

Lab 2.3

Auto-generating Java classes using JAXB

Outline

- Perform a blast search using your favorite protein, and collect the results in XML format. Read the outline for the DOCTYPE fix.
- Use the XML Schema provided to auto-generate the Java classes needed by the JAXB runtime environment.
- Create a class which:
 - Initializes the JAXB runtime environment with your content tree.
 - Unmarshalls your blast results from file to the content tree.
 - Validates the content tree (unmarshall-time).
 - Outputs a summary of the results to console.

Getting the source code

- Use CVS!
- Module name is “blastoutput”

Making eclipse happy (optional – you can use command line)

- You may have conflicting versions of xerces.
- This can be overridden by overriding the ANT classpath
- Under the Window->preferences menu, expand the Ant item, and choose Runtime.
- Add the following External JARs to the classpath:
 - /opt/diro/jwsdp-1.3/jaxp/lib/endorsed/xercesImpl.jar
 - /opt/diro/jwsdp-1.3/jaxp/lib/endorsed/dom.jar
- Create a classpath variable called JWSDP_HOME and point it to /opt/diro/jwsdp-1.3

Where did this XML Schema come from?

- NCBI makes the Document Type Definition (DTD) for BlastOutput available through their website
 - <http://www.ncbi.nih.gov/dtd>
- Converted to XML Schema using Datatool program (part of the NCBI C++ toolkit)
 - Note: Autogeneration
- NOTE: BioJava Blast Parser will be introduced in lecture 3.1

Reality Check

- XML Schema specification is not fully mature.
- The decimal datatype cannot handle scientific notation (Example: 1.6E-311).
- A tool developer must always take note of the special cases.
- XML Schema was modified to handle this type by hand

Solution

- A SimpleType was created.

```
<xs:simpleType name="sn-double">
  <xs:restriction base="xs:double">
    <xs:pattern value="-?\d\.\d+((E|e)(\+|\-)\d{2,3})?" />
    <xs:pattern value="\d+" />
    <xs:pattern value="-?\d+\.\d+" />
  </xs:restriction>
</xs:simpleType>
```

Step1: Using ANT

- Use the build.xml ANT script provided
- Targets are:
 - generate
 - compile
 - run
 - docs
- `ant -Djwsdp.home="" <target>`

Step2: Generating The Source

- ant generate
 - This target will run the JAXB Binding Compiler and generate your content tree Java source files.
- ant compile
 - This target will compile all sources in the src subdirectory.
- ant run
 - This target will run the `ca.bioinformatics.blastoutput.main.Main` class.
- ant docs
 - This target generates the javadoc for your content tree

Step3: Creating the Main class

- Modify the `ca.bioinformatics.dtt.main.Main` class.
- This will be your startup class which will initialize the JAXB environment, unmarshall, validate, and output a summary.

Step 4: Generate output

- Your application should generate the following information about your blast result.
- Accession of hit
 - E-value and score of HSPs

Results of BLAST search

```
Accession:NP_055202
    evalue=1.07425E-77      score=744.0
Accession:CAA11218
    evalue=1.76723E-58      score=579.0
Accession:NP_110480
    evalue=7.72818E-34      score=366.0
Accession:NP_034819
    evalue=3.83768E-31      score=343.0
Accession:AAH52340
    evalue=1.23418E-30      score=338.0
Accession:BAC40554
    evalue=1.75215E-30      score=337.0
Accession:NP_004723
    evalue=9.85871      score=72.0
[marc@blueprint-14 bin] $
```