





NERC Environmental Bioinformatics Centre

The proliferation in the am ountof data available on line has given rise to many new challenges for those wishing to explicit in their research. Simply finding the data can be a problem. The sheer number and diversity of sources makes it very difficult for an individual to be aware of all of the possible data sets that might be of interest to them. A second significant issue is that of data-integration. Successful integration requires that unam biguous metadata descriptions are available to ensure that disparate data sets are com parable.



An important developm entwhich promises to help in both of these cases is the semantic web and the attendant rise in interest in ontologies. Ontologies offer the potential to assist in both the *searching* for (by enabling smarter matching and automatic generation of search terms) and the *interpretation* of data sets (where the unambiguous nature of ontological annotation facilitates the discovery of suitable mappings from one data set to another).



Figure 1 – Term matching

Emplying onto bgies in the annotation of data should not be unnecessarily burdensome to the user. Similarly, users cannot be expected to invest significant time in becoming intimately familiar with specific onto bgies, many of which contain thousands of terms. To this end, we are investigating methods for assisting non-expert users in annotating their data.

W e presenta toolthatautom atically detects onto by calterns in free text. Figure 1 illustrates the algorithm used by this tool. Once candidate terms have been identified the results are displayed either overlaid on the original text or in a list organised by onto by and frequency. Examples of these representations are shown in Figure 2.



The Term inizer services currently is built using the ontobgies from the OBO Foundry, a collection of over 40 biological ontobgies in a common format. Coupled with the GAZ gazetteer, the database presently contains 390,000 terms and 150,000 synonyms.We will shortly be expanding this to include ontobgies from the National Center for Biomedical Ontobgy. The user can interactively acceptor reject each match, or try to find a more appropriate match by exploring the network of ontobgy concepts them selves. In typical ontobgical resources, the parent(s) of a term represent broader concepts whilst the children of a term representmore specific concepts. In this way, the suggested match can used as a starting point for the user to find a more suitable term . Figure 3 illustrates the ontobgy brow ser interface, which has both a textual and a graphical mode.

9	Biological process	Biological process
	prowse mode : 🔍 tabular 🛛 graphical	browse mode : 🔘 tabular 🛛 🔍 graphical
		Current Selection
		steroid biosynthetic process
		"The chemical reactions and pathways resulting in the formation of steroids, compounds with 1,2,cyclopentanoperhydrophenanthrene nucleus; includes de novo formation and steroid interconversion by modification [GOC:go_curato]
GO Term ergosterol biosynthetic	strocess GO Term cholesterol biosynthetic	
	15 a	Broader Terms (less specific)
15_8	GO Term	lipid biosynthetic process
	sterol biosynthetic process	
GO Term	GO Term	
ecdysteroid biosynthetic process	is glucocorticoid biosynthetic	
	is a	Narrower Terms (more specific)
is a	GO Term Is a steroid biosynthetic process	sterol biosynthetic process
GO Term androgen biosynthetic process	ic a	glucocorticoid biosynthetic process
		androgen biosynthetic process
CO Tama	GO Term bile acid biosynthetic process is_a	ecdysteroid biosynthetic process
C21steroid hormone biosynthetic process	50	Internatioconicolo biosynthetic process
7	phytosteroid bic	Breatly Colorial Terma
		necenily selected remis
	iya	androgen biosynthetic process
		steroid biosynthetic process

In addition to the interactive mode, the software is also available as a W eb service. Both the term detection service and the interactive presentation layer can be incorporated within other W eb sites or programs.

The Tem inizer system has been built using the omixed fram ework, an architecture for supporting the rapid deployment of collaborative databases. More information about om ixed and Tem inizer, including a live demonstration of the service, is available on our website:

http://terminizer.org/



Figure 3 – The ontobgy explorers

