

Bioclipse gets a new extension point

Egon Willighagen 

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Abstract

I hacked in a new extension point for Bioclipse yesterday, based on a proposal I made earlier. The new extension point (EP) is called ChildResourceCreator and allows creating child resources for a given IBioResource. One application where this is very useful is the CMLRSS application (earlier blog), or any RSS or Atom enriched with any other XML language.

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chem-bla-ics

I hacked in a new extension point for [Bioclipse](#) yesterday, based on a [proposal](#) I made earlier. The new extension point (EP) is called `ChildResourceCreator` and allows creating child resources for a given `IBioResource`. One application where this is very useful is the [CMLRSS application](#) ([earlier blog](#)), or any [RSS](#) or [Atom](#) enriched with any other XML language. Here, child resources are created for each feed entry resource with as content the foreign XML, e.g. the CML bits in the blog.

Other applications involve complex documents, which is basically most existing documents. Take, for example, the [PDB format](#) from the [PDB database](#). These PDB files contain a plethora of information including one or more protein structures, sequences and bibliographic information. Bioclipse supports each of those using the [CDK](#), [BioJava](#) and [JabRef](#) libraries.

By making extension for the `ChildResourceCreator` EP, I am able to setup a general `PDBResource` (with Bioclipse's syntax highlighted PDB editor), and child resources for the different bits of information. [Bioclipse 1.0](#), however, only allow looking at the molecular structure(s) in the file, not at the sequence, nor the references. Will post the obligatory screenshot asap.