Attendees: Scott, Alasdair, Erich, Joachim, Michel, David, Kerstin

Restarting back after the summer recess.

Scott provided a summary of the goals of the work; describing datasets for use for SPARQL access and reasoners.

David joined from the EU Eureka project about clinical data. Common goals. Need to capture various provenance properties, but this largely seems to be in common with the properties that are already captured.

Do we need to have a clinical data extension or can it be captured dcat:theme?
Should be expressible through the vocabularies, themes, and the statistics about the predicate/subject/etc that appear in the data.
The vocabularies and statistics can be auto-generated from the data using SPARQL queries.
An example is available from Bio2RDF at http://download.bio2rdf.org/release/2/gene/gene.html

Open PHACTS have already created a VoID generation tool and validation tool which could be extended to the HCLS Note. The existing tools are available from
http://openphacts.cs.man.ac.uk/Void-Editor/
http://openphacts.cs.man.ac.uk:9091/Validator or
http://openphacts.cs.man.ac.uk:9094/Validator (These are slightly different versions the first is for the Open PHACTS profile while the second is being developed for the HCLS profile.)

The VoID Editor is the subject of a third year project which will look at auto-generating as much of the metadata as possible and using Q&A to gather the rest of the required metadata. The student will be starting this month and running until Easter.

Actions:

- Scott to come up with a timeline to generating the note
- Doodle poll about whether these meetings are weekly or bi-weekly