Domain Concept-Based Queries for Cancer Research Data Sources

Alejandra González Beltrán

Joint work with Anthony Finkelstein (UCL), J Max Wilkinson (NCRI) and Jeff Kramer (ICL)





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Overview

ONIX (ONcology Information eXchange) – UK NCRI platform to facilitate access to distributed cancer-research data sources

□ Interoperability with the caGrid infrastructure

Support for non-caGrid resources

- caGrid metadata infrastructure and query language
 - Queries based on the structure of the resource
 - No support for concept-based queries

Goal: support for high-level and descriptive queries of cancer-research data sources, expressed using domain concepts and their relationships

- Architecture & Approach: caGrid + Semantic Web technologies
 - Concept-based queries and data integration
 - Extensible to non-caGrid resources

Conclusions















UCL

caGrid infrastructure

caGrid Query Language (CQL)

Simple object-oriented query language (semantic annotations are not considered)

Procedural (client must be aware of resource's structure)



Structural layer



Motivating example

A biomedical scientist is interested in finding single nucleotide polymorphisms (SNPs) associated with the gene Transforming Growth Factor Beta 1 (TGFB1)

caGrid query language (CQL)



Architecture

Web Ontology Language (OWL): W3C recommendation for knowledge representation

Reasoning: inference capabilities

OWL generation service: develops OWL ontologies from information models

Semantic query service: transforms concept-based queries to CQL using the generated ontologies + reasoning

Prototype: OWLAPI and Pellet





Approach

1) Generate an ontology from the data service metadata (annotated UML to OWL transformation)



Approach

2) Express the concept-based query over the generated ontology

□ Find objects that have concept *Single_Nucleotide_Polymorphism* and have an association with objects whose concept is *Gene*, which in turn have an attribute with concept *Gene_Symbol*, whose value is "TGFB1"

Approach

2) Express the concept-based query over the generated ontology

- □ Find objects that have concept *Single_Nucleotide_Polymorphism* and have an association with objects whose concept is *Gene*, which in turn have an attribute with concept *Gene_Symbol*, whose value is "TGFB1"
- hasConcept some Single_Nucleotide_Polymorphism and hasAssociation some (hasConcept some Gene and hasAttribute some (hasConcept some Gene_Symbol and hasValue value "TGFB1")

Description Logic query (DL-query) in Manchester OWL Syntax

Approach

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Description Logic query (DL-query) in Manchester OWL Syntax

□ **Concept-based** query: it can be used for any resource annotated with the same vocabulary

- □ **High-level** query: it is not based on the structure of a particular target resource
- **Descriptive**: it gives the criteria for the desired data

Approach

3) Transform the query using the generated ontology into the caGrid query language



Conclusions

- Support for high-level descriptive queries based on domain concepts for caGrid data services
- Approach generates ontologies from caGrid metadata and uses ontologies+reasoning to translate concept-based queries to CQL
 - □ Same concept-based query applicable to all the relevant resources
 - □ No need for the user to be aware of the structure of the target resource
- General approach: it is applicable to other resources exposing metadata and semantic annotations for query translation and data integration
 - James P. McCusker, Joshua A. Phillips, Alejandra González Beltrán, Anthony Finkelstein, Michael Krauthammer "Semantic web data warehousing for caGrid" BMC Bioinformatics SWAT4LS Supplement, in press 2009.
- Prototype Demo http://tinyurl.com/o6uw7z



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Thank you!



Questions?