Populating the i2b2 database with data from the electronic medical record:

Ontology-Based Form Data Integration for i2b2



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INTRODUCTION

Case report form (CRF) data capture systems and the electronic medical record (EMR) store precious data that is tempting to reuse in medical research [1]. However, while technical access to the systems' databases is easy, it is very difficult to reuse the data structured way, it is typically not encoded with standard terminologies like the UMLS or SNOMED. Furthermore, single medical concepts are often expressed by a combination of multiple CRF components (text input fields, checkboxes and radio data integration tools, the time-consuming work of remodeling data relationships for export is typically "buried" inside SQL scripts, which are difficult to maintain. What is even worse is that the tediously modeled knowledge is lost for further processing.

METHODS

We have developed a system [2] to extract, transform and load heterogeneous CRF/EMR data into an i2b2 research database [3]. In contrast to standard data integration approaches, our system is **RESULTS** designed to address the knowledge-related issues mentioned above. Instead of working at the database level, relationships between form components, data transformation and filtering rules are expressed in "higher-level" OWL ontologies [4]. This allows easier maintenance and the semantic interpretation of the created knowledge from a technical, administrative and medical perspective. The system works by describing both the source prostate cancer research network in Germany.

system (EMR) and the target dataset independently

with ontologies. The source ontology is an abstract,

technical description of the source system that

provides information on how to access the data

semantic features available in i2b2 (e.g. a mono- mapping was impossible, two mappings were hierarchy of medical concepts, data type, medication and lab value ranges - if applicable). The target ontology is also used to automatically create **DISCUSSION** the i2b2 ontology.

for research purposes. Although stored in a In order to link the source system (EMR) to the target system (i2b2) and to perform the export, a mapping ontology has to be created. This ontology contains manually created semantic relationships between the source and the target ontology, which can be processed by a software component to button groups). These relationships are, however, perform the data transfer. Relationships can either only apparent to humans and are not stored be "simple" one-to-one or "complex" mappings, with image on the right side. Intermediate nodes can also and standards for medical terminologies [5, 6, 7]. be cascaded into full expression trees to allow comprehensive data filtering and arbitrary data. By using i2b2 and extending it with our ontology automatically generated to perform the data transfer data from routine care. from the source systems to the target i2b2

We have evaluated the system's capabilities in a cross-institutional data sharing project between the university hospitals in Erlangen and Münster, Germany. Its objective was to establish a joint i2b2 research database with annotation data from prostate cancer patients for the *Deutsches* Prostatakarzinom Konsortium (DPKK), a large

The challenge was to map two distinct EMR systems to a central i2b2 database with a common dataset. As illustrated in the table below, we were able to records behind each form component, done in a map almost all of the data elements available in the machine-interpretable way. The target ontology cancer documentation of the two EMRs (Siemens describes the collection of medical concepts to be Soarian® in Erlangen and Agfa Orbis® in Münster)

made available in the i2b2 system, along with to the set of 166 medical concepts. Only one impractical to create with our current system.

The current implementation must still be considered prototypical as it offers room for improvements. For example, the system is currently designed to import data to i2b2 only, while being limited to the functionality and semantics of i2b2. We are confident, however, that we will be able to describe generic target database systems and that we will be able to better differentiate between metadata and facts data. This would allow us to derive and reuse explicitly in the database schema. Using standard full data filtering and transformation rules. The latter collected mapping information in other without ones are defined by using intermediate operation transforming actual data. We also hope to extend nodes, as illustrated with the ADD node on the the system to follow commonly accepted desiderata

> transformation. By sequentially processing these suite we feel confident that we have made a step nodes in a correct order, SQL statements are forward in efficiently accessing and reusing EMR

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Source **System** Gleason1 Gleason3 Gleason2 **PSA** PSA_value Mapping **Target Target Ontology Source Ontology** Ontology **Dataset i2b2** QuickMapp OntoExport **OntoEdit** (with Target Dataset)

Above: the Erlangen mapping concept with some of the developed tools.

No. of concepts directly mapped (simple 1:1) No. of concepts mapped through transformations No. of concepts not documented in source system No. of mappings not supported / impractical Generated SQL statements / execution time: Number of facts / patients in source table: Obtained facts / patients for DPKK i2b2:

Erlangen Hospital Münster Hospital 10 (4 with a workaround) 548 / ~15 seconds 284 / ~ 3 seconds 29,721,416 / 161,512 5,100 / 500 (test data) 3,686 / 155 2,585 / 487 (test data)

Table above: Result after mapping the two distinct EMRs to a set of 166 common medical concepts. Our approach has proven to work well, because for almost all of the target concepts a mapping could be created.

Münster Erlangen **EMR EMR**

> **DPKK Research Database**

i2b2

Background image: visualization of the 780 forms in the Erlangen EMR with a total of 42,000 (potentially) semantically distinct data elements (including versioning).

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