Developing Ontology Support for Human Malaria Control Initiatives
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Abstract
Malaria is one of the most common infectious diseases and an enormous public health problem in Sub-Sahara Africa, Asia and parts of America. In this paper, we discuss the development of the Human Malaria Control Ontology (HMCO) which contains general information on Malaria and epidemiological information that can help in the formulation of effective malaria control policies. The HMCO is aimed at providing interoperability support for the knowledge management of malaria control initiatives, and serve as an open semantic web infrastructure for malaria research and treatment.

Introduction
Ontology is a formal explicit representation of the conceptualization of a domain that provides a platform for the sharing and reuse of knowledge across heterogeneous platforms. An ontology contains semantic descriptions of the features of a domain using concepts and relationship abstractions in a way that is readable by both man and machine. In recent times, the use of ontology have gained increasing relevance in the biomedical domain in that it enables researchers to stay abreast of current biomedical knowledge and promotes the understanding of such information. They also facilitate the sharing and reuse of biomedical knowledge across heterogeneous platforms for the delivery of medical services and implementation of health-related policies.

Malaria is one of most worrisome vector-borne diseases that affect humans. It is caused by the parasite Plasmodium falciparum. Malaria is endemic in the tropical regions and sub-tropical regions of the world which are mainly in the South East Asia, Middle East Asia, Central and South America, and Sub-Sahara Africa. Millions of malaria cases are reported each year, killing over one million per year in Sub-Sahara Africa. Generally, the control of a vector-borne disease such as malaria pose a critical challenge due to a number of reasons: 1) the complex nature of its transmission which involves three entities, which are the host (human), vector (female anopheles mosquito) and pathogens (Plasmodium specie); 2) the complicated epidemiology through the vector; and 3) social problems (poverty), geography, and resistance of pathogens to insecticides. All of these challenges compel the need to complement existing biomedical approaches of tackling the spread of vector-borne diseases with readily accessible, interoperable, and semantically-rich knowledge management support. This challenge motivated our pursuit of developing an ontology-based support for human malaria control. The Human Malaria Control Ontology (HMCO) contains information on human malaria that can be leveraged for the formulation of human malaria control initiatives in Sub-Sahara Africa. In terms of benefits the HCMO is expected to: 1) provide an interoperable platform for accessing malaria epidemiology information over the web; 2) provide information support for malaria control research and formulation of malaria control policy initiatives; and 3) Create an interoperable platform for the sharing and reuse of knowledge on malaria.

The outline of the rest of this paper is given as follows. In Section 2 an overview of related work on medical ontologies is presented. Section 3 is a short description of the design and implementation of the HMCO. Section 4 is a discussion of the possible application of the HMCO, while the paper is concluded in Section 5 with an outlook of future work.

Related Work
Medical Ontologies have played useful roles in facilitating the re-use, dissemination and sharing of patient information across disparate platforms. Also, they have been used in semantic-based statistical analysis of medical data. Examples of medical ontologies include GALEN, UMLS, MeSH, OMICs, and Tambis. The Systematized Nomenclature of Medicine, Foundational Model of Anatomy, MENELAS ontology, Gene Ontology and LinKBase. The NBCO’s Bio-portal consists of more than 50 bio-ontologies that span several aspects of bio-medicine including diseases, biological processes, plant, human, bio-medical resources etc. However, none of the ontologies in the bio-portal is specifically dedicated to malaria control. The work by Hadzic and Chang was based on providing interoperability support for research in, and diagnosis of human disease using ontology-based approach. A prototype Generic Human Disease Ontology (GenDO) that contains common general information regarding human diseases was created which captured the information in four dimensions. However, the dimension of diseases control was not included.

The Infectious Diseases Ontology (IDO) is designed to make infectious disease-relevant data derived from different sources comparable and computable. It also provides coverage of entities that are common to many infectious diseases. The Vector-borne disease ontology is an ongoing project that is designed to provide an integrated interoperable platform for the sharing and reuse of knowledge about a group of vector-borne diseases in which the MalIDO (Malaria IDO) is a first step. The MalIDO incorporates several information dimensions such as gene models for A.gambia, Anatomy of mosquito, insecticide-resistance, and physiological processes of mosquito. As a contribution our work in the HMCO specifically focuses on the creation of an interoperable platform that gives access to epidemiological information on malaria in Sub-Saharan Africa that can be used for the formulation of malaria control policies.

**Description of the HMCO**

The HMCO captures information on human malaria in 7 dimensions. These are: (1) Malaria vectors (2) Malaria types, (3) Malaria parasites, (4) Malaria Symptoms, (5) Malaria treatment (prevention, therapy), (6) Epidemiology data on malaria, and (7) Malaria Control. The design of HMCO was based on the Open Biomedical Ontologies (OBO) foundry principles, while OBO foundry naming conventions were also adopted significantly in naming its concepts. It was implemented as an OWL ontology using the Protégé 3.4 Ontology tool. The conceptual taxonomy of the HMCO consists of 97 class abstractions that cover the seven dimensions of our interest. Nine disjoint subclasses comprising vector, treatment, continent, type, parasite, epidemiology-info, symptom, year-data, and control were modelled as constituents of the superclass human_malaria using ‘belongTo’ object property. The subclasses for the three classes: symptom, treatment, and continent were modelled as OWL value partitions (viz. each of the classes was represented as comprising disjunctive subclasses that cover all known instances of each class). Concepts relationships among classes (concepts) in the HMCO class hierarchy were represented using object property abstractions that define the nature of association between the classes. These include associations between parasite and vector (‘hasVector’), type and symptom (‘hasSymptom’), type and malaria_therapy (‘hasTherapyDrug’), type and malaria_prevention (‘hasPreventionDrug’), type and parasite (‘isCausedby’), parasite and malaria_therapy (‘isCuredby’), parasite and malaria_prevention (‘isPreventedby’), vector and continent (‘isFrom’), epidemiology_info and year_data (‘hasEpidemiymdata’) etc. The class hierarchy and description of entities in the HMCO is shown in Table 1 (appendix).

Also, specific object properties and datatype properties in the HMCO have appropriate cardinality restrictions imposed on them in order to effectively capture the semantics of relationships among the classes in the HMCO. Figure 1 (appendix) is a view of the class hierarchy of the HMCO.

**Application of the HMCO**

Currently, the HMCO knowledgebase is being populated with available data (at present we have epidemiological data on malaria from the year 1999 to 2003 for 16 countries in
Sub-Saharan Africa), while the data gathering process is still ongoing. At the completion of the first version of the HMCO, it is expected to serve as a web-based repository for accessing epidemiology information on malaria in Sub-Saharan Africa. As a first step to attaining this, a prototype semantic web application has been built that can be used to query the HMCO knowledgebase.

The Java programming language implementation technology was engaged in building the prototype semantic web application using the NetBeans Java IDE. The Web GUI that facilitates client interaction with the HMCO knowledgebase was implemented using Macro Media Flash and Dream Weaver web design tools, and Java Server Pages (JSP). The business logic for querying the HMCO was implemented as an Enterprise Java Beans (EJB) component that is invoked from a Java Servlet class running on Sun Application Web Server 9.0. The EJB makes use of Protégé ontology Java APIs to access the HMCO knowledgebase for information retrieval. The Pellet 1.5 Descriptive Logics (DL) reasoner was used as the OWL DL reasoner to facilitate semantic web reasoning (entailment, subsumption, and ABox reasoning) on the classes and individuals in the HMCO.

Conclusion

In this paper, a description of the Human Malaria Control Ontology (HMCO) has been presented. The HMCO offers as its contribution, an interoperable platform for accessing epidemiological information on malaria as viable knowledge management infrastructure for malaria control policy formulation and research in Sub-Saharan Africa. Though still an ongoing work, a preliminary test of the usability of the HMCO has been undertaken with promising results. Subsequently, the HMCO will incorporate other dimensions of information on malaria by importing relevant ontologies like the IDO and the Vector-borne diseases ontology (particularly the MalIDO) and will be presented for submission to the bio-ontology portal for open access and evaluation.

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• human_malaria
  o control
    drainage
    mosquito_net
    insecticide
  o epidemiology_info
    Drc_Data; Kenya_Data;
    Angola_Data; Nigeria_Data;
    Mozambique_Data; Namibia_Data;
    Ghana_Data; Sudan_Data;
    CAR_Data; Mali_Data;
    Uganda_Data; Malawi_Data;
    Tanzania_Data; Zambia_Data
  o Parasite
    P. falciparum
    P. ovale
    P. malariae
    P. vivax
  o Symptom
    rigor
    vomiting
    convulsion
    haemoglobinuria
    anemia
    fever
    arthralgia
    shivering
    abnormal_posturing

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Figure 1. A view of class hierarchy of the HMCO

| Class name          | Description                                                                 |
|---------------------|-----------------------------------------------------------------------------|
| retina_damage       |                                                                             |
|                      | treatment                                                                   |
|                      | malaria_prevention                                                          |
|                      | hydroxyxhloriquine;                                                        |
|                      | chloroquine; proguanil;                                                     |
|                      | doxycycline; mefloquine;                                                    |
|                      | atavoquone- proguanil                                                       |
|                      | malaria_therapy                                                             |
|                      | artesunate- sulfadoxine-pyrimethamine;                                     |
|                      | artesunate- amodiaquine;                                                    |
|                      | artesunate- mefloquine;                                                    |
|                      | artemether- lumefantrine;                                                   |
|                      | quinine; cotrifazid;                                                        |
|                      | sulfadoxine- pyrimethamine;                                                 |
|                      | primaquine                                                                   |
|                      | type                                                                        |
|                      | severe_malaria                                                              |
|                      | chronic_malaria                                                             |
|                      | vector                                                                      |
|                      | A.gambiae; A.freeboni,                                                       |
|                      | A.culicifacous; A. fluviatilis; A. minimus; A. phillipinesis; A. stephensi; |
|                      | A. leucosphyrushave                                                         |
|                      | Continent                                                                   |
|                      | Africa; Antarctica; Asia; Australia; Europe; North_America; South_America |
|                      | year_data_1990 ... _2008                                                   |
|   | Class Name          | Description                                                                 | Examples                                                                 |
|---|---------------------|------------------------------------------------------------------------------|--------------------------------------------------------------------------|
| 1 | human_malaria       | Main ontology superclass                                                     | control, epidemiology_info, parasite, symptom, treatment, type, vector, continent, year_data |
| 2 | control             | Defines the concepts under malaria control                                  | severe_malaria, chronic_malaria                                          |
| 3 | epidemiology_info   | Captures information on epidemiology data of 16 countries in Sub-Sahara Africa | Drc_Data, Kenya_Data, Angola_Data, Nigeria_Data, Mozambique_Data, Namibia_Data, Ghana_Data, Sudan_Data, Central African Republic (CAR)_Data, Mali_Data, Uganda_Data, Malawi_Data, Tanzania_Data, Zambia_Data |
| 4 | parasite            | Defines the different types of human malaria pathogens                       | P. falciparum, P. ovale, P. malariae, P. vivax                           |
| 5 | symptom             | Defines the different types of malaria symptoms as a set of disjointed classes. | rigor, vomiting, convulsion, haemoglobinuria, Anemia, Fever, arthralgia, shivering, abnormal_posturing (children), retina_damage (children) |
| 6 | treatment           | Defines the types drugs for prevention and treatment of human malaria         | malaria_prevention, malaria_therapy                                      |
| 7 | type                | Defines the types of human malaria                                            | severe_malaria, chronic_malaria                                          |
| 8 | vector              | Defines the different types of human malaria vectors                          | A.gambiae, A.freeboni, A.culicifacious, A.fluvialis, A. minimus, A. philippensis, A. stephensi, A. leucosphyrushave |
| 9 | continent           | Defines continental regions to which specific malaria vectors belong.       | Africa, Antarctica, Asia, Australia, Europe, North_America, South_America |
| 10| year_data           | Defines information on malaria endemics on a yearly basis for 19 years.     | _1990 ... _2008                                                          |

Table 1. An Overview of Classes in the HMCO