Are SNOMED CT Browsers Ready for Institutions? Introducing MySNOM

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Abstract. SNOMED Clinical Terms (SNOMED CT) is one of the most widespread ontologies in the life sciences, with more than 300,000 concepts and relationships, but is distributed with no associated software tools. In this paper we present MySNOM, a web-based SNOMED CT browser. MySNOM allows organizations to browse their own distribution of SNOMED CT under a controlled environment, focuses on navigating using the structure of SNOMED CT, and has diagramming capabilities.

Keywords: SNOMED CT, Terminologies, Ontologies, Browser

1 Introduction

The Systematized Nomenclature of Medicine - Clinical Terms (SNOMED CT) has become a reference terminology and an active research topic in both the healthcare and the Semantic Web communities. An effective way to advance research in ontologies for the life sciences in general, and in SNOMED CT in particular, is by providing appropriate software to researchers and practitioners. In particular, visualization and browsing tools can provide a better understanding of the structure and complexity of a biomedical terminology. There are lists of existing browsers, such as the ones provided by The National Library of Medicine of the United States and the United Kingdom’s National Health Service, in order to provide researchers and practitioners with a reference. These lists, however, are not comprehensive, not up to date, and have not been fully ranked or evaluated. These lists can also be misleading for practitioners: as an example, Protégé is listed as a SNOMED CT browser. In 2008, Rogers performed a study of existing SNOMED CT browsers and developed a general catalog of desirable browsing features.

In this paper we present a SNOMED CT web-based browser called MySNOM, aiming at showing the structure of SNOMED, with support for rich diagrams.

1 www.nlm.nih.gov/research/umls/Snomed/snomed_browsers.html
2 www.connectingforhealth.nhs.uk/systemsandservices/data/snomed/browser
3 protege.stanford.edu
2 Architecture Overview

MySNOM is a web application following a layered approach to decouple orthogonal functionalities such as access control, graphical interface, business logic and persistence. The architecture is shown in Figure 1.

![Architecture Diagram]

**Fig. 1.** MySNOM architecture.

MySNOM is hosted in a CGI-enabled web server and accessed using a web browser. It relies on the lightweight and standard HTTP Digest protocol [5] for authentication of users, and SQLite [4] to provide fast access to the terminology.

3 User Interface and Diagramming Support

MySNOM uses a clean interface to explore all the relationships related to a given concept, including reverse relationships. It places the current concept in the middle of the screen, with referring concepts on the left and referred concepts on the right. This structure follows the extended left-to-right direction of flow and shows the natural directed graph structure of SNOMED CT. Every concept and relationship is clickable. Each clicked concept or relationship leads to a new screen, becoming the current term. The current concept is shown only once on screen. Users can hover their mouse over any given concept to further inspect the concept (e.g. see the SNOMED CT code). A screenshot of MySNOM browsing the term ‘chronic disease’ is shown in Figure 2.

MySNOM supports the generation of diagrams that might clarify the context of a given concept or serve as a graphic resource to be embedded in SNOMED

[4] www.sqlite.org
CT-related documents. It uses different colors to emphasize several concepts: the main concept is marked in yellow and hierarchy relationships are colored in red. The diagramming module creates an intermediate representation of the browsed concept using the DOT language \[6\]. The DOT language is a plain text description that can be read by humans and processed by visualization tools. MySNOM uses Graphviz \[7\], an Open Source, fast, graph generation software, to generate the final diagram. Generated diagrams reflect the ontological nature of SNOMED CT, as a directed graph, showing all concepts and relationships of a given concept. A sample diagram is shown in Figure 3.
4 Conclusion and Future Work

We have presented an access-controlled, web-based, lightweight browser in order to provide structural navigation and concise diagrams for SNOMED CT, using an authorized standard distribution of SNOMED CT. The browser can be viewed and tested at [8] and freely obtained by contacting the author. As future work we plan to improve the prototype and add modularization support. Modularization support will let a user store and export selected portions of SNOMED CT by using ontology modularization techniques [9]. A tool with these characteristics will allow a deep research of SNOMED CT’s structure and capabilities.

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References

1. Freitas, F., Schulz, S.: Survey of current terminologies and ontologies in biology and medicine: Elect. J. Commun. Inf. Innov. Health vol. 3, no. 1, pp. 7-18. (2009)
2. Spackman, K., Reynoso, G.: Examining SNOMED from the perspective of formal ontological principles: Some preliminary analysis and observations. In: Proceedings of the 1st International Workshop on Formal Biomedical Knowledge Representation, pp. 81-87. (2004)
3. Héja, G., Surján, G., Varga, P.: Ontological analysis of SNOMED CT: BMC Medical Informatics and Decision Making vol. 8, suppl. 1, S8. (2008)
4. Rogers, J., Bodenreider, O.: SNOMED CT: Browsing the Browsers. In: Proceedings of the 3rd International Conference on Knowledge Representation in Medicine, pp. 30–36. (2008)
5. Franks, J., Hallam-Baker, P., Hostetler, J., Leach, P., Luotonen, A., Sink, E., Stweart, L. RFC2069: An Extension to HTTP : Digest Access Authentication. RFC. (1997)
6. Eleftherios Koutsofios, E., North, S. C.: Drawing Graphs With Dot (1996)
7. Ellson, J., Gansner, E., Koutsofios, L., North, S., Woodhull, G.: Graphviz Open Source Graph Drawing Tools. In: Mutzel, Petra and Jnger, Michael and Leipert, Sebastian (eds.). LNCS, vol. 2265, pp. 594–597. Springer, Heidelberg (2002)
8. BDI Group, MySNOM SNOMED CT Browser demo, [http://bdid.si.ehu.es/mysnom/](http://bdid.si.ehu.es/mysnom/) (user: demo, password: demo).
9. Stuckenschmidt et al.: Modular Ontologies: Concepts, Theories and Techniques for Knowledge Modularization: LNCS. Springer, Heidelberg (2009)