

caCDE-QA: A quality assurance platform for cancer study common data elements

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Introduction

- Semantic interoperability among terminologies, data elements, and information model is fundamental and critical for sharing information from the scientific bench to the clinical bedside and back among systems.
- Domain-specific Common Data Elements (CDEs) are emerging as an effective approach to standards-based clinical research data storage and retrieval and have been broadly adopted.



Introduction

- National Cancer Institute (NCI) created the Cancer Data Standards Repository (caDSR) based on the ISO/IEC 11179 standard for metadata repositories.
- In the ISO/IEC 11179, a data element is defined as a unit of data for which the definition, identification, representation and permissible values are specified by means of a set of attributes.
- The binding of controlled terminology provides the basis for semantic scaling of the CDEs.



Challenges

- The potential of terminology-based scaling has not yet been fully explored.
- In particular, there is a very limited toolbox at present for quality assurance (QA) of meta-data registered in such a repository like the caDSR.
- CDE content errors can have a significant negative impact on downstream CDE uses.

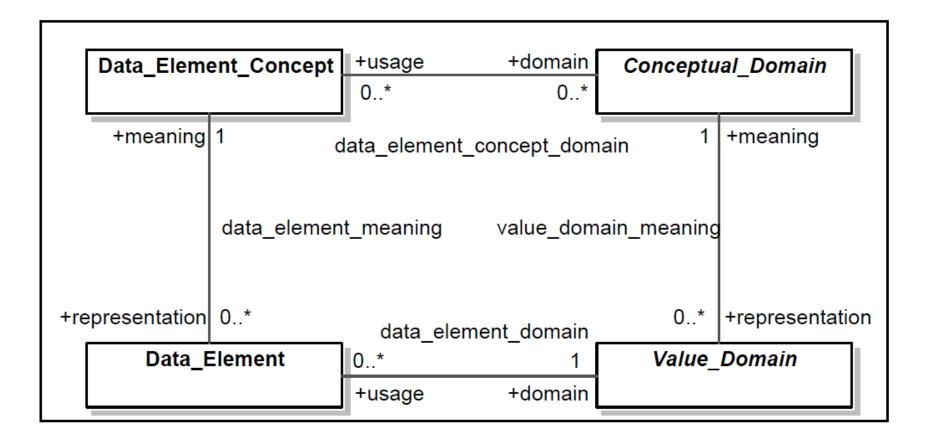


Specific Aims

- Aim 1: To develop a suite of QA tools for validation and harmonization of cancer study CDEs;
 - UMLS Semantic Network-based approaches
 - Semantic Web-based approaches
- Aim 2: To apply the QA tools to audit experimental cancer study CDEs represented in a semantic web framework;
 - NCI caDSR
 - Preferred sets of CDEs from TCGA data dictionary
- Aim 3: To deploy and evaluate a QA web-portal for collaborative CDE review and harmonization.
 - Specification for Standard CDE Services

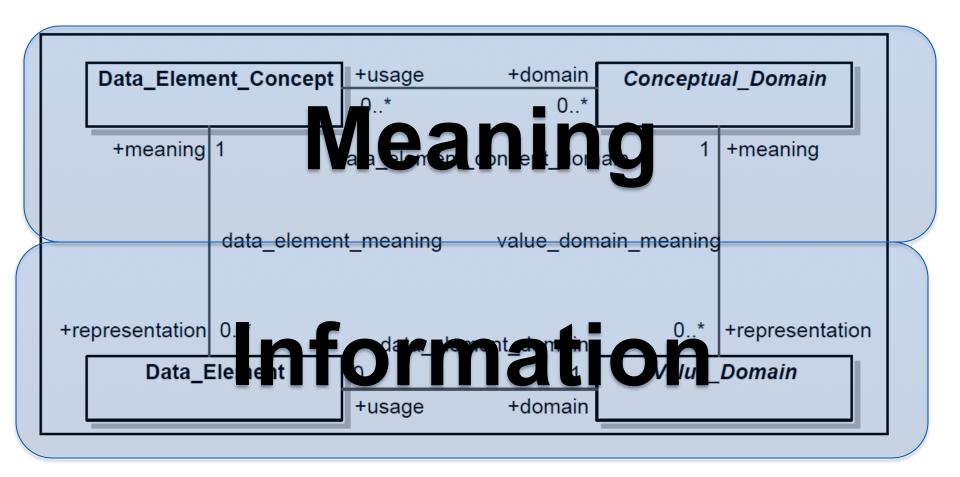


High-level data description meta-model in ISO 11179 specification

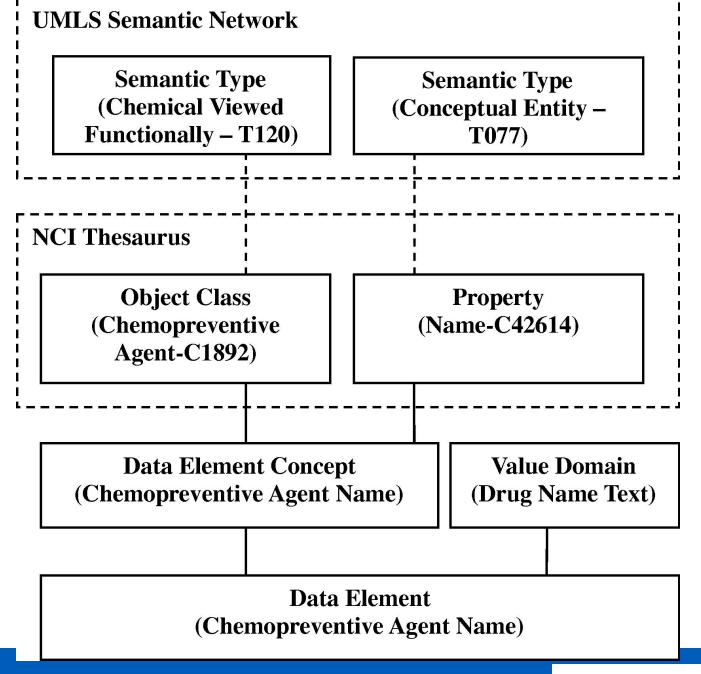




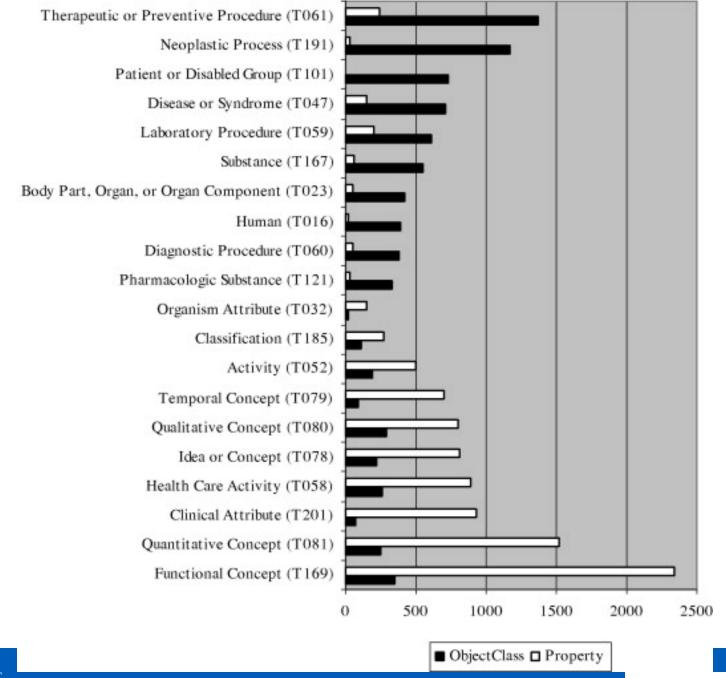
Information and Meaning



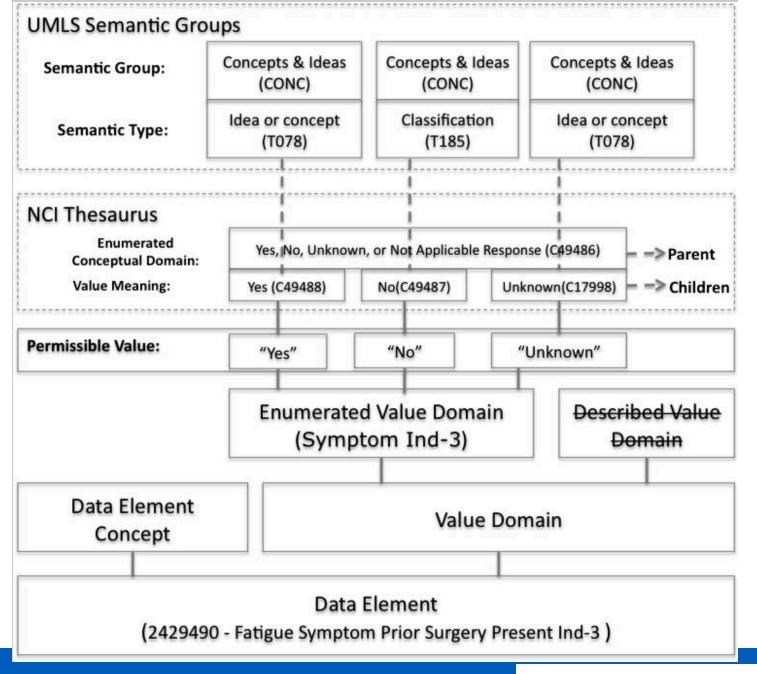














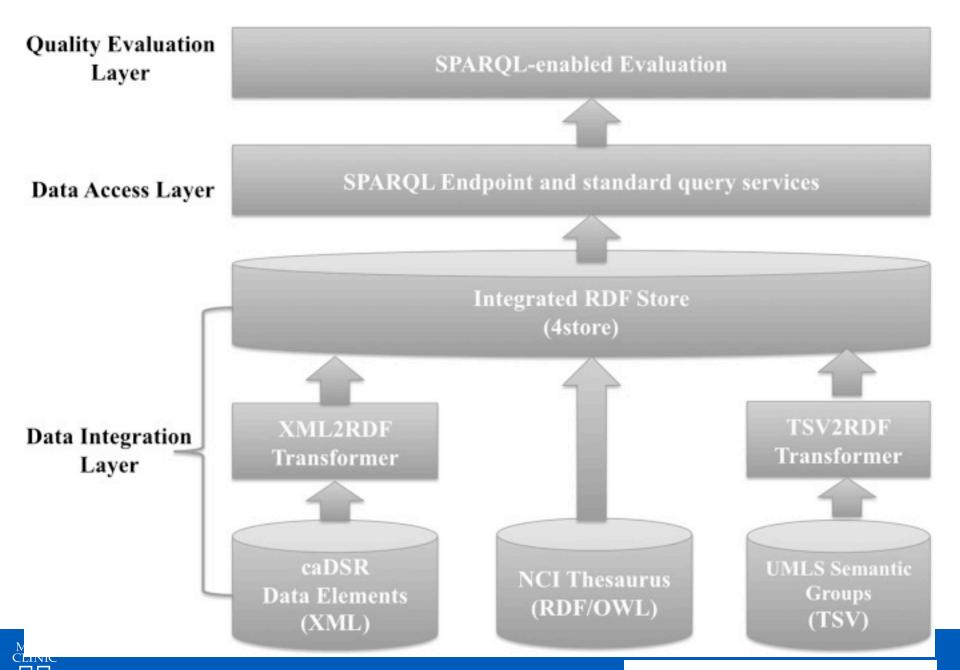


Table 3

Permissible values of the first example from the data elements identified with inconsistent codes (highlighted in bold and italic)

Valid value	Value meaning	NCIt code (meaning concept)	NCIt concept label	Semantic type	Type code	Semantic group	Group code
Palpatation	Palpation	C16950	Palpation	Diagnostic Procedure	T060	Procedures	PROC
CT	Computed Tomography	<u>C17204</u>	Computed Tomography	Diagnostic Procedure	T060	Procedures	PROC
Chest x-ray	Chest Radiography	C38103	Chest Radiography	Diagnostic Procedure	T060	Procedures	PROC
Spiral CT	Spiral CT	C20645	Spiral CT	Diagnostic Procedure	T060	Procedures	PROC
Plain x-ray	X-Ray	<u>C17262</u>	X-Ray	Natural Phenomenon or Process	T070	Phenomena	PHEN
MRI	Magnetic Resonance Imaging	<u>C16809</u>	Magnetic Resonance Imaging	Diagnostic Procedure	T060	Procedures	PROC

Data element: Malignant Neoplasm Measurable Disease Evaluation Method Clinical Trial Eligibility Criteria Type (3179024). Value domain: Malignant Neoplasm Measurable Disease Evaluation Method Type (3179022).

NCIt, National Cancer Institute Thesaurus.

C17262|X-Ray -> **C38101**|Radiography

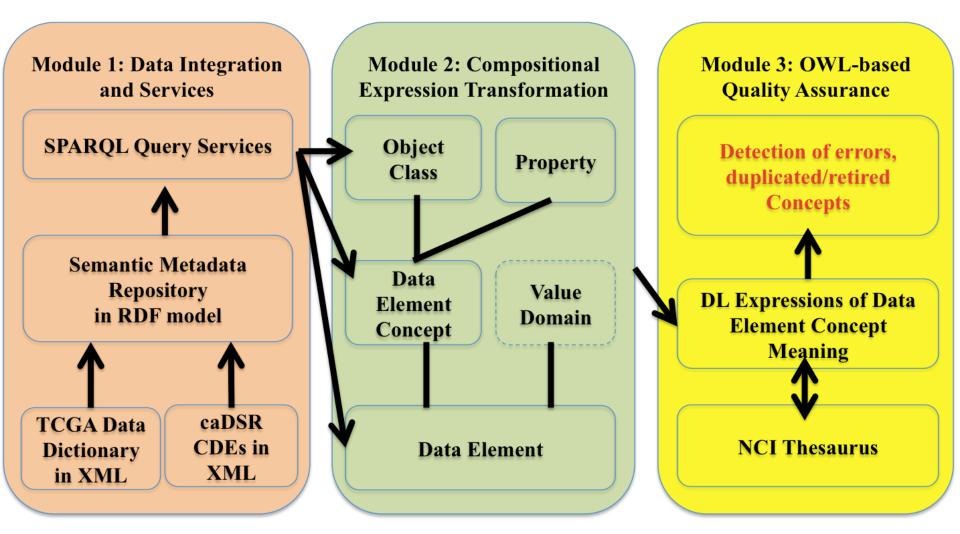
A radiographic procedure using the emission of X-Rays to form an image of the structure penetrated by the radiation



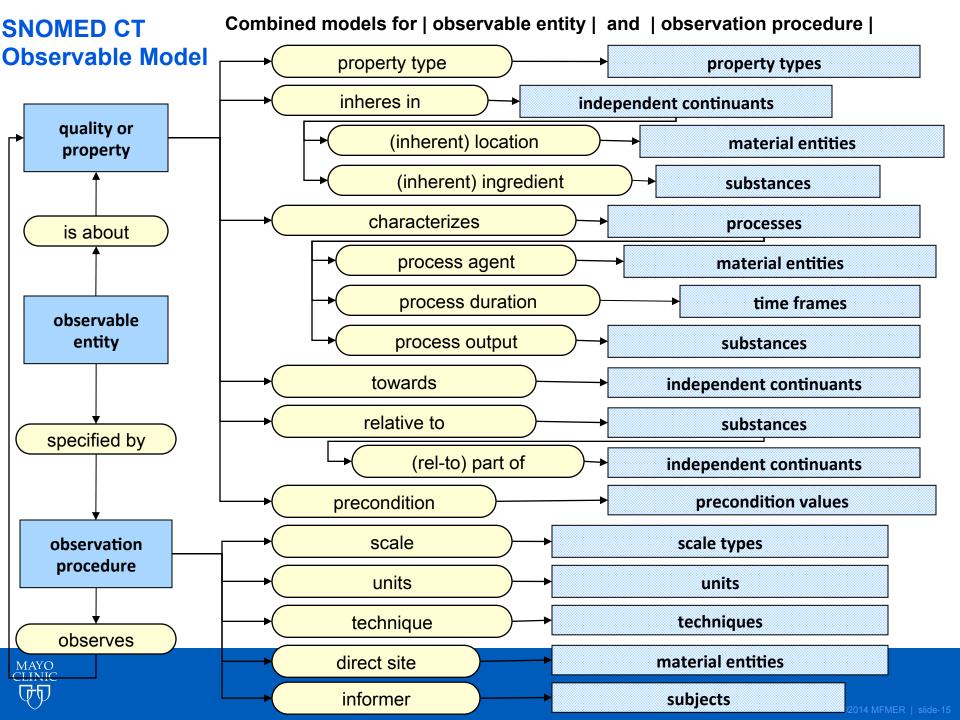


Progress to date

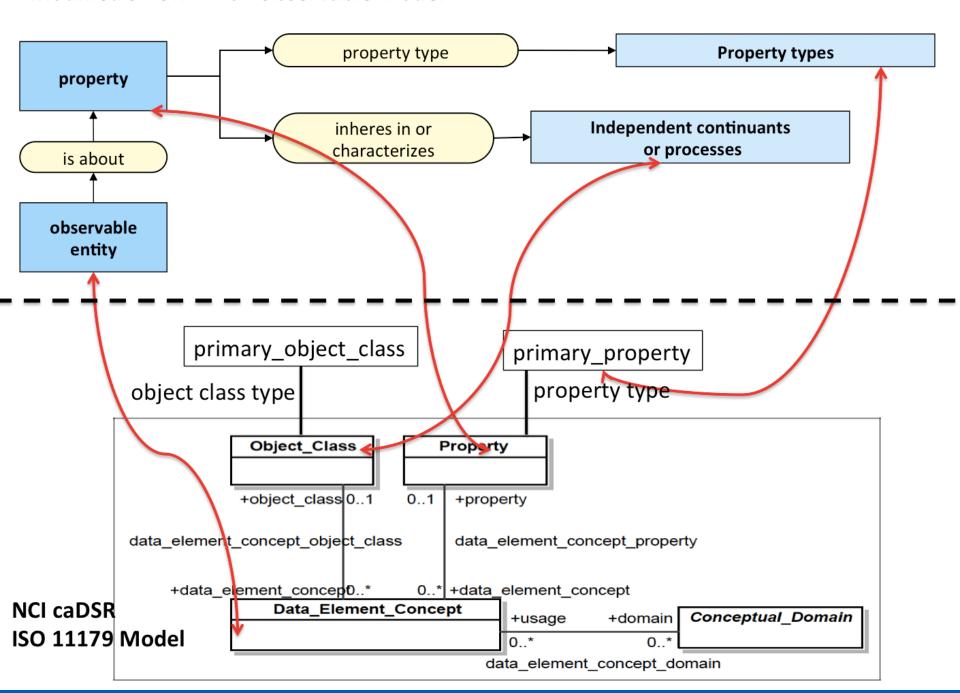
QA of cancer study CDEs using a post-coordination approach







Modified SNOMED CT Observable Model



An example of compositional expression

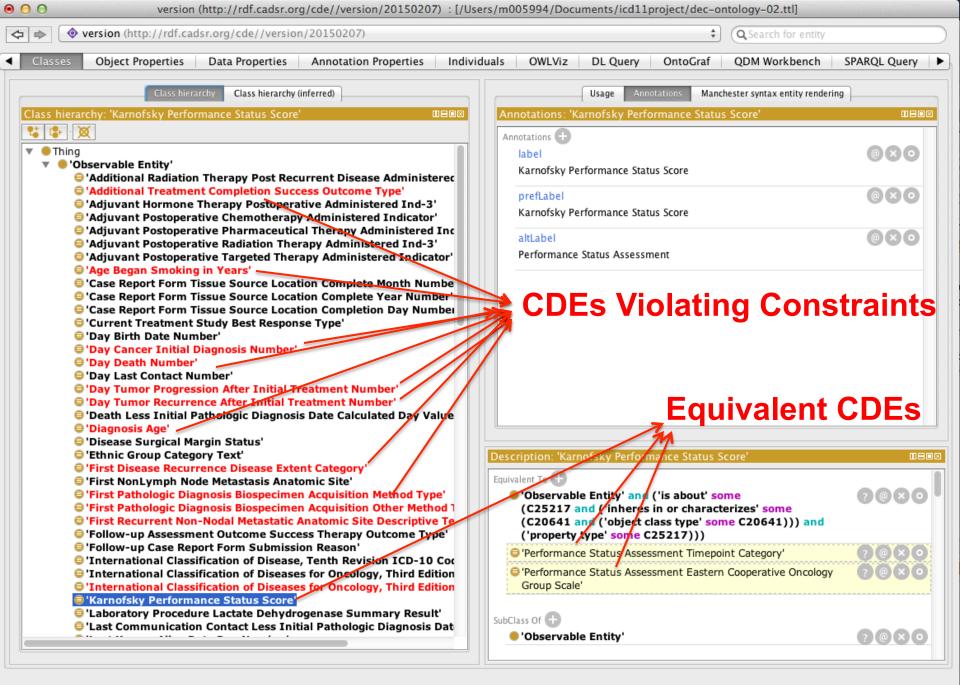
•	
Original Data Recorded in caDSR	Transformed Compositional Expression
Public Id: 3378323	Class: 'Clinical Trial Drug Classification Name'
CDE Name: Clinical Trial Drug Classification Name	Annotations:
Property Code: C25161	label "Clinical Trial Drug Classification Name"
Property Name: Classification	EquivalentTo:
Primary Property: C25161	'Observable Entity'
Object Class code: C71104:C1708	and ('is about' some (Classification
Object Class Name: Clinical Trial Agent	and ('inheres in or characterizes' some ('Clinical Trial Agent'
Primary Object Class: C1708	and ('object class type' some Agent)))
	and ('property type' some Classification)))
	SubClassOf:
	'Observable Entity'



OWL-based constraints

- Disjointness constraint (owl:disjointWith)
 - The disjointness of a set of classes guarantees that an individual (or a subclass) that is a member (or a subclass) of one class cannot simultaneously be an instance (or a subclass) of a specified other class
- Domain/range constraint (rdfs:domain, rdfs:range)
 - restricts each object property linking between the instances of asserted classes





Future Plan for OWL-based QA tool

- 1) developing a systematic QA approach leveraging upper level ontologies;
- 2) refining the compositional expression model by aligning with SNOMED observable model;
- 3) making the reasoning-based explanation more user-friendly; and
- 4) incorporating the value domain in the scope.



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Building A Semantic Web-based Metadata Repository for Facilitating Detailed Clinical Modeling in Cancer Genome Studies

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Abstract.

Detailed Clinical Models (DCMs) have been regarded as the basis for retaining computable meaning when data are exchanged between heterogeneous computer systems. To better support clinical cancer data capturing and reporting, there is an emerging need to develop informatics solutions for standards-based clinical models in cancer study domains. The objective of the study is to develop and evaluate a use case-driven approach that enables a Semantic Webbased cancer study metadata repository based on both ISO11179 metadata standard and Clinical Information Modeling Initiative (CIMI) Reference Model (RM). We used the common data elements (CDEs) defined in The Cancer Genome Atlas (TCGA) data dictionary, and extracted the metadata of the CDEs

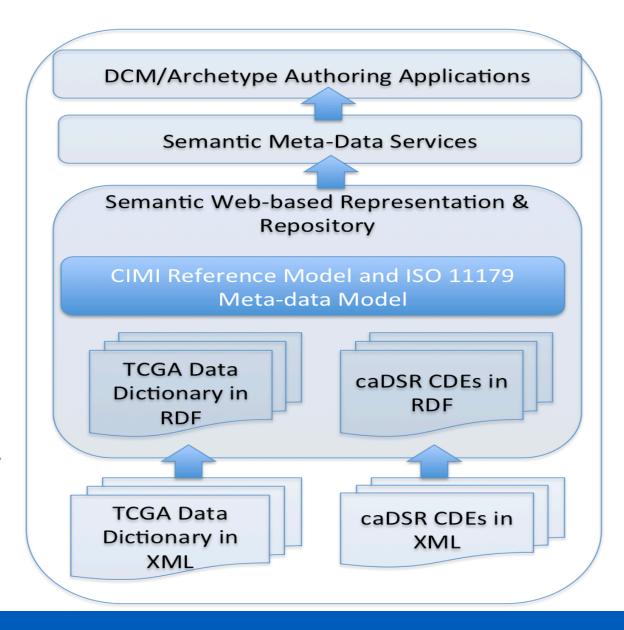


Application Layer

Services Layer

Persistence Layer

Transformation Layer





Study Plan for QA of cancer detailed clinical models

- Goal
 - To build QA tool for cancer detailed clinical models
- Leveraging existing CIMI modeling efforts
 - CIMI full reference model
 - CIMI archetype modeling language (AML)
 - CTS2-based terminology binding
- Looking into HL7 FHIR modeling efforts
 - FHIR resources and profiles
 - FHIR terminology binding (value sets)
 - FHIR Ontology (HL7 RDF workgroup)
- Collaboration with clinical cancer research community
 - DeepPhe project (PI: Drs. Guergana Savova and Rebecca Crowley)
 - Mayo cancer registries (e.g., breast cancer, ovarian cancer)

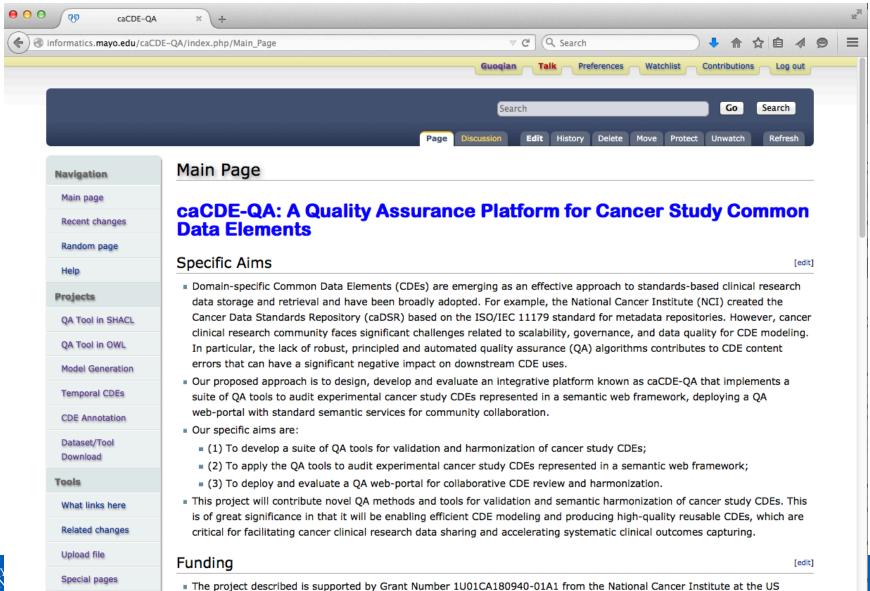


Collaboration study plan

- CDE discovery tool
 - Collaboration with Dr. Chunhua Weng from Columbia University
 - ClinicalTrials.gov
- CDE temporal pattern identification
 - Collaboration with Dr. Cui Tao from UTHealth
 - CNTRO Temporal Ontology
- QA tool using Shape Constraint Language
 - Dr. Tim Berners-Lee W3C/MIT
- CDISC / CIMI
- NCI caDSR team and TCGA project



http://informatics.mayo.edu/caCDE-QA/index.php/Main_Page





Questions & Discussion