

PMD code checking: checking for programming against CDK interfaces

Egon Willighagen 

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chem-bla-ics

I wrote recently about [programming against interfaces](#) , which led to some discussion. [Rajarshi commented](#):

patch 2 seems to replace usage of Molecule with IMolecule, rather than IMolecule

Now, this particularly problem was [Eclipse](#) being overly active cleaning up the imports, but we reached the point that it is actually interesting to have [PMD](#) warn about these issues. PMD is a cool piece of software used by the CDK project for a long time one. One particular cool thing is that you can write *code warnings* as [XPath](#) queries.

So, I promised Rajarshi to have a look at a custom rule to look at using IMolecule instead of Molecule. The code is actually pretty simple, using the aforementioned XPath feature:

```
<rule name="ReplaceMoleculeWithIMolecule"
  language="java"
  message="Use the IMolecule interfaces instead of the Molecule implementation"
  class="net.sourceforge.pmd.rules.XPathRule">
  <description>
    Programming against the CDK interfaces allows users to pick their favorite
    implementation (nonotify, datadebug).
  </description>
  <priority>3</priority>
  <properties>
    <property name="xpath">
      <value>
<![CDATA[
  //Type/ReferenceType/ClassOrInterfaceType[@Image='Molecule' ]
]]>
      </value>
    </property>
  </properties>
  <example>
    <![CDATA[
      Molecule mol = builder.newInstance(IMolecule.class);
    ]]>
  </example>
</rule>
```

Patch pending...