Attendees: Erich Gombocz, Joachim Baran, Michel Dumontier, Scott Marshall
Excused: Alasdair Gray
Scribe: Joachim Baran

- recapture of last week's meeting
    - use of IANA, ontologies, etc.
- discussion of "availability" related linked data items
    - use of dcat:distribution for downloadable data
        - Michel provided an example (part of vocab-dcat)
        - general mechanism for linking any file type
    - RDF dump
        - for RDF only, having both void:dataDump and dcat:Dataset
    - SPARQL endpoint
        - use void:sparqlEndpoint
    - REST API -- renamed "API documentation page"
        - documentation web-page
        - also applicable for SOAP based services
- discussion of "statistics" related linked data items
    - "should" properties: number of statements, entities, subjects, properties and objects, distinct predicates and types
    - "may" properties: number of named graphs
        - initially decided to have one graph per dataset at a previous BioHackathon
        - several sources offer multiple graphs though (UniProt, for example, at http://beta.sparql.uniprot.org)
        - discuss with Jerven Bolleman to get data provider feedback
    - Michel points out that Virtuoso can also calculate these void:* numbers
    - Scott remarks that we should later provide examples for calculating these stats generically (across various tools); Joachim concurs
    - void:LinkSet to describe interdataset link statistics (predicate-object links, etc.)
        - Michel gives Bio2RDF example: number of GO terms in a dataset