Ontology-Enriched Query Answering on Relational Databases

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The Thirty-Third Annual Conference on Innovative Applications of Artificial Intelligence (IAAI-21)

February 4-6, 2021

How it started...



A. Quamar, C. Lei, D. Miller, F. Özcan, J. Kreulen, R. Moore, V. Efthymiou. An Ontology-Based Conversation System for Knowledge Bases. *SIGMOD 2020*

How it started...

	IBM Micromedex Watson Assistant	- ×	
Well-forme Query (SQ	Headache × adult × ed L)	Clear what treats cephalalgia	
	I cannot answer that. "cephalalgia" is unknown!	Was this helpful?	logs
Relational Database	Type something		

Many of the unknown terms are contained in SNOMED CT and our DB actually has the data to answer such queries

By using SNOMED, we could also provide more complete answers to some queries

DL Ontologies

Based on Description Logic Body • a family of logic-based knowledge Structure representation formalisms Finding **Substance** decidable fragments of FOL Cell Head structure Pre-Finding Head Describe domains in terms of: embryonic site finding structure concepts (aka classes) Medicament • roles (aka binary relationships) Zygote Cephalalgia Antidote Frequent Dental Anesthetic Migraine Headache Headache

Main Challenges

- Identify and reuse only the parts of SNOMED CT that are relevant
 - we used existing tools from different AI communities
 - ontology creation from a DB, ontology matching, module extraction
 - we designed a flexible framework that goes beyond our use case
- Answer queries expressed over the vocabulary of SNOMED CT using our data
 - Two main approaches exist:
 - Materialization:
 - Materialize a universal solution (once) using the *chase* procedure from the data exchange community;
 - compute the certain answers on arbitrary conjunctive queries over the target schema using the materialized universal solution
 - Query Rewriting: Keep the original data, but rewrite every query when it comes before evaluating it

Background – Data Exchange



Data exchange setting $\mathbf{M} = (\mathbf{S}, \mathbf{T}, \boldsymbol{\Sigma}_{st}, \boldsymbol{\Sigma}_{t})$, where

Background – Examples of tgds and egds



$$\begin{split} \boldsymbol{\Sigma}_{st} &= st\text{-tgd} : \forall did, dn, dc(Drug(did, dn, dc) \rightarrow Medicament(did) \land DNames(did, dn)) \\ \boldsymbol{\Sigma}_{t} &= \begin{bmatrix} \texttt{-tegd} : \forall mid, x, y \ (DNames(mid, x) \land DNames(mid, y) \rightarrow x = y) \\ \texttt{-ttgd} : \forall mid, dn(DNames(mid, dn) \rightarrow \exists a(AIngred(mid, a)) \end{bmatrix} \end{split}$$

Background – Data Exchange



The certain answers of a query q over **T** on I, wrt a data exchange setting **M** are defined as:

cert(q,I,**M**) = \bigcap {q(J): J is a solution for I }

Fagin et al. 2005: if J is a universal solution for I w.r.t. **M**, then the certain answers of every conj. query q over **T** can be obtained by evaluating q on J and then removing all tuples containing null values

Problem: the chase may not always terminate!

Background – \mathcal{ELH} terminologies

Concepts constructs: $C \coloneqq A \mid \top \mid \exists r. C \mid C \sqcap D$

An \mathcal{FLH} terminology is a set of

- concept definitions $A \equiv C$,
- concept inclusions $A \sqsubseteq C$, and
- role inclusions $r \sqsubseteq s$

Contributions

Adoption of AI and data exchange methods and tools in real medical use case

Backing our use case with concrete theoretical guarantees

 $^\circ$ we define acyclic \mathcal{FLH}^{fdr} and show it is C-stratified

> the standard chase always terminates in polynomial time

A reference framework architecture for ontology-enriched query answering

available on github (<u>https://github.com/IBM/ontology-enriched-query-answering</u>)

Experimental evaluation showing the benefits of our framework

• more query answers by exploiting SNOMED CT as an external reference ontology

Framework Architecture

- Step 1: Ontology Creation
- Step 2: Matchings Generation
- Step 3: Module Extraction
- Step 4: Unifying the TBox
- Step 5: Query Answering via the Chase



Ontology Creation [Lei et al. 2018]



Matchings Generation



- Several methods tested, including SOTA in ontology matching (LogMap, AML) with unsatisfactory results
- Ended up providing matchings with manual inspection

Matchings C = {(Drug, Medicament)}

Module Extraction [Cuenca Grau et al. 2009]



Unifying the TBox

•To produce the unified TBox *T* :

- for every matching $(N_1, N_2) \in C$ rename every occurrence of N_1 in Ontology 1 with N_2
- $^{\circ}\,$ return the union of Ontology 1 (after renaming) and the S-module from Ontology 2



Expressivity of the Unified TBox (use case)

SNOMED CT belongs to the acyclic \mathcal{FLH} fragment of Description Logics

•Ontology generated from the DB falls under acyclic \mathcal{FL} , extended with domain and range restrictions, as well as functionality assertions

> The unified TBox can be expressed in acyclic *ELHfdr*

• \mathcal{ELH}^{fdr} is \mathcal{ELH} extended with domain and range restrictions, and limited functionality

- (simplified): no functional roles are allowed on the right-hand side of axioms
- Acyclicity intuition (proper definitions in the paper)
 - $\circ \mathcal{ELH}$ acyclicity: prevents a concept from directly or indirectly referring to (aka using) itself
 - *ELHfdr* acyclicity: need additional conditions to take care of domain & range restrictions and functionality
 - Example: $A \sqsubseteq \exists r$ $rng(r) \sqsubseteq A$ (acyclic under the \mathcal{ELH} acyclicity conditions, but results in infinite chase)

•Our schema exchange setting $\mathbf{M} = (\mathbf{S}, \mathbf{T}, \boldsymbol{\Sigma}_{st}, \boldsymbol{\Sigma}_{t})$:

- use the relational schema S of the input DB as the source schema S
- \circ use the unified TBox T as the target schema T

•Use the following rule to generate st-tgds from every relation R of S:

$$R(\underline{x_1}, \dots, x_n) \to R'(x_1) \wedge R'^{1,2}(x_1, x_2) \wedge \dots \wedge R'^{1,n}(x_1, x_n),$$

where x_1 is the primary key of R, and R', $R'^{1,j}$ are fresh relation names. If $(R, R'') \in C$, we replace $R'(x_1)$ above with $R''(x_1)$, i.e., we rename R as R''

Every functional role r gives rise to the t-egd:

 $r(x,y) \wedge r(x,z) \rightarrow y = z$

•Fact: for every \mathcal{FL} concept C, there is a conj. query $q_C(x)$ with a free variable x, s.t. $C(x) \equiv q_C(x)$

- **Case 1:** $q_{\mathcal{C}}(x) \coloneqq \exists \overline{y} \varphi_{\mathcal{C}}(\overline{y}, x)$, where \overline{y} is a non-empty tuple of variables
- Case 2: $q_C(x) \coloneqq A_1(x) \land \dots \land A_n(x)$, where $A_1(x), \dots, A_n(x)$ are concept names

The tgds arising from an \mathcal{ELH}^{fdr} terminology have one of the following seven types*:

1) $A(x) \rightarrow \exists \bar{y} \varphi_C(\bar{y}, x)$ (arises from $A \sqsubseteq C$, where C is of Case 1)2) $A(x) \rightarrow A_1(x) \land \dots \land A_n(x)$ (arises from $A \sqsubseteq C$, where C is of Case 2)3) $\varphi_C(\bar{y}, x) \rightarrow A(x)$ (arises from $C \sqsubseteq A$, where C is of Case 1)4) $A_1(x) \land \dots \land A_n(x) \rightarrow A(x)$ (arises from $C \sqsubseteq A$, where C is of Case 2)5) $r_1(x, y) \rightarrow r_2(x, y)$ (arises from $r_1 \sqsubseteq r_2$)6) $r(x, y) \rightarrow A(x)$ (arises from $dom(r) \sqsubseteq A$)7) $r(x, y) \rightarrow A(y)$ (arises from $rng(r) \sqsubseteq A$)

*We treat each axiom $A \equiv C$ as two inclusions $A \sqsubseteq C$ and $C \sqsubseteq A$

Chase Termination

Theorem: Let *T* be an acyclic \mathcal{ELH}^{fdr} terminology and let $\Sigma(T)$ be the associated set of tgds and egds. Then $\Sigma(T)$ is C-stratified.

[Meier, Schmidt, and Lausen 2009]: if a set Σ of tgds and egds is C-stratified, then, on every input database instance J, the standard chase w.r.t. Σ terminates in time bounded by a polynomial in the size of J.

Evaluation

Input DB (MDB): 62 relations of arities from 2 to 11, 158 FKs, 500k+ tuples, 62.3MB

•Ontologies:

- MDB ontology (Step 1): 49 concepts, 170 roles (all with domain & range), 156 functional
- SNOMED CT: 356k concepts, 119 roles (none is functional or with domain/range restrictions)
- 12 matchings identified (with manual inspection, after running LogMap, AML)
 - signature S given for module extraction contains 12 elements
- S-module in SNOMED CT: 35 concepts, 7 roles
- Unified TBox: 72 concepts, 177 roles, 156 functional, 170 with domain & range restrictions

Chase:

- Number of tgds: 62 st-tgds, 154 t-tgds, 156 t-egds
- Chase execution time: 1,676ms (870ms for st-tgds, 806ms for t-tgds and t-egds)
- Chase space overhead: 24% (62.3 MB used for the source instance; 77.5 MB used for the chase output)

Evaluation Results



Query-answering times ranged from 1ms (for q_{13} - q_{15}) to 576ms (for q_3), averaging **64ms**.

Thank you!

The source code of this work is publicly available:

https://github.com/IBM/ontology-enriched-query-answering