format



## Transformation: Relations -> XML

```
select xmllag(xmlelement("speciesReference",
xmlelements(reactants.reactant as species))
from reactants where reaction.id=reactants.id)
```

```
<speciesReference species="insulin">
```

### Transformation: Relations -> XML

```
select xmlelement("reaction", xmlattributes(reaction.reaction as name, reaction.id),
    xmlelement("listOfReactants",
        (select xmlagg(xmlelement("speciesReference",
            xmlattributes(reactants.reactant as species)
        from reactants where reaction.id=reactants.id))
        xmlelement("listOfProducts",
            (select xmlagg(xmlelement("speciesReference",
                xmlattributes(products.product as species))
            from products where reaction.id=products.id))
        from reaction
```

### Transformation: Relations -> XML

```
select speciesReference.reactant as species
from reactants speciesReference
where reaction.id = speciesReference.id
for XML auto, type
```

```
<speciesReference species="insulin">
```

### Transformation: Relations -> XML

```
select reaction.reaction as name, reaction.id,
    (select (select speciesReference.reactant as species from
        reactants speciesReference where reaction.id =
        speciesReference.id
        for XML auto, type)
    from emptyXML as listOfReactants for XML auto, type),
    (select (select speciesReference.product as species from
        products speciesReference where reaction.id =
        speciesReference.id
        for XML auto, type)
    from emptyXML as listOfProducts for XML auto, type) ,
    from reaction
    for XML auto, type
```

### Lab exercises:

Construct queries for your relational and XML data.

Record what is hard or easy.

Answer questions, compare and write report.

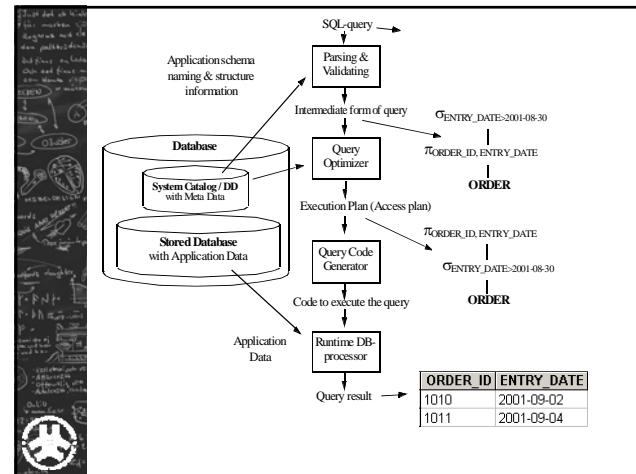
For +: XSLT and XML generation for XML

**Types of query languages**

Typically:

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Typical user	Every day		Expert

For efficient query processing,  
the data structure is important.



**Heuristic optimisation**

Idéa: Do selection and projection first, join as late as possible

Algorithm:

- Break up conjunctive select into cascades
- Move down select as far as possible in the tree
- Rearrange select operations – most restrictive first
- Convert cross product to join with the appropriate join condition from a selection
- Move down project operations as far as possible in the tree
- Identify subtrees that can be executed by a single algorithm

**SQL**

Incompleteness – NULL – Inner and outer join.

Name	Phone	Address
Ludvig	12345	NULL
Lena	23456	NULL

## Incompleteness of XQuery

- (a) node ids (they can be replaced by node variables);
- (b) node labels (they can be replaced by wildcards);
- (c) precise vertical relationship between nodes (we can use descendant edges in addition to child edges);
- (d) precise horizontal relationship between nodes (using younger-sibling edges instead of next-sibling).

## Data Guides

A structural summary over a databank that is used as a dynamic schema

Is used in query formulation and optimization

Is often created a posteriori

Properties:

- concise
- accurate
- convenient

## Data Guides - definitions

Label path: sequence of labels  $L_1.L_2. \dots .L_n$

Data path: alternating sequence of labels and oids:  $L_1.o_1.L_2.o_2. \dots .L_n.o_n$

Data path  $d$  is an instance of label path  $l$  if the sequences of labels are identical in  $l$  and  $d$ .

## Data Guides - definition

A data guide for object  $s$  is an object  $d$  such that every label path of  $s$  has exactly one data path instance in  $d$ , and each label path in  $d$  is a label path of  $s$ .



## Data Guides

A databank can have several data guides

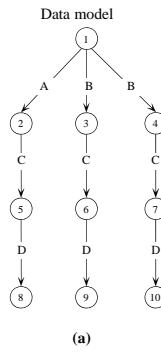
### Minimal data guides

the smallest data guides

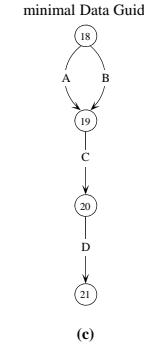


## Data Guides - example

Data model



minimal Data Guide



## Minimal Data Guides

Concise

May be hard to maintain

Example: child node for 10 with label E



## Strong Data Guides

Intuitively:

"label paths that reach the same set of objects in the data model = label paths that reach the same objects in the data guide"



## Strong Data Guides - definitions

An object  $o$  can be reached from  $s$  via  $l$  if there is a data path of  $s$  that is an instance of  $l$  and that has  $o$  as last oid  
 $(L_1.o_1.L_2.o_2. \dots L_n.o)$

The target set for label path  $l$  in object  $s$  is the set of objects that can be reached from  $s$  via  $l$ . Notation:  
 $T(s,l)$

$L(s,l)$ : set of label paths of  $s$  that have the same target set in  $s$  as  $l$ .

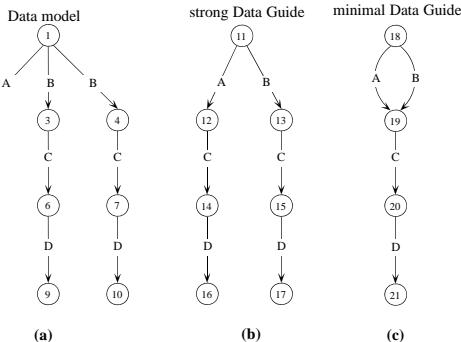
## Strong Data Guides - definitions

### Definition:

$d$  is a strong data guide for  $s$  if  
for all label paths  $l$  of  $s$  it holds that  $L(s,l) = L(d,l)$

There is a 1-1-mapping between target sets in the data model and nodes in a strong data guide.

## Data Guides - example



## Strong Data Guides - algorithm

### Implementation:

Traverse data model depth-first.

Each time you find a new target set for label path  $l$ , create a new object in the data guide.

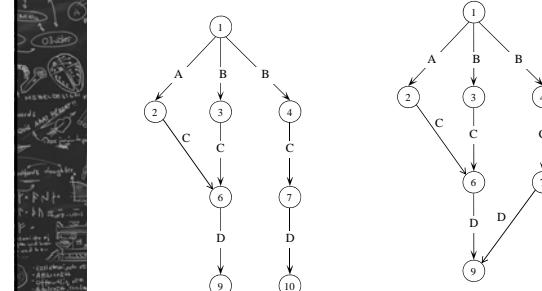
If the target set is already represented in the data guide, do not create a new object, but link to the existing object.

## Strong Data Guides - use

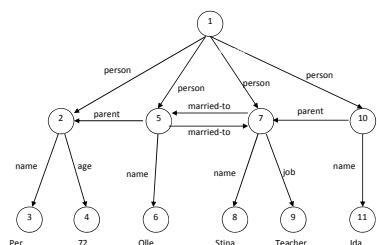
Easier to maintain

Used as path index for query optimization

## Construct the strong data guide



## Exercise: Construct the minimal and strong data guides.



## Exercise

Draw the strong Data Guide for the restaurant guide data model below.

