Theoretically Optimal Datalog Rewritings for OWL 2 QL Ontology-Mediated Queries

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Abstract. We show that, for *OWL 2 QL* ontology-mediated queries with (*i*) ontologies of bounded depth and conjunctive queries of bounded treewidth, (*ii*) ontologies of bounded depth and bounded-leaf tree-shaped conjunctive queries, and (*iii*) arbitrary ontologies and bounded-leaf tree-shaped conjunctive queries, one can construct and evaluate nonrecursive datalog rewritings by, respectively, LOGCFL, NL and LOGCFL algorithms, which matches the optimal combined complexity.

1 Introduction

Ontology-based data access (OBDA) via query rewriting [18] reduces the problem of finding answers to conjunctive queries (CQs) mediated by OWL 2 QL ontologies to standard database query answering. The question we are concerned with here is whether this reduction is optimal with respect to the combined complexity of query evaluation. Figure 1 (a) summarises what is known about the size of positive existential (PE), nonrecursive datalog (NDL) and first-order (FO) rewritings of OWL 2 QL ontologymediated queries (OMQs) depending on the existential depth of their ontologies and the shape of their CQs [13, 9, 12, 3]. Figure 1 (b) shows the combined complexity of OMQ evaluation for the corresponding classes of OMQs [5, 14, 12, 3]. Thus, we see, for example, that PE-rewritings for OMQs with ontologies of bounded depth and CQs of bounded treewidth can be of super-polynomial size, and so not evaluable in polynomial time, while the evaluation problem for these OMQs is decidable in LOGCFL \subseteq P. On the other hand, the OMQs in this class enjoy polynomial-size NDL-rewritings. However, these rewritings were defined using an argument from circuit complexity [3], and it has been unclear whether they can be constructed and evaluated in LOGCFL. The same concerns the class of OMQs with ontologies of bounded depth and bounded-leaf treeshaped queries, which can be evaluated in NL, and the class of OMQs with arbitrary ontologies and bounded-leaf tree-shaped queries, which can be evaluated in LOGCFL.

In this paper, we consider OMQs in these three classes and construct NDL-rewritings that are theoretically optimal in the sense that the rewriting and evaluation can be carried out by algorithms of optimal combined complexity, that is, from the complexity classes LOGCFL, NL and LOGCFL, respectively. Such algorithms are known to be space efficient and highly parallelisable. We compared our optimal NDL rewritings with those produced by query rewriting engines Clipper [8] and Rapid [6], using a sequence of OMQs with linear CQs and a fixed ontology of depth 1.

The full version is available at http://tinyurl.com/LogNDL-DL.



Fig. 1. (a) Size of OMQ rewritings; (b) combined complexity of OMQ evaluation.

2 Preliminaries

We give OWL 2 QL in the DL syntax with *individual names* a_i , *concept names* A_i , and *role names* P_i $(i \ge 1)$. *Roles* R and *basic concepts* B are defined by

$$R ::= P_i \mid P_i^-, \qquad B ::= A_i \mid \exists R.$$

A *TBox*, T, is a finite set of inclusions of the form

$$B_1 \sqsubseteq B_2, \qquad B_1 \sqcap B_2 \sqsubseteq \bot, \qquad R_1 \sqsubseteq R_2, \qquad R_1 \sqcap R_2 \sqsubseteq \bot.$$

An *ABox*, \mathcal{A} , is a finite set of atoms of the form $A_k(a_i)$ or $P_k(a_i, a_j)$. We denote by $\operatorname{ind}(\mathcal{A})$ the set of individual names in \mathcal{A} , and by $\mathbb{R}_{\mathcal{T}}$ the set of role names occurring in \mathcal{T} and their inverses. We use $A \equiv B$ for $A \sqsubseteq B$ and $B \sqsubseteq A$. The semantics for *OWL 2 QL* is defined in the usual way based on interpretations $\mathcal{I} = (\Delta^{\mathcal{I}}, \mathcal{I})$ [2].

For every role $R \in \mathbb{R}_{\mathcal{T}}$, we take a fresh concept name A_R and add $A_R \equiv \exists R$ to \mathcal{T} . The resulting TBox is said to be in *normal form*, and we assume, without loss of generality, that all our TBoxes are in normal form. The subsumption relation induced by \mathcal{T} is denoted by $\sqsubseteq_{\mathcal{T}}$: we write $S_1 \sqsubseteq_{\mathcal{T}} S_2$ if $\mathcal{T} \models S_1 \sqsubseteq S_2$, where S_1, S_2 are both either concepts or roles. We write $R(a, b) \in \mathcal{A}$ if $P(a, b) \in \mathcal{A}$ and R = P, or $P(b, a) \in \mathcal{A}$ and $R = P^-$; we also write $(\exists R)(a) \in \mathcal{A}$ if $R(a, b) \in \mathcal{A}$ for some b. An ABox \mathcal{A} is called *H*-complete with respect to \mathcal{T} in case

 $P(a,b) \in \mathcal{A}$ if $R(a,b) \in \mathcal{A}$, for roles P and R with $R \sqsubseteq_{\mathcal{T}} P$, $A(a) \in \mathcal{A}$ if $B(a) \in \mathcal{A}$, for a concept name A and basic concept B with $B \sqsubseteq_{\mathcal{T}} A$.

A conjunctive query (CQ) q(x) is a formula $\exists y \varphi(x, y)$, where φ is a conjunction of atoms $A_k(z_1)$ or $P_k(z_1, z_2)$ with $z_i \in x \cup y$ (without loss of generality, we assume that CQs do not contain constants). We denote by $\operatorname{var}(q)$ the variables $x \cup y$ of qand by $\operatorname{avar}(q)$ the answer variables x. An ontology-mediated query (OMQ) is a pair $Q(x) = (\mathcal{T}, q(x))$, where \mathcal{T} is a TBox and q(x) a CQ. A tuple a in $\operatorname{ind}(\mathcal{A})$ is a certain answer to Q(x) over an ABox \mathcal{A} if $\mathcal{I} \models q(a)$ for all models \mathcal{I} of \mathcal{T} and \mathcal{A} ; in this case we write $\mathcal{T}, \mathcal{A} \models q(a)$. If $x = \emptyset$, then a certain answer to Q over \mathcal{A} is 'yes' if $\mathcal{T}, \mathcal{A} \models q$ and 'no' otherwise. We often regard a CQ q as the set of its atoms. Every consistent OWL 2 QL knowledge base (KB) $(\mathcal{T}, \mathcal{A})$ has a canonical model $\mathcal{C}_{\mathcal{T},\mathcal{A}}$ with the property that $\mathcal{T}, \mathcal{A} \models q(a)$ iff $\mathcal{C}_{\mathcal{T},\mathcal{A}} \models q(a)$, for any CQ q and any a in ind(\mathcal{A}). Thus, CQ answering in OWL 2 QL amounts to finding a homomorphism from the given CQ into the canonical model. Informally, $\mathcal{C}_{\mathcal{T},\mathcal{A}}$ is obtained from \mathcal{A} by repeatedly applying the axioms in \mathcal{T} , introducing fresh elements as needed to serve as witnesses for the existential quantifiers. According to the standard construction (cf. [16]), the domain $\Delta^{\mathcal{C}_{\mathcal{T},\mathcal{A}}}$ of $\mathcal{C}_{\mathcal{T},\mathcal{A}}$ consists of words of the form $aR_1 \dots R_n$ ($n \ge 0$) with $a \in ind(\mathcal{A})$ and $R_1 \dots R_n \in \mathbb{R}^*_{\mathcal{T}}$ such that (i) $\mathcal{T}, \mathcal{A} \models \exists R_1(a)$ and (ii) $\exists R_i^- \sqsubseteq_{\mathcal{T}} \exists R_{i+1}$ and $R_i^- \nvDash_{\mathcal{T}} R_{i+1}$, for $1 \le i < n$. We let $\mathbf{W}_{\mathcal{T}}$ consist of all words $R_1 \dots R_n \in \mathbb{R}^*_{\mathcal{T}}$ satisfying condition (ii). A TBox \mathcal{T} is of depth ω if $\mathbf{W}_{\mathcal{T}}$ is infinite, and of depth $d < \omega$, if d is the maximum length of the words in $\mathbf{W}_{\mathcal{T}}$.

A datalog program, Π , is a finite set of Horn clauses $\forall z \ (\gamma_0 \leftarrow \gamma_1 \land \cdots \land \gamma_m)$, where each γ_i is an atom S(y) with $y \subseteq z$ or an equality (z = z') with $z, z' \in z$. (As usual, when writing clauses, we omit $\forall z$.) The atom γ_0 is the *head* of the clause, and $\gamma_1, \ldots, \gamma_m$ its *body*. All variables in the head must also occur in the body, and = can only occur in the body. The predicates in the heads of clauses in Π are *IDB predicates*, the rest (including =) *EDB predicates*. A predicate *S depends* on *S'* in Π if Π has a clause with *S* in the head and *S'* in the body; Π is a *nonrecursive datalog* (NDL) *program* if the (directed) *dependence graph* of the dependence relation is acyclic.

An NDL query is a pair $(\Pi, G(\mathbf{x}))$, where Π is an NDL program and $G(\mathbf{x})$ a predicate. A tuple \mathbf{a} in ind (\mathcal{A}) is an answer to $(\Pi, G(\mathbf{x}))$ over an ABox \mathcal{A} if $G(\mathbf{a})$ holds in the first-order model with domain ind (\mathcal{A}) obtained by closing \mathcal{A} under the clauses in Π ; in this case we write $\Pi, \mathcal{A} \models G(\mathbf{a})$. The problem of checking whether \mathbf{a} is an answer to $(\Pi, G(\mathbf{x}))$ over \mathcal{A} is called the query evaluation problem. The arity of Π is the maximal arity, $r(\Pi)$, of predicates in Π . The depth of $(\Pi, G(\mathbf{x}))$ is the length, $d(\Pi, G)$, of the longest directed path in the dependence graph for Π starting from G. NDL queries are equivalent if they have exactly the same answers over any ABox.

An NDL query $(\Pi, G(\mathbf{x}))$ is an NDL-rewriting of an OMQ $\mathbf{Q}(\mathbf{x}) = (\mathcal{T}, \mathbf{q}(\mathbf{x}))$ over H-complete ABoxes in case $\mathcal{T}, \mathcal{A} \models \mathbf{q}(\mathbf{a})$ iff $\Pi, \mathcal{A} \models G(\mathbf{a})$, for any H-complete ABox \mathcal{A} and any tuple \mathbf{a} in ind (\mathcal{A}) . Rewritings over arbitrary ABoxes are defined by dropping the condition that the ABoxes are H-complete. Let $(\Pi, G(\mathbf{x}))$ be an NDL-rewriting of $\mathbf{Q}(\mathbf{x})$ over H-complete ABoxes. Denote by Π^* the result of replacing each predicate Sin Π with a fresh predicate S^* and adding the clauses $A^*(x) \leftarrow B'(x)$, for $B \sqsubseteq_{\mathcal{T}} A$, and $P^*(x, y) \leftarrow R'(x, y)$, for $R \sqsubseteq_{\mathcal{T}} P$, where B'(x) and R'(x, y) are the obvious first-order translations of B and R (for example, $B'(x) = \exists y R(x, y)$ if $B = \exists R$). It is easy to see that $(\Pi^*, G^*(\mathbf{x}))$ is an NDL-rewriting of $\mathbf{Q}(\mathbf{x})$ over arbitrary ABoxes.

It is well-known [4] that, without loss of generality, we can only consider NDL-rewritings of OMQs $(\mathcal{T}, q(x))$ over ABoxes \mathcal{A} that are *consistent* with \mathcal{T} .

We call an NDL query $(\Pi, G(x_1, \ldots, x_n))$ ordered if each of its IDB predicates S comes with fixed variables x_{i_1}, \ldots, x_{i_k} $(1 \le i_1 < \cdots < i_k \le n)$, called the *parameters* of S, such that (i) every occurrence of S in Π is of the form $S(y_1, \ldots, y_m, x_{i_1}, \ldots, x_{i_k})$, (ii) the x_i are the parameters of G, and (iii) if x' are all the parameters in the body of a clause, then the head has x' among its parameters. The width $w(\Pi, G)$ of a ordered (Π, G) is the maximal number of non-parameter variables in a clause of Π . All our NDL-rewritings in Secs. 4–6 are ordered, so we now only consider ordered NDL queries.

3 NL and LOGCFL Fragments of Nonrecursive Datalog

In this section, we identify two classes of (ordered) NDL queries with the evaluation problem in the complexity classes NL and LOGCFL for combined complexity. Recall [1] that an NDL program is called *linear* if the body of its every clause contains at most one IDB predicate (remember that equality is an EDB predicate).

Theorem 1. Fix some w > 0. The combined complexity of evaluating linear NDL queries of width at most w is NL-complete.

Proof. Let $(\Pi, G(\mathbf{x}))$ be a linear NDL query. Deciding whether $\Pi, \mathcal{A} \models G(\mathbf{a})$ is reducible to finding a path to $G(\mathbf{a})$ from a certain set X in the grounding graph $\mathfrak{G}(\Pi, \mathcal{A}, \mathbf{a})$ constructed as follows. The vertices of the graph are the ground atoms obtained by taking an IDB atom from Π , replacing each of its parameters by the corresponding constant from \mathbf{a} , and replacing each non-parameter variable by some constant from \mathcal{A} . The graph has an edge from $S(\mathbf{c})$ to $S'(\mathbf{c}')$ iff the grounding of Π contains a clause $S'(\mathbf{c}') \leftarrow S(\mathbf{c}) \wedge E_1(\mathbf{e}_1) \wedge \cdots \wedge E_k(\mathbf{e}_k)$ with $E_j(\mathbf{e}_j) \in \mathcal{A}$, for $1 \leq j \leq k$ (we assume that $(c = c) \in \mathcal{A}$). The set X consists of all vertices $S(\mathbf{c})$ with IDB predicates S being of in-degree 0 in the dependency graph of Π for which there is a clause $S(\mathbf{c}) \leftarrow E_1(\mathbf{e}_1) \wedge \cdots \wedge E_k(\mathbf{e}_k)$ in the grounding of Π with $E_j(\mathbf{e}_j) \in \mathcal{A}$ ($1 \leq j \leq k$). Bounding the width of (Π, G) ensures that $\mathfrak{G}(\Pi, \mathcal{A}, \mathbf{a})$ is of polynomial size and can be constructed by a deterministic Turing machine with separate input, write-once output and logarithmic-size working tapes.

The transformation of NDL-rewritings over H-complete ABoxes into rewritings for arbitrary ABoxes in Section 2 does not preserve linearity. However, we can still show that it suffices to consider the H-complete case:

Lemma 2. For any fixed w > 0, there is an L^{NL}-transducer that, given a linear NDLrewriting of an OMQ Q(x) over H-complete ABoxes that is of width at most w, computes a linear NDL-rewriting of Q(x) over arbitrary ABoxes whose width is at most w + 1.

The complexity class LOGCFL can be defined in terms of *nondeterministic auxiliary pushdown automata* (NAuxPDAs) [7], which are nondeterministic Turing machines with an additional work tape constrained to operate as a pushdown store. Sudborough [19] proved that LOGCFL coincides with the class of problems that are solved by NAuxPDAs running in logarithmic space and polynomial time (the space on the pushdown tape is not subject to the logarithmic bound).

We call an NDL query (Π, G) skinny if the body of any clause in Π has ≤ 2 atoms.

Lemma 3. For any skinny NDL query $(\Pi, G(\mathbf{x}))$ and ABox \mathcal{A} , query evaluation can be done by an NAuxPDA in space $\log |\Pi| + w(\Pi, G) \cdot \log |\mathcal{A}|$ and time $2^{O(\mathsf{d}(\Pi,G))}$.

Proof. Let $\Pi^{a}_{\mathcal{A}}$ be the set of ground clauses obtained by first replacing each parameter in Π by the corresponding constant from a, and then performing the standard grounding of Π using the constants from \mathcal{A} . Consider the monotone Boolean circuit $C(\Pi, \mathcal{A}, a)$ constructed as follows. The output of $C(\Pi, \mathcal{A}, a)$ is G(a). For every atom γ occurring in the head of a clause in $\Pi^{a}_{\mathcal{A}}$, we take an OR-gate whose output is γ and inputs are the bodies of the clauses with head γ ; for every such body, we take an AND-gate whose inputs are the atoms in the body. We set an input gate γ to 1 iff $\gamma \in \mathcal{A}$. Clearly, $C(\Pi, \mathcal{A}, a)$ is a semi-unbounded fan-in circuit (where OR-gates have arbitrarily many inputs, and ANDgates two inputs) with $O(|\Pi| \cdot |\mathcal{A}|^{w(\Pi,G)})$ gates and depth $O(d(\Pi,G))$. It is known that the nonuniform analog of LOGCFL can be defined using families of semi-unbounded fanin circuits of polynomial size and logarithmic depth. Moreover, there is an algorithm that, given such a circuit C, computes the output using an NAuxPDA in logarithmic space in the size of C and exponential time in the depth of C [20, pp. 392–397]. Observing that $C(\Pi, \mathcal{A}, a)$ can be computed by a deterministic logspace Turing machine, we conclude that the query evaluation problem can be solved by an NAuxPDA in space $\log |\Pi| + w(\Pi, G) \cdot \log |\mathcal{A}|$ and time $2^{O(d(\Pi,G))}$.

A function ν from the predicate names in Π to \mathbb{N} is a *weight function for* an NDLquery $(\Pi, G(\boldsymbol{x}))$ if $\nu(P) > 0$, for any IDB P in Π , and $\nu(P) \ge \nu(Q_1) + \cdots + \nu(Q_n)$, for any $P(\boldsymbol{z}) \leftarrow Q_1(\boldsymbol{z}_1) \land \cdots \land Q_n(\boldsymbol{z}_n)$ in Π .

Lemma 4. If $(\Pi, G(\mathbf{x}))$ has a weight function ν , then it is equivalent to a skinny NDL query $(\Pi', G(\mathbf{x}))$ such that $|\Pi'|$ is polynomial in $|\Pi|$, $d(\Pi', G) \leq d(\Pi, G) + \log \nu(G)$ and $w(\Pi', G) \leq w(\Pi, G)$.

Proof. The proof is by induction on $d(\Pi, G)$. If $d(\Pi, G) = 0$, we take $\Pi' = \Pi$. Suppose Π contains a clause ψ of the form $G(\mathbf{z}) \leftarrow P_1(\mathbf{z}_1) \land \cdots \land P_k(\mathbf{z}_k)$ and, for each $1 \leq j \leq k$, we have an NDL query (Π'_{P_j}, P_j) which is equivalent to (Π, P_j) and such that

$$\mathsf{d}(\Pi'_{P_i}, P_j) \le \mathsf{d}(\Pi_{P_i}, P_j) + \log \nu(P_j) \le \mathsf{d}(\Pi, G) - 1 + \log \nu(P_j).$$
(1)

We construct the Huffman tree [11] for the alphabet $\{1, \ldots, k\}$, where the frequency of j is $\nu(P_j)/\nu(G)$ (by definition, $\nu(G) > 0$). The Huffman tree is binary and has kleaves, denoted $1, \ldots, k$, and k - 1 internal nodes (including the root, g), and the length of the path from g to any leaf j at most $\lceil \log(\nu(G)/\nu(P_j)) \rceil$. For each internal node vof the tree (but the root), we take a predicate $P_v(z_v)$, where z_v is the union of z_u for all descendants u of v; for the root g, we take $P_g(z_g) = G(z)$. Let Π'_{ψ} be the extension of the union of Π'_{P_j} , for $1 \le j \le k$, with clauses $P_v(z_v) \leftarrow P_{u_1}(z_{u_1}) \land P_{u_2}(z_{u_2})$, for each v with immediate successors u_1 and u_2 . The number of the new clauses is k - 1. Consider the NDL query $(\Pi'_{u_i}, G(z))$. By (1), we have:

$$d(\Pi'_{\psi}, G) \le \max_{j} \{ \lceil \log(\nu(G)/\nu(P_{j})) \rceil + d(\Pi'_{P_{j}}, P_{j}) \} \le \max_{j} \{ \log(\nu(G)/\nu(P_{j})) + d(\Pi, G) + \log\nu(P_{j}) \} = \log\nu(G) + d(\Pi, G).$$

Let Π' be the result of applying this transformation to each clause in Π with head G(z). It is readily seen that (Π', G) is as required; in particular, $|\Pi'| = O(|\Pi|^2)$.

Theorem 5. Fix $c \ge 1$, $w \ge 1$ and a polynomial p. Query evaluation for NDL queries $(\Pi, G(\mathbf{x}))$ with a weight function ν such that $\nu(G) \le p(|\Pi|)$, $w(\Pi, G) \le w$ and $d(\Pi, G) \le c \log \nu(G)$ is in LOGCFL for combined complexity.

Proof. By Lemma 4, (Π, G) is equivalent to a skinny NDL query (Π', G') with $|\Pi'|$ polynomial in $|\Pi|$, $w(\Pi', G) \le w$, and $d(\Pi', G') \le (c+1) \log \nu(G)$. By Lemma 3, query evaluation for (Π', G') over \mathcal{A} is solved by an NAuxPDA in space $\log |\Pi'| + w(\Pi', G) \cdot \log |\mathcal{A}| = O(\log |\Pi| + \log |\mathcal{A}|)$ and time $2^{O(d(\Pi', G'))} \le 2^{O(\log \nu(G))} = (\nu(G))^{O(1)} \le p'(|\Pi|)$, for some polynomial p'.

Corollary 6. Suppose there is an algorithm that, given any OMQ Q(x) from some class C, constructs its NDL-rewriting $(\Pi, G(x))$ over H-complete ABoxes having a weight function ν with $\nu(G) \leq |\mathbf{Q}|$ and $d(\Pi, G) \leq c \log \nu(G)$, and such that $w(\Pi, G) \leq w$ and $|\mathbf{Q}| \leq |\Pi| \leq p(|\mathbf{Q}|)$, for some fixed constants c, w and polynomial p. Then the evaluation problem for the NDL-rewritings $(\Pi^*, G^*(x))$ of the OMQs in C over arbitrary ABoxes (defined in Section 2) is in LOGCFL for combined complexity.

4 Bounded Treewidth CQs and Bounded-Depth TBoxes

With every CQ q, we associate its *Gaifman graph* \mathcal{G} whose vertices are the variables of q and edges are the pairs $\{u, v\}$ such that $P(u, v) \in q$, for some P. We call q tree-shaped if \mathcal{G} is a tree; q is connected if the graph \mathcal{G} is connected. A tree decomposition of an undirected graph $\mathcal{G} = (V, E)$ is a pair (T, λ) , where T is an (undirected) tree and λ a function from the set of nodes of T to 2^V such that the following conditions hold:

- for every $v \in V$, there exists a node t with $v \in \lambda(t)$;
- for every $e \in E$, there exists a node t with $e \subseteq \lambda(t)$;
- for every $v \in V$, the nodes $\{t \mid v \in \lambda(t)\}$ induce a connected subtree of T.

We call the set $\lambda(t) \subseteq V$ a *bag for* t. The *width* of (T, λ) is $\max_{t \in T} |\lambda(t)| - 1$. The *treewidth of a graph* \mathcal{G} is the minimum width over all tree decompositions of \mathcal{G} . The *treewidth of a CQ* is the treewidth of its Gaifman graph.

Its natural tree decomposition of treewidth 1 is based on the chain T of 7 vertices, which are represented as bags as follows:



Fix a connected CQ q(x) and a tree decomposition (T, λ) of its Gaifman graph $\mathcal{G} = (V, E)$. Let D be a subtree of T. The *size* of D is the number of nodes in it. We call a node t of D boundary if T has an edge $\{t, t'\}$ with $t' \notin D$, and let the *degree* deg(D) of D be the number of its boundary nodes. Note that T itself is the only subtree of T of degree 0. We say that a node t splits D into subtrees D_1, \ldots, D_k if the D_i partition D without t: each node of D except t belongs to exactly one D_i .

Lemma 8 ([3]). Let D be a subtree of T of size m > 1. If deg(D) = 2, then there is a node t splitting D into subtrees of size $\leq m/2$ and degree ≤ 2 and, possibly, one subtree of size < m - 1 and degree 1. If deg(D) ≤ 1 , then there is t splitting D into subtrees of size $\leq m/2$ and degree ≤ 2 .

In Example 7, t splits T into T_1 and T_2 depicted below:



We define recursively a set sub(T) of subtrees of T, a binary relation \prec on sub(T)and a function σ on sub(T) indicating the splitting node. We begin by adding T to sub(T). Take $D \in sub(T)$ that has not been split yet. If D is of size 1 then let $\sigma(D)$ be the only node of D. Otherwise, by Lemma 8, we find a node t in D that splits it into D_1, \ldots, D_k . We set $\sigma(D) = t$ and, for each $1 \le i \le k$, add D_i to sub(T) and set $D_i \prec D$; then, we apply the procedure recursively to each of D_1, \ldots, D_k . In Example 7 with t splitting T, we have $\sigma(T) = t$, $T_1 \prec T$ and $T_2 \prec T$.

For each $D \in sub(T)$, we recursively define a set of atoms q_D by taking

$$\boldsymbol{q}_D = \{ S(\boldsymbol{v}) \in \boldsymbol{q} \mid \boldsymbol{v} \subseteq \lambda(\sigma(D)) \} \cup \bigcup_{D' \prec D} \boldsymbol{q}_{D'}.$$

By the definition of tree decomposition, $q_T = q$. Denote by x_D the subset of $\operatorname{avar}(q)$ that occur in q_D . In our running example, $x_T = \{x_0, x_7\}$, $x_{T_1} = \{x_0\}$ and $x_{T_2} = \{x_7\}$. Denote by ∂D the union of all $\lambda(t) \cap \lambda(t')$ for a boundary node t of D and its unique neighbour t' in T outside D. In our example, $\partial T = \emptyset$, $\partial T_1 = \{x_3\}$ and $\partial T_2 = \{x_4\}$.

Let \mathcal{T} be a TBox of finite depth k. A type is a partial map w from V to $W_{\mathcal{T}}$; its domain is denoted by dom(w). By ε we denote the unique partial type with dom $(\varepsilon) = \emptyset$. We use types to represent how variables are mapped into $\mathcal{C}_{\mathcal{T},\mathcal{A}}$, with w(u) = w indicating that u is mapped to an element of the form aw (for some $a \in ind(\mathcal{A})$), and with $w(u) = \varepsilon$ that u is mapped to an ABox individual. We say that a type w is *compatible* with a bag t if, for all $u, v \in \lambda(t) \cap dom(w)$, we have

- if $v \in \operatorname{avar}(q)$, then $w(v) = \varepsilon$;
- if $A(v) \in q$, then either $w(v) = \varepsilon$ or w(v) = wR with $\exists R^{-} \sqsubseteq_{\mathcal{T}} A$;
- if $R(v, u) \in q$, then $w(v) = w(u) = \varepsilon$, or w(u) = w(v)R' with $R' \sqsubseteq_{\mathcal{T}} R$, or w(v) = w(u)R' with $R' \sqsubseteq_{\mathcal{T}} R^-$.

In the sequel, we abuse notation and use sets of variables in place of sequences assuming that they are ordered in some (fixed) way. For example, we use x_D for a tuple of variables in the set x_D (ordered in some way). Also, given a tuple a in $ind(\mathcal{A})$ of length $|x_D|$ and $x \in x_D$, we write a(x) to refer to the element of a that corresponds to x (that is, to the component of the tuple with the same index).

Let Π_Q be an NDL program that, for any $D \in sub(T)$, types w and s such that $dom(w) = \partial D$, $dom(s) = \lambda(\sigma(D))$, s is compatible with $\sigma(D)$ and agrees with w on their common domain, contains the clause

$$G_D^{\boldsymbol{w}}(\partial D, \boldsymbol{x}_D) \leftarrow \operatorname{At}^{\boldsymbol{s}} \wedge \bigwedge_{D' \prec D} G_{D'}^{(\boldsymbol{s} \cup \boldsymbol{w}) \restriction \partial D'}(\partial D', \boldsymbol{x}_{D'}),$$
(2)

where x_D are the parameters of predicate G_D^w , $(s \cup w) \upharpoonright \partial D'$ is the restriction¹ of the union $s \cup w$ of s and w to $\partial D'$, and At^s is defined as follows:

$$\mathsf{At}^{\boldsymbol{s}} = \bigwedge_{\substack{A(u) \in \boldsymbol{q} \\ \boldsymbol{s}(u) = \varepsilon}} A(u) \land \bigwedge_{\substack{R(u,v) \in \boldsymbol{q} \\ \boldsymbol{s}(u) = \varepsilon}} R(u,v) \land \bigwedge_{\substack{R(u,v) \in \boldsymbol{q} \\ \boldsymbol{s}(u) \neq \varepsilon \text{ or } \boldsymbol{s}(v) \neq \varepsilon}} \bigwedge_{\substack{R(u,v) \in \boldsymbol{q} \\ \boldsymbol{s}(v) \neq \varepsilon \text{ or } \boldsymbol{s}(v) \neq \varepsilon}} \bigwedge_{\substack{s(u) = Sw' \\ \text{ for some } w'}} A_{S}(u). (3)$$

The first two conjunctions in At^s ensure that atoms all of whose variables are assigned ε are present in the ABox. The third conjunction ensures that if one of the variables in a role atom is not mapped to ε , then the images of the variables share the same initial individual. Finally, atoms in the final conjunction ensure that if a variable is to be mapped to aSw', then the individual a satisfies $\exists S$ (so aSw' is part of the domain of $C_{\mathcal{T},\mathcal{A}}$).

Example 9. Now we fix an ontology \mathcal{T} with the following axioms:

$$A \equiv \exists P, \quad P \sqsubseteq S, \quad P \sqsubseteq R^-, \qquad B \equiv \exists Q, \quad Q \sqsubseteq R, \quad Q \sqsubseteq S^-.$$

Since $\lambda(t) = \{x_3, x_4\}$, there are only three types compatible with t:

$$s_1 \colon x_3 \mapsto \varepsilon, x_4 \mapsto \varepsilon, \qquad s_2 \colon x_3 \mapsto P, x_4 \mapsto \varepsilon \quad \text{and} \quad s_3 \colon x_3 \mapsto \varepsilon, x_4 \mapsto Q.$$

So, $At^{s_1} = R(x_3, x_4)$, $At^{s_2} = A(x_3) \land (x_3 = x_4)$, $At^{s_3} = B(x_4) \land (x_3 = x_4)$. Thus, predicate G_T^{ε} is defined by the following clauses, for s_1 , s_2 and s_3 , respectively:

$$\begin{aligned} G_{T}^{\varepsilon}(x_{0}, x_{7}) &\leftarrow G_{T_{1}}^{x_{3} \mapsto \varepsilon}(x_{3}, x_{0}) \wedge R(x_{3}, x_{4}) \wedge G_{T_{2}}^{x_{4} \mapsto \varepsilon}(x_{4}, x_{7}), \\ G_{T}^{\varepsilon}(x_{0}, x_{7}) &\leftarrow G_{T_{1}}^{x_{3} \mapsto P}(x_{3}, x_{0}) \wedge A(x_{3}) \wedge (x_{3} = x_{4}) \wedge G_{T_{2}}^{x_{4} \mapsto \varepsilon}(x_{4}, x_{7}), \\ G_{T}^{\varepsilon}(x_{0}, x_{7}) &\leftarrow G_{T_{1}}^{x_{3} \mapsto \varepsilon}(x_{3}, x_{0}) \wedge B(x_{4}) \wedge (x_{3} = x_{4}) \wedge G_{T_{2}}^{x_{4} \mapsto Q}(x_{4}, x_{7}). \end{aligned}$$

By induction on \prec on sub(T), we show that $(\Pi_Q, G_T^{\varepsilon})$ is a rewriting of Q(x).

Lemma 10. For any ABox \mathcal{A} , any $D \in \operatorname{sub}(T)$, any type w with dom $(w) = \partial D$, any $b \in \operatorname{ind}(\mathcal{A})^{|\partial D|}$ and $a \in \operatorname{ind}(\mathcal{A})^{|\boldsymbol{x}_D|}$, we have $\Pi_{\boldsymbol{Q}}, \mathcal{A} \models G_D^{\boldsymbol{w}}(\boldsymbol{b}, \boldsymbol{a})$ iff there is a homomorphism $h: \boldsymbol{q}_D \to \mathcal{C}_{\mathcal{T},\mathcal{A}}$ such that

$$h(x) = \boldsymbol{a}(x), \text{ for } x \in \boldsymbol{x}_D, \text{ and } h(v) = \boldsymbol{b}(v)\boldsymbol{w}(v), \text{ for } v \in \partial D.$$

Fix now k and t, and consider the class of OMQs $Q(x) = (\mathcal{T}, q(x))$ with \mathcal{T} of depth $\leq k$ and q of treewidth $\leq t$. Let T be a tree decomposition of q of treewidth $\leq t$. We take the following weight function: $\nu(G_D^w) = |D|$. Clearly, $\nu(G_T^\varepsilon) \leq |Q|$. By Lemma 8, $d(\Pi_Q, G_T^\varepsilon) \leq 2 \log |T| = 2 \log \nu(G_T^\varepsilon) \leq 2 \log |Q|$. Since $|\operatorname{sub}(T)| \leq |T|^2$ and there are at most $|\mathcal{T}|^{2tk}$ options for w, there are polynomially many predicates G_D^w and so, Π_Q is of polynomial size. Thus, by Corollary 6, the obtained NDL-rewriting over arbitrary ABoxes can be evaluated in LOGCFL. Finally, we note that a tree decomposition of treewidth $\leq t$ can be computed using an $\mathsf{L}^{\mathsf{LOGCFL}}$ -transducer [10], and so the NDL-rewriting can also be constructed by an $\mathsf{L}^{\mathsf{LOGCFL}}$ -transducer.

¹ By construction, dom $(\boldsymbol{s} \cup \boldsymbol{w})$ covers $\partial D'$, and so the domain of $(\boldsymbol{s} \cup \boldsymbol{w}) \upharpoonright \partial D'$ is $\partial D'$.

5 Bounded-Leaf CQs and Bounded-Depth TBoxes

We next consider OMQs with tree-shaped CQs in which both the depth of the ontology and the number of leaves in the CQ are bounded. Let \mathcal{T} be a TBox of finite depth k, and let q(x) be a tree-shaped CQ with at most ℓ leaves. Fix one of the variables of q as root, and let M be the maximal distance to a leaf from the root. For $n \leq M$, let z^n denote the set of all variables of q at distance n from the root; clearly, $|z^n| \leq \ell$. We call the z^n slices of q and observe that they satisfy the following: for every $R(u, v) \in q$ with $u \neq v$, there exists $0 \leq n < M$ such that either $u \in z^n$ and $v \in z^{n+1}$ or $u \in z^{n+1}$ and $v \in z^n$. For $0 \leq n \leq M$, we denote by $q_n(z_{\exists}^n, x^n)$ the query consisting of all atoms S(u) of q such that $u \subseteq \bigcup_{n \leq m \leq M} z^m$, where $x^n = var(q_n) \cap x$ and $z_{\exists}^n = z^n \setminus x$.

By type of a slice z^n , we mean a total map w from z^n to W_T . Analogously to Section 4, we define what it means for a type (or pair of types) to be compatible with a slice (pair of adjacent slices). We call w locally compatible with z^n if for every $z \in z^n$:

- if $z \in \operatorname{avar}(q)$, then $w(z) = \varepsilon$;
- if $A(z) \in q$, then either $w(z) = \varepsilon$ or w(z) = wR with $\exists R^- \sqsubseteq_T A$;
- if $R(z, z) \in q$, then $w(z) = \varepsilon$.

If w, s are types for z^n and z^{n+1} respectively, then we call (w, s) compatible with (z^n, z^{n+1}) if w is locally compatible with z^n , s is locally compatible with z^{n+1} , and for every atom $R(z^n, z^{n+1}) \in q$, one of the following holds: (i) $w(z^n) = s(z^{n+1}) = \varepsilon$, (ii) $s(z^{n+1}) = w(z^n)R'$ with $R' \sqsubseteq_T R$, or (iii) $w(z^n) = s(z^{n+1})R'$ with $R' \sqsubseteq_T R^-$.

Consider the NDL program Π'_{Q} defined as follows. For every $0 \le n < M$ and every pair of types (w, s) that is compatible with (z^n, z^{n+1}) , we include the clause:

$$P_n^{\boldsymbol{w}}(\boldsymbol{z}_{\exists}^n, \boldsymbol{x}^n) \leftarrow \mathsf{At}^{\boldsymbol{w} \cup \boldsymbol{s}}(\boldsymbol{z}^n, \boldsymbol{z}^{n+1}) \wedge P_{n+1}^{\boldsymbol{s}}(\boldsymbol{z}_{\exists}^{n+1}, \boldsymbol{x}^{n+1})$$

where x^n are the parameters of P_n^w and $At^{w \cup s}(z^n, z^{n+1})$ is the conjunction of atoms (3), as defined in Section 4, for the union $w \cup s$ of types w and s. For every type w locally compatible with z^M , we include the clause:

$$P_M^{\boldsymbol{w}}(\boldsymbol{z}_\exists^M, \boldsymbol{x}^M) \leftarrow \mathsf{At}^{\boldsymbol{w}}(\boldsymbol{z}^M).$$

(Recall that z^M is a disjoint union of z_{\exists}^M and x^M .) We use G, with parameters x, as the goal predicate and include $G(x) \leftarrow P_0^w(z^0, x)$ for every predicate $P_0^w(z^0, x^0)$ occurring in the head of one of the preceding clauses.

The following lemma (which is proved by induction) is the key step in showing that $(\Pi'_{Q}, G(\boldsymbol{x}))$ is a rewriting of $(\mathcal{T}, \boldsymbol{q})$ over H-complete ABoxes:

Lemma 11. For any *H*-complete ABox \mathcal{A} , any $0 \le n \le M$, any predicate $P_n^{\boldsymbol{w}}$, any $\boldsymbol{b} \in \operatorname{ind}(\mathcal{A})^{|\boldsymbol{z}_{\exists}^n|}$ and any $\boldsymbol{a} \in \operatorname{ind}(\mathcal{A})^{|\boldsymbol{x}^n|}$, we have $\Pi'_{\boldsymbol{Q}}, \mathcal{A} \models P_n^{\boldsymbol{w}}(\boldsymbol{b}, \boldsymbol{a})$ iff there is a homomorphism $h: \boldsymbol{q}_n \to \mathcal{C}_{\mathcal{T}, \mathcal{A}}$ such that

$$h(x) = \boldsymbol{a}(x), \text{ for } x \in \boldsymbol{x}^n, \text{ and } h(z) = \boldsymbol{b}(z)\boldsymbol{w}(z), \text{ for } z \in \boldsymbol{z}_{\exists}^n.$$
 (4)

It should be clear that Π'_Q is a linear NDL program of width at most 2ℓ . Moreover, when ℓ and k are bounded by fixed constants, it takes only logarithmic space to store a type w, which allows us to show that Π'_Q can be computed by an L^{NL}-transducer. We can apply Lemma 2 to obtain an NDL rewriting for arbitrary ABoxes, and then use Theorem 1 to conclude that the resulting program can be evaluated in NL.

6 Bounded-Leaf CQs and Arbitrary TBoxes

For OMQs with bounded-leaf CQs and ontologies of unbounded depth, our rewriting utilises the notion of tree witness [15]. Let $Q(x) = (\mathcal{T}, q(x))$ with $q(x) = \exists y \varphi(x, y)$. For a pair $t = (t_r, t_i)$ of disjoint sets of variables in q, with $t_i \subseteq y$ and $t_i \neq \emptyset$, set

$$\boldsymbol{q}_{\mathsf{t}} = \left\{ S(\boldsymbol{z}) \in \boldsymbol{q} \mid \boldsymbol{z} \subseteq \mathsf{t}_{\mathsf{r}} \cup \mathsf{t}_{\mathsf{i}} \text{ and } \boldsymbol{z} \not\subseteq \mathsf{t}_{\mathsf{r}} \right\}$$

If q_t is a minimal subset of q for which there is a homomorphism $h: q_t \to C_T^{A_R(a)}$ such that $t_r = h^{-1}(a)$ and q_t contains every atom of q with at least one variable from t_i , then we call $t = (t_r, t_i)$ a *tree witness for* Q generated by R. Note that the same tree witness $t = (t_r, t_i)$ can be generated by different roles R. By $q \setminus t$ we denote the query obtained from q by removing the atoms in q_t and having $x \cup t_r$ as answer variables, and for every $v \in var(q)$, we let $\mathsf{TW}_Q(v)$ denote the set of all tree witnesses t for Q such that $v \in t_i$.

The logarithmic depth NDL-rewriting for bounded-leaf queries and ontologies of unbounded depth is based upon the following observation.

Lemma 12. Every tree T of size m has a node splitting it into subtrees of size $\leq m/2$.

We will use repeated applications of this lemma to decompose the input CQ into smaller and smaller subqueries. Formally, for every tree-shaped CQ q, we use v_q to denote a vertex in the Gaifman graph \mathcal{G} of q that satisfies the condition of Lemma 12. Then, starting from an OMQ $Q_0 = (\mathcal{T}, q_0(x_0))$, we define SQ_{Q0} as the smallest set of queries that contains $q_0(x_0)$ and is such that for every $q(z) \in SQ_{Q_0}$ with |var(q)| > 1, the following queries also belong to SQ_{Q0}:

- for every u_i that is adjacent to v_q in G: the query q_i(z_i) consisting of all role atoms linking v_q and u_i, as well as all atoms whose variables cannot reach v_q in G without passing by u_i, and with z_i equal to (z ∩ var(q_i)) ∪ {v_q};
- for every t ∈ TW_{Q0}(v_q) with t_r ≠ Ø: the queries q^t₁(z^t₁),...,q^t_{mt}(z^t_{mt}) corresponding to the connected components of q \ t, with z^t_i equal to var(q^t_i) ∩ avar(q \ t).

The NDL program Π''_{Q_0} uses IDB predicates P_q , for $q \in SQ_{Q_0}$, with arity |avar(q)| and parameters $var(q) \cap x$. For each $q(z) \in SQ_{Q_0}$ with |var(q)| > 1, we include the clause

$$P_{\boldsymbol{q}}(\boldsymbol{z}) \quad \leftarrow \bigwedge_{A(v_{\boldsymbol{q}}) \in \boldsymbol{q}} A(v_{\boldsymbol{q}}) \quad \wedge \bigwedge_{R(v_{\boldsymbol{q}},v_{\boldsymbol{q}}) \in \boldsymbol{q}} R(v_{\boldsymbol{q}},v_{\boldsymbol{q}}) \quad \wedge \bigwedge_{1 \leq i \leq n} P_{\boldsymbol{q}_i}(\boldsymbol{z}_i),$$

where $q_1(z_1), \ldots, q_n(z_n)$ are the subqueries induced by the neighbours of v_q in \mathcal{G} . We also include, for every $t \in \mathsf{TW}_Q(v_q)$ with $t_r \neq \emptyset$ and role R generating t, the clause

$$P_{\boldsymbol{q}}(\boldsymbol{z}) \quad \leftarrow \bigwedge_{u,u' \in \mathsf{t}_{\mathsf{r}}} (u = u') \quad \land \bigwedge_{u \in \mathsf{t}_{\mathsf{r}}} A_{R}(u) \quad \land \bigwedge_{1 \leq i \leq m_{\mathsf{t}}} P_{\boldsymbol{q}_{i}^{\mathsf{t}}}(\boldsymbol{z}_{i}^{\mathsf{t}})$$

where $q_1^t, \ldots, q_{m_t}^t$ are the connected components of $q \setminus t$. For every $q \in SQ_{Q_0}$ with |var(q)| = 1, we include the clause $P_q(z) \leftarrow q$. Finally, if q_0 is Boolean, we include clauses $P_{q_0} \leftarrow A(x)$ for all concept names A such that $\mathcal{T}, \{A(a)\} \models q_0$.

The program $\Pi_{Q_0}^{\prime\prime}$ is inspired by a similar construction from [12]. By adapting results from the latter paper, we can show that $(\Pi_{Q_0}^{\prime\prime}, P_{q_0}(x))$ is indeed a rewriting:

Lemma 13. For any tree-shaped OMQ $Q_0(x) = (\mathcal{T}, q_0(x))$, any $q(z) \in SQ_{Q_0}$, any *H*-complete ABox \mathcal{A} , and any tuple a in ind (\mathcal{A}) , $\Pi''_{Q_0}, \mathcal{A} \models P_q(a)$ iff there exists a homomorphism $h: q \to C_{\mathcal{T},\mathcal{A}}$ such that h(z) = a.

Now fix $\ell > 1$, and consider the class of OMQs $Q(x) = (\mathcal{T}, q(x))$ with tree-shaped q(x) having at most ℓ leaves. The size of Π''_Q is polynomially bounded in |Q|, since bounded-leaf CQs have polynomially many tree witnesses and also polynomially many tree-shaped subCQs. It is readily seen that the function ν defined by setting $\nu(P_{q'}) = |q'|$ is a weight function for (Π''_Q, P_q) such that $\nu(P_q) \leq |Q|$. Moreover, by Lemma 12, $d(\Pi, G) \leq \log \nu(P_q) + 1$. We can thus apply Corollary 6 to conclude that the obtained NDL-rewritings can be evaluated in LOGCFL. Finally, we note that since the number of leaves is bounded, it is in NL to decide whether a vertex satisfies the conditions of Lemma 12, and it is in LOGCFL to decide whether $\mathcal{T}, \{A(a)\} \models q_0$ [3] or whether a (logspace) representation of a possible tree witness is indeed a tree witness. This allows us to show that (Π''_Q, P_q) can be generated by an L^{LOGCFL}-transducer.

7 Conclusions

As shown above, for three important classes of OMQs, NDL-rewritings can be constructed and evaluated by theoretically optimal NL and LOGCFL algorithms. To see whether these rewritings are viable in practice, we generated three sequences of OMQs with the ontology from Example 9 and linear CQs of up to 15 atoms as in Example 7. We compared our NL and LOGCFL rewritings from Sections 5 and 4 (called LIN and LOG) with those produced by Clipper [8] and Rapid [6]. The barcharts below show the number of clauses in the rewritings over H-complete ABoxes. While LIN and LOG grow linearly (in accord with theory), Clipper and Rapid failed to produce rewritings for longer CQs.



We evaluated the rewritings over a few randomly generated ABoxes using off-the-shelf datalog engine RDFox [17]. The experiments (see the full version) show that our rewritings are usually executed faster than Clipper's and Rapid's when the number of answers is relatively small ($\leq 10^4$); for queries with $\geq 10^6$ answers, the execution times are comparable. The version of RDFox we used did not seem to take advantage of the structure of the NL/LOGCFL rewritings, and it would be interesting to see whether their nonrecursiveness and parallelisability can be utilised to produce efficient execution plans.

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