Algorithmic study of Kernel Contraction in \mathcal{EL}

by

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Abstract

Kernel contraction is an interesting problem that can be used as a step towards belief revision. Kernels were introduced to determine why a given belief is accepted by the knowledge base. The aim of using kernels is to invalidate the reasons why that given belief is accepted, and hence rejecting it. In this study we will explain what kernels are and how we can use them to contract beliefs in an \mathcal{EL} TBox. We will introduce some algorithms that performs belief contraction in different ways. We will also see how to determine the rationality of belief contraction by the rules introduced in the AGM framework.

Keywords: KEYWORDS

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Chapter 1

Introduction

Changing one's mind is a process that happens very frequently as part of one's daily routine. Seeing the sunrise in the morning, one would intuitively change his mind and believe that it is not dark any more; that would logically imply that one gives up the current belief that it is dawn time and replace it with a new belief that it is sunrise time. This is an example of belief revision. Revision is a kind of change in which the new belief ("it is sunrise") is conflicting with the current state of mind (believing that "it is dawn").

Mind changing can be thought of as the manipulation of beliefs according to perception. When humans perceive any change in their world, they change their knowledge accordingly. So changing the knowledge (or beliefs¹) of an agent² can be seen as what we informally call changing the mind of the agent. One example of belief change is **Revision**, which involves changing the knowledge base in order to add a conflicting belief. This can actually be broken down into two processes: changing the knowledge base to account for the conflict, and adding the new belief.

The first process involves removing the beliefs that are causing that conflict. This process is called **Belief Contraction**. The second process involves adding the new belief and expanding the knowledge base accordingly. This process is called **Belief Expansion**. Contraction is the type of change we are mainly concerned with in this study. Kernel contraction is one approach to contraction that performs by getting all kernels and removing some statements from each of them.

¹Throughout this study we will be using the terms "knowledge" and "beliefs" interchangeably, although they are not exactly the same. Knowledge is usually assumed to be a special kind of beliefs. However, for convenience, when we use the word "knowledge" we will be referring to "belief."

²The word "agent" here means humans, computers, or any thing that has knowledge base and perception.

1.1 Kernel contraction by example

The following chapters will explain kernel contraction in more details, but it is helpful to get a brief idea on what it is before moving on. A kernel is a minimal set of beliefs that imply a certain belief. If our knowledge contains the following beliefs:

- We are in Canada
- The trees are green
- The temperature is 30 Celsius,

and we know that whenever these three statements are true then "it is summer," we can imply that it is indeed summer time. In this case, the three beliefs together form a kernel that implies "it is summer." That kernel can be used for contraction purpose. If we want to give up the belief "it is summer," we can remove a belief from the kernel so that the remaining two cannot be enough to imply that it is summer. In some other cases we can have more than one kernel, and the same rules can still apply: remove a belief from each kernel in order to perform kernel contraction. The beliefs used in this example are quite subjective, but the aim is to give a brief and simple example. The following chapters will talk more formally about kernel contraction.

1.2 The language

The knowledge of an agent can be represented as a set of beliefs. An agent is assumed to believe in A if A is a member of its belief set³. In this study, the language used to represent belief sets is Description Logic. Description Logic is a family of formalisms that are used to represent knowledge. They use concepts to represent classes of individuals and roles to represent relationships between them. They have different expressive powers and different reasoning mechanism with different complexities. They vary according to the set of logical operators they use. Here we use \mathcal{EL} , which is a member of the Description Logic family.

1.3 Scope of this study

We aim at studying implementations of kernel contraction for knowledge bases represented in \mathcal{EL} description logic. We will look at some basic implementations as well as some advanced ones. Our focus is on investigating the algorithmic aspects of those different approaches. The time complexity will be taken into considerations, and more importantly the optimality of the solutions found. Optimality can be measures in different ways, among

³Belief sets will be explained in the next chapter

them is what makes more sense as a solution to humans and what is closer to solutions humans generally prefer.

The starting point of optimality seeking in this study will be from a syntactic point of view based on the hitting set problem. The hitting set problem is simply the problem of finding the smallest set of elements that intersect with given set of sets of elements (this will be explained in more details later). We adopt the hitting set approach because what we have is a set of kernels (sets of elements) that we need to remove one from each. By doing this we hope to achieve syntactical optimality that is only measured by the size of the solution: the smaller the better.

One of the significant contributions of this study is the investigation of algorithms that perform kernel contraction achieving semantic optimality based on \mathcal{EL} semantics. These approaches exploit the structure of \mathcal{EL} knowledge bases to find most reasonable solutions. Both approaches that seek semantic and syntactic optimality can be combined together to achieve better solutions. We will study two heuristics that can be used to make decisions on which beliefs to remove based on the meaning of the logical operators and the structure of the language. These heuristics can be useful in cases where other approaches (such as the minimum hitting set algorithm) result in more than one solution with the same size, to select the solution that is more reasonable based on the meaning of the beliefs. Approaches that only consider the size of the solution do not always make a meaningful choice on which beliefs to remove. So we can then use the heuristics as a tie-breaker step when we get more than one solution.

We start the study by discussing the basic types of belief change, but before that, we build a ground for them by defining the framework that was introduced by Gardenfors. We define epistemic states and attitudes that will help in understanding the mechanics of belief change. We then explain some postulates introduced in the AGM framework; those postulates are considered rationality rules for belief change operations. Then we explain what description logic is and what logical operators are used. In that discussion, we focus on the most relevant variation to this study, which is \mathcal{EL} .

After that, we will go over some belief contraction techniques and show how syntactic optimality can be achieved, and what the cost can be (in terms of time complexity.) We will talk more formally about contraction using kernels. Our new approach to kernel contraction is by reducing the problem of contraction to a graph problem. We will give a brief introduction to this new technique and will discuss it in details with a few different examples. We will use the algorithm for network flow problems to produce a solution to kernel contraction that will be guaranteed to be smallest. This algorithm cannot be used for any \mathcal{EL} knowledge base; it only works on knowledge bases in some certain settings. This will be discussed in more details later, and the examples will show the cases in which the graph technique is useful and the cases in which it is not. Then, we will turn to the approaches that consider the semantics of the language in finding best solutions to the kernel

contraction problem. These heuristics are called Localization and Specificity. Later, we will see what they are and how useful they can be.

Chapter 2

Background

Learning is one of the most interesting functions of the human mind. If we think of the human memory as a storage of beliefs, we may think of learning as the manipulation of these beliefs. In that sense, learning something new would correspond to the addition of new beliefs. But learning is not only about new things. We sometimes learn things that contradict other things we learned before. In that sense, learning could be thought of as the update or revision of beliefs.

In this chapter, we aim at explaining the context of the ideas developed in this study, and giving some background of the main building blocks of our work. We will discuss three different ways of changing beliefs and we will use the AGM framework[12] to define and explain these changes. After explaining the AGM rules that govern our approaches to rational belief change, we will explain a Description Logic called \mathcal{EL} , which will be used in this study as the knowledge representation language.

2.1 Belief change

The AGM framework is named after Alchourron, Gardenfors, and Makinson, the three scientists who invented it in the 1980s. Since that time, AGM has been the most adopted framework for belief change. In [7], Gardenfors came up with some very useful notations to describe the change of beliefs, epistemic states and attitudes. We can think of the epistemic states as the states of mind (from the beliefs perspective), where epistemic states of an agent are defined by the epistemic attitudes agents have towards the concepts they know about. In this case, changing one attitude towards one concept implies a new epistemic state. Therefore, belief change is simply transitions between different epistemic states. Before moving on to explaining the AGM paradigm, it is useful to shed some light on the notions of epistemic states and attitudes.

2.1.1 Epistemic states

As the word "change" suggests, our concern is about transitions between epistemic states. One can think of an epistemic state as the state of beliefs of an agent (or a human), and of the change as the move from a given state to a new state. The beliefs of an agent can be modelled -as the epistemological theory suggested in [7]- as a set of propositions (or beliefs), with some assumptions on the attitudes towards each of the propositions (will be discussed in more details in 2.1.2.) Those beliefs are not meant to be psychological propositions expressing beliefs in human mind; they are epistemological idealizations of psychological propositions that a human mind can believe or reject. It is reasonable to consider that every agent should always seek an equilibrium state, where the epistemic state is consistent; and if some concepts are contradicting, the agent ought to revise its beliefs to reach a consistent state.

In this study, we use the words "belief sets" and "epistemic state", and they are not exactly the same, although they are related; A **Belief Set** is the set of all things that an agent believe in, while an **Epistemic State** is the complete state of the agent's epistemic attitudes towards all propositions, including the ones that the agent does not believe in. Depending on the epistemological model we follow, we might consider propositions that are not in the belief set to be rejected, or we might assume that the agent neither believes in them nor rejects them.

Another important notion is the *belief base*, which we use to refer to the (possibly incomplete) set of *explicit* beliefs of an agent. It is different from a belief set because we typically assume that belief sets are close under logical consequence, while belief bases are not. In that sense, we say that the belief set includes some *implicit* beliefs that follow from some explicit ones.

The AGM paradigm will use the notion of belief sets. Later in 2.2, we will discuss Description Logics (DLs) as formalisms to express knowledge, and start using DL *concepts* to represent belief bases. All following discussions about *contraction* algorithms will use such representation.

2.1.2 Epistemic attitudes

A belief of an agent can be interpreted according to the concepts in an epistemic state. Suppose the concept C is defined as:

$$C = \text{It's Monday}$$

If C exists in the belief set of an agent, we say that the agent believes in C (believes that today is Monday), or accepts C [7], which also means that accepting C is part of the epistemic state of the agent. However, accepting a belief is not the only attitude an agent can have towards a concept. An agent can also reject a concept C if the negation of that

concept is in the belief set (i.e. It is not Monday.) In that epistemic state, we say that the agent rejects C. It is also possible that an agent stays undetermined (or ignorant) about a concept if neither the concept nor its negation is in the belief set (e.g. an agent has no idea whether today is Monday or not.) There can be more attitudes if we consider other models of beliefs such as the probabilistic model, where an agent might have different levels of beliefs. But for the scope of this study we are only interested in the three attitudes: accept, reject and $no\ clue$.

2.1.3 Basic types of change

The AGM framework defines three types of belief change: revision, expansion, and contraction. If one sees a flying penguin, it introduces a new belief (penguins might fly), which contradicts another belief people these days have: penguins can't fly. In this case, just accepting the new belief will make the belief system of the agent (In this case, a human) inconsistent, since two beliefs are contradicting. So revising the old beliefs, and possibly removing the belief about penguins can't fly, is the rational action humans tend to take. This is referred to as belief revision.

In the case when the new belief does not contradict any of the old beliefs, the new belief is accepted without having to remove any of the old beliefs. This is called belief expansion. When a belief is removed, or given up, without accepting a new belief, we call this process belief contraction. A good example of belief contraction is to give up a belief for the sake of argument, to reach a common ground with an opponent, and argue based on this ground.

Before going further into details of those types of change of belief, some important notions need to be explained.

belief set We represent the beliefs of an agent by a set of sentences, where $\{A\}$ represents a belief set of only one belief A. So an agent who has $\{A\}$ as his belief set only believes in A, and is ignorant everything else. The belief set is closed under deductive inference; if K is a belief set, then $K \vdash \alpha$ if and only if $\alpha \in K$.

Given a belief set K and a sentence A, we say $K \vdash A$ if and only if A is a consequence of the belief set K. We refer to the set of all consequences of K by Cn(K). Since K is a belief set, then it is closed under deduction. Hence, K = Cn(K), where Cn(K) is the deductive closure of K. For any belief set K and a sentence A, we can represent the three aforementioned epistemic attitudes as follows [7]:

- If $A \in K$, A is accepted and -A is rejected.
- If $-A \in K$, A is rejected, and -A is accepted.
- $A \notin K$ and $-A \notin K$ means that A is undetermined.

We can think of belief change about A as a change of the attitude towards A. Because we have three attitudes, we have six possible changes. However, because of the symmetry of some pairs of change we only consider three types of change:

- 1. Change from being undetermined to either accepting or rejecting A (can only be obtained through expansion).
- 2. Change from accepting to rejecting and vice versa (revision).
- 3. Change from accepting or rejecting A to being undetermined (contraction)[7].

Belief Expansion

As part of the learning process and gaining knowledge an agent expands its belief set by accepting new beliefs. If K is a belief set and A is a statement, then K_A^+ is the new belief set resulting from accepting A. So the expansion function + is a function that takes a belief set K and a statement A and returns another belief set K_A^+ .

Belief Contraction

Contraction is different from expansion. Expansion by A is the change from the state of not accepting nor rejecting it to the state of accepting it, while contraction is the change from accepting or rejecting A to being undetermined of it. We denote the belief set resulting from contracting the belief set K with respect to sentence A by K_A^- . Since contraction is the most relevant change to the scope of this study, we need to discuss some of the rationality criteria (or postulates) of such change. We consider the contraction function — as a function from a set of beliefs K and a sentence A to a new set of beliefs K_A^- as in [7] and [12].

In the AGM framework, the following eight postulates ((K - 1) to (K - 8)) are introduced as basis for ensuring the rationality of contraction functions (more details can be found in [12] and [7]):

(K - 1) For any sentence A and any belief set K, K_A^- is a belief set.

 K_A^- is obtained by removing some beliefs. So

$$(\mathbf{K} - \mathbf{2}) K_A^- \subseteq K.$$

If A is not already in the belief set, the contraction should not change the belief set.

$$(K-3)$$
 If $A \notin K$, then $K_A^- = K$.

Unless A is logically valid, A should not be in the belief set after contraction.

$$(K - 4)$$
 If not $\vdash A$, then $A \notin K_A^-$.

The main idea of contraction is to give up some belief (maybe temporarily). If we need to give B up we can remove it from the belief set, and make sure it is not derivable from the remaining beliefs. To do that we need to look for statements that together entail B and remove at least one of them. This will be elaborated more in chapter 3 when we discuss kernels. So we do not just remove A from K, but remove the minimum number of sentences that would entail A. We remove the "minimum" number of sentences because it is better to give up as least as possible (minimum change will be explained in more details in chapter 3). Also, while performing expansion with A we compute all possible consequences that can be reached after accepting A. Therefore, all beliefs are recoverable by expansion after contraction:

$$(K - 5)$$
 If $A \in K$, then $K \subseteq (K_A^-)_A^+$.

If A and B are equivalent sentences, then contracting K for the belief A should result in the same belief set as contracting it for B.

$$(K - 6)$$
 If $\vdash A \leftrightarrow B$, then $K_A^- = K_B^-$.

If we contract K by A&B, we should contract K by either A or B, but nothing else.

$$(K - 7) K_A^- \cap K_B^- \subseteq K_{A\&B}^-.$$

Motivated by the concept of minimal change, the last postulate states that if A was removed while contracting K by A&B, then at least A was given up – possibly along with B.

(K - 8) If
$$A \notin K_{A\&B}^-$$
, then $K_{A\&B}^- \subseteq K_A^-$.

Belief Revision

Revision is different from expansion and contraction, as it is considered to be non-monotonic. New beliefs are added with -possibly- some of the old beliefs removed. Revision takes place when the attitude is changed from accepting to rejecting or from rejecting to accepting. We can think of revision as a combination of both contraction and expansion. The revision function * is a function from a belief set K and a sentence A to a belief set K_A^* . If $-A \in K$, then $K_A^* = (K_{-A}^-)_A^+$.

In the next section we look at a class of languages called Description Logic that is being used these days to represent knowledge. We use it as a model and implement contraction in a DL-based knowledge bases.

2.2 Description logic

One of the earliest approaches to representing knowledge is using logic. Logic has been suitable for general-purpose applications. Another approach is what is so-called semantic

networks, which is a graph (or a network)-based approach. Semantic networks use graphs to represent knowledge[4], where nodes represent *concepts* and edges represent *relationships* between them.

Description Logic (DL) comes as an evolution of semantic networks to a logic-based approach with some new flavours. DL is a class of logics based on -as the name suggests-describing sets of individuals using concepts and describing the relationships between them using roles. Usually, two types of knowledge need to be represented: intensional and extensional knowledge. Intensional knowledge is general knowledge about a problem or a domain, while extensional knowledge is knowledge about a specific problem instance. For this purpose, a knowledge base¹ is composed of two main components: Terminological Box, which we will refer to by TBox, and Assertions Box, which we refer to by ABox.

To get an idea about what DL looks like, we look at two categories of symbols: logical and non-logical symbols. **Non-logical** symbols include the following:

Concepts that are similar to category nouns (e.g. *Human*, *Mother*, *Animal*, *College*, etc.). They are used to represent classes (or sets) of individuals. So we can use the concept *Animal* to refer to the set of animals, and the concept *Man* to represent the class of men.

Roles that are like relational nouns (e.g. MotherOf, HeightOf, etc.). They can be used to specify attributes of concepts.

Constants and they are used to represent individuals, and they are similar to proper nouns, e.g. Adam, Sally, etc.

Concepts and roles can be used with the help of some logical symbols to construct complex expressions. **Logical** symbols include the following:

- □, □, ¬ used as propositional constructors (conjunction, disjunction, and negation respectively).
- \forall , \exists used for restriction and quantification.
- \top , \bot (\top represents the set of all individuals, while \bot represents the empty set of individuals).

Logical and non-logical symbols can be used together to construct complex expressions. To get a sense of how to build complex expressions, suppose C and D are concepts, and R is a role:

- $C \sqcap D$, $C \sqcup D$, and $\neg C$ are concepts (also called concept expressions).
- $\forall R.C$, and $\exists R.\top$ are concepts (or concept expressions).

¹By the word *Knowledge base* we refer to a belief base − a set of sentences that represent the belief of an agent.

2.2.1 ABox

As we said at the beginning of this section, to represent knowledge we usually look at intensional and extensional knowledge. DL knowledge bases usually consist of two main components: TBox and ABox. **ABoxes** are built from assertions (extensional knowledge) about a specific problem or domain, e.g.

Girl(Sally) may represent the fact that Sally is a Girl and, FatherOf(Adam, Sally) may represent the fact that Adam is the Father Of Sally

where Sally and Adam are constants, Girl is a concept name, and FatherOf is a role name. Using such representation, ABoxes can be built to represent assertions about a problem or a domain.

2.2.2 TBox

Suppose that Human is a concept that refers to all humans, and Male is the concept that refer to all male beings. We can use conjunction (\Box) in

$$Human \sqcap Male$$

to represent all individuals that are both humans and male beings. We can also define a new concepts Man to represent those individuals by introducing the definition

$$Man \doteq Human \sqcap Male$$

So every man has to satisfy that definition. The TBox is composed of concept definitions and General Concept Inclusion rules (GCIs). GCIs are weaker than definitions; the rule

$$Man \sqsubseteq Human$$

states that Man is subsumed by Human, which means that every man is a human². Every definition can be safely broken down into two GCIs. For example, the definition

$$Man \doteq Human \sqcap Male$$

can be broken down into the two GCIs:

$$Man \sqsubseteq Human \sqcap Male$$

 $Human \sqcap Male \sqsubseteq Man$

Tboxes are used to represent general knowledge (intensional knowledge) about a class of problems or domains, using axioms (or terminologies). A typical TBox is composed of definitions and GCIs – sometimes for convenience all definitions are broken down into GCIs. The following is an example of a -somewhat incomplete- DL TBox:

 $^{^{2}}$ or more precisely, every member of the set represented by Man is also a member of the set represented by Human.

- 1. $Man \doteq Human \sqcap Male$.
- 2. $Woman \sqsubseteq Human \sqcap \neg Man$.
- 3. $Father \doteq Man \sqcap \exists ParentOf. \top$
- 4. $Mother \doteq Woman \sqcap \exists ParentOf. \top$
- 5. $FatherWithoutSon \doteq Father \sqcap \forall ParentOf. \neg Man.$
- 6. $Parent \doteq Father \sqcup Mother$
- 7. $GrandFather \sqsubseteq Father \sqcap \exists ParentOf.Parent$

where they can be interpreted such that #1 defines Man to be a human male, #2 states that every Woman is a human and not a man, #3 defines Father to be a man that is a parent of something (since \top includes everything), #4 defines Mother similarly, #5 defines a FatherWithoutSon to be a father which every individual that is in a "ParentOf" relationship with is not a man, #6 defines a Parent to be a father or a mother, and #7 states that every GrandFather is a father and in a "ParentOf" relation ship with a parent.

Along with some other symbols and constructors, these symbols are the building blocks of DL formalisms. There are many members of the DL family, that vary in their expressivity power and the complexity of their reasoning algorithms. Each member of the family includes a subset of the DL symbols, and is uniquely identified by the containment of those symbols. In the following section we discuss a very famous member of the DL family, \mathcal{EL} , and use it as a knowledge representation language for the rest of the study.

2.2.3 \mathcal{EL} language

One of the DLs that recently became famous and attracted much attention is \mathcal{EL} . \mathcal{EL} is a light-weight Description Logic, though it is used in some well-known ontologies such as SNOMED CT, which is a medical ontology that contains around 380000 concepts[3]. \mathcal{EL} only contains a subset of the concept constructors that we discussed in the previous section, and they are:

- The conjunction symbol \sqcap
- Existential restriction \exists
- The top concept \top

One of the advantages of \mathcal{EL} , besides being simple and easy to use, is its polynomial-time subsumption algorithm. The subsumption problem, which is the most important problem in \mathcal{EL} , is actually a classification problem. The subsumption algorithm classifies the TBox depending on the subsumption relation expressed by \sqsubseteq . It is basically checking whether a

specific subsumption relation holds or not (e.g. whether $C \sqsubseteq D$ holds or not) in a given knowledge base. Checking whether a subsumption relation holds or not is also checking whether the concept (or concept expression) on the left-hand side of the subsumption relation is subsumed by the concept on the right-hand side.

It is helpful to sketch the subsumption algorithm before moving on into the core of this study, as it will be used later on. But we will use an example to see how the algorithm can be applied. Let's assume we have a knowledge base TBox \mathcal{T} that consists of only two subsumption relations:

$$\mathcal{T} = \{ \text{ Haddock} \sqsubseteq \text{Fish, Fish} \sqsubseteq \text{Animal.} \}$$

The first one states that every Haddock is a Fish, and the second one states that every Fish is an Animal. Now, given that \mathcal{T} , we need to check that every Haddock is an Animal:

$$Haddock \sqsubseteq Animal.$$

This is not explicitly stated in \mathcal{T} , but we can imply it using the subsumption algorithm. The algorithm proceeds in four steps:

- 1. Normalize the TBox.
- 2. Translate the normalized TBox into a graph.
- 3. Complete the graph using completion rules.
- 4. Read off the subsumption relationships from the normalized graph.

We can follow the four steps and apply them one by one to our example and get the solution that we need.

Normalize the TBox

We can only call a TBox normalized if all the GCIs it contains are of any of the following forms:

- $A_1 \sqcap A_2 \sqsubseteq B$
- $A \sqsubseteq \exists r.B$
- $\exists r.A \sqsubseteq B$

where A, A_1 , A_2 , and B are arbitrary concepts, and r is an arbitrary role. A TBox can be normalized in polynomial time. $A \sqcap \top$ is equivalent to A. So, the normalized version of the \mathcal{T} look like:

$$\{ \text{ Haddock} \sqcap \top \sqsubseteq \text{ Fish, Fish } \sqcap \top \sqsubseteq \text{ Animal } \}.$$

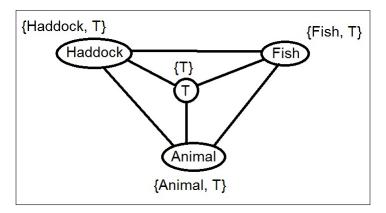


Figure 2.1: Graph transformation of the TBox

Translate the normalized TBox into graph

The second step is to translate the TBox into a graph. This is done by creating a node that corresponds to each concept in the TBox (including \top), and have an edge between every pair of nodes. We use S to denote the set of nodes' labels, and R to denote the set of roles' labels. S will contain the subsumers of a node, e.g. S(A) contains B if $A \sqsubseteq B$. Likewise, R(A,B) contains r if $A \sqsubseteq \exists r.B$. Initially, the set S starts by containing the nodes and \top , i.e. $S(Haddock) = \{Haddock, \top\}$, $S(Fish) = \{Fish, \top\}$, and $S(Animal) = \{Animal, \top\}$. The set R will initially be empty. The graph can be visualised in Figure 2.1. Translating the TBox into a graph can be done in polynomial time.

Complete the graph using completion rules

The following three rules are then used to extend the sets S and R:

- If $(A_1 \sqcap A_2 \sqsubseteq B) \in \mathcal{T}$ and $A_1, A_2 \in S(A)$ then add B to S(A)
- If $(A_1 \sqsubseteq \exists r.B) \in \mathcal{T}$ and $A_1 \in S(A)$ then add r to R(A,B)
- If $(\exists r.B_1 \sqsubseteq A_1) \in \mathcal{T}$ and $B_1 \in S(B), r \in S(A, B)$ then add A_1 to S(A)

If we apply the completion rules to \mathcal{T} , S will become as follows:

- $S(Animal) = \{Animal, \top\}$
- $S(Fish) = \{Fish, \top, Animal\}$
- $S(Haddock) = \{Haddock, \top, Fish, Animal\}$

The new graph (after the completion rules are applied) can be visualised in Figure 2.2.

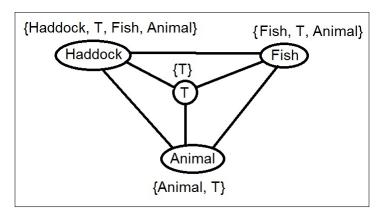


Figure 2.2: Completed graph of the TBox

Read off the subsumption relationships

Now that the graph is complete, we can look at the sets S and R and determine whether the subsumption relationship in question holds or not. Since that subsumption relationship was $Haddock \sqsubseteq Animal$, we can look at S(Haddock) and see if it contains Animal or not, since the set S(Haddock) contains all subsumers of Haddock. So, indeed, $Haddock \sqsubseteq Animal$ holds. This brief example hopefully shows a simple application of the subsumption algorithm of \mathcal{EL} . We didn't discuss the complexity of the algorithm in details, but it is enough to say it is polynomial in the size of the TBox. More details on the algorithm can be found in [5] and [3].

Now that we had a quick introduction to the basic types of belief change and the framework of AGM, as well as the \mathcal{EL} language and reasoning algorithm, we can start talking about the core of this study, which is kernel contraction. Throughout this study, we only consider DL knowledge bases represented in \mathcal{EL} . In the next chapter we look at an implementation of belief contraction using kernels as introduced in [11]. We will discuss some basic and general approaches as well as some more sophisticated ones. Then we will consider a language-specific approach to exploit the structure of \mathcal{EL} and discuss heuristics that will provide more meaningful solutions.

Chapter 3

Kernel contraction

Belief contraction is the process of removing beliefs from belief sets. It can be done by either of two ways. The first way is selecting one of the largest subsets of the belief set that do not logically imply the belief to be removed. This is called the *remainder set* approach. It works by finding the largest subsets of the belief set, such that those subsets do not imply the removed belief. For example, the knowledge base:

$$\{a, a \rightarrow b, b, c\}$$

has two remainder sets that do not logically imply b:

$$\{a,c\},\{a\to b,c\}$$

And they are the largest such subsets. This means that there are no larger subset of the original set that do not imply b. So we can choose one of them to be the new (contracted) belief set.

The other approach is done by incising all the minimal subsets that logically imply the removed belief; by "incising" we mean removing a member from the set. This second approach is the one we are going to focus on in this study. If a set K of beliefs imply a belief α , and K is the minimal such set (no proper subset of K implies α), removing one element of K will guarantee that it does not imply α . Contraction is done here by finding all such minimum subsets and removing an element from each of them. We call such minimal sets kernels. Kernels will be discussed in more details in the coming section. But we need to go over some example first.

Consider the same set:

$$\{a, a \rightarrow b, b, c\}$$

To contract the set by b, it is not enough to remove b from the set. If we do so, we get:

$$\{a, a \rightarrow b, c\}$$

which still implies b. We need to prevent the resulting set of beliefs from implying b. The minimal subset that implies b is

$$\{a, a \to b\}$$

We call it a b-kernel. Since the kernel is a minimal subset that implies b, removing only one element of that kernel is enough to make it not imply b. The only other b-kernel of the original set is $\{b\}$; we can only break this kernel by removing b. Breaking the other kernel can be done either by removing a, or by removing $a \to b$. So the two possible resulting belief sets after contraction are:

$$\{a,c\}$$
 and $\{a \to b,c\}$

This is how kernel contraction works. Every b-kernel is one way to infer b. To contract b every kernel should be broken by removing at least one belief.

Kernel contraction can be composed of two main steps: computing kernels, and cutting them. In this chapter, we will discuss kernels and see how they can be defined formally. We will discuss briefly how to compute kernels using an algorithm called *pinpointing*. Assuming we have a set of kernels that we computed, we will discuss previous work done on cutting kernels that will give us a good basis for introducing the new approaches afterwards. Then we will discuss the hitting-set approach to contraction in the context of *minimal change* requirement. Last topic to be discussed in this chapter will be our new graph approach kernel contraction. The algorithm will combine the two steps of kernel contraction (computing and cutting). It will be followed by some analysis and comparisons between different examples using it.

3.1 What are kernels?

Kernel contraction was introduced by Hansson[9] as a variant of an older approach called "safe contraction"[2]. In both approaches, contracting a knowledge base \mathbf{K} by α is done by discarding beliefs that contribute to make \mathbf{K} imply α . Beliefs that contribute to make \mathbf{K} imply α are members of some α -kernels of \mathbf{K} , and all members of every α -kernel contribute to make \mathbf{K} imply α . In the previous example, there were two b-kernels: $\{b\}$ and $\{a, a \to b\}$. We need to remove at least one element from each kernel to contract the original set by b. We refer to the set of α -kernels of \mathbf{K} by $\mathbf{K} \perp \alpha$.

Definition 3.1. (Kernel Set) According to [8], for a belief set **K** and a belief α , **K** $\perp \alpha$ is a set such that $X \in \mathbf{K} \perp \alpha$ if and only if:

- 1. $X \subseteq \mathbf{K}$,
- 2. $X \vdash \alpha$, and

3. if $Y \subset X$ then $Y \not\vdash \alpha$.

 $\mathbf{K} \perp \alpha$ is a kernel set that contains all α -kernels of \mathbf{K} .

Kernels are the smallest subsets of the knowledge base that imply a specific belief. Therefore, in order to give up that belief, every one of those kernels needs to be incised. If those kernels are not "minimal", they might contain some beliefs that do not contribute to make the kernel imply that specific belief. For example, in $\{\alpha, \beta\}$, β is a belief that does not contribute to make the set imply α . In that case, removing β only would not help in giving up α .

So, because of the minimality of the kernels, every belief in the kernel is significant to implying the belief that we want to give up (α) . And that's why removing one belief from a kernel is sufficient to make that kernel not imply α . So, we use a function that cuts every kernel in the kernel set. We call such function an *incision function*; it takes a set of kernels and removes an element from each kernel.

The incision function is defined in [8] as follows:

Definition 3.2. (Incision function) An incision function σ is a function such that for all beliefs α :

- 1. $\sigma(A \perp \alpha) \subseteq \bigcup (A \perp \alpha)$
- 2. If $\phi \neq X \in A \perp \alpha$, then $X \cap \sigma(A \perp \alpha) \neq \phi$

Contraction is done by removing the beliefs that are selected by the incision function from the original knowledge base. Contraction \approx_{σ} using the incision function σ can be defined as:

Definition 3.3. [8] (Contraction)

$$A \approx_{\sigma} \alpha = A \setminus \sigma(A \perp \alpha)$$

3.2 Computing kernels

Kernel contraction is about contracting knowledge bases using kernels. Kernels are the minimal subsets of the knowledge base that imply a certain consequence. The incision function can perform the contraction by cutting through all kernels of a specific belief. So, the first step in contraction is computing all kernels that imply the belief that needs to be removed. For that purpose, we use the pinpointing algorithm introduced in [6].

To show how the algorithm works, we use the example introduced in [6]. Given an \mathcal{EL} $TBox \mathcal{T}$ consisting of the following four GCI axioms:

$$\mathcal{T} = \{ax_1 : A \sqsubseteq \exists r.A, \quad ax_2 : A \sqsubseteq Y, \quad ax_3 : \exists r.Y \sqsubseteq B, \quad ax_4 : Y \sqsubseteq B\},$$

we can see that $A \sqsubseteq B$ holds according to \mathcal{T} , i.e. $A \sqsubseteq_{\mathcal{T}} B$. Let:

$$\alpha = A \sqsubseteq B$$
.

According to the definition of kernel sets:

$$\mathcal{T} \perp \alpha = \{ \{ax_2, ax_4\}, \{ax_1, ax_2, ax_3\} \}$$

The algorithm introduced in [6] computes all kernels using a modified version of the \mathcal{EL} subsumption algorithm. It works by finding a monotone boolean formula called "pinpointing formula". The propositions of the pinpointing formula are GCIs of the TBox, and a propositional valuation represents the kernels. In that sense, a valuation is a set of propositional variables that satisfy the formula, and these variables are the GCIs that constitute a kernel. So, by finding all valuations of the pinpointing formula we can get all kernels.

In the worst case, this approach takes exponential time (w.r.t the size of the TBox) to find all kernels. This is the case when there are exponentially many kernels. However, [6] also gives a polynomial-time algorithm that computes only one kernel. Because it is not part of the scope of this study, we are not going to discuss and analyze details of the pinpointing algorithm. All that matters is the worst-case time complexity, as it will affect the complexity of the kernel contraction algorithm that uses it.

3.3 Previous work

We now know we can use the pinpointing algorithm to get the set of all kernels. We then need to remove one element from each kernel to perform contraction. This is the role of the incision function; it picks an element from each kernel so that it cuts through all kernels. In this section, we look at some of the incision function implementations discussed in [11]. The goal of this study is to continue the work started in [11], and to revise some of what has already been done.

Given an \mathcal{EL} TBox \mathcal{T} and a belief α , $(\mathcal{T} \perp \alpha)$ is the set of α -kernels. The main contraction algorithm is shown in Algorithm 1.

Algorithm 1 Contraction algorithm

- 1: procedure CONTRACT (\mathcal{T}, A)
- $kernelset = PINPOINT(\mathcal{T}, A)$ 2:
- giveUpSet = Cut(kernelset)3:
- $\mathcal{T} = \mathcal{T} / \text{giveUpSet}$
- 5: end procedure

"kernelset" refers to $(\mathcal{T} \perp \alpha)$, and give UpSet is the set of beliefs selected by the incision function (CUT) to be removed. The algorithm is straightforward. It works by generating the set of kernels using the pinpoint formula as discussed in the previous section. Then it calls the incision function that picks up beliefs from every kernel, ensuring that it cuts all kernels. The beliefs are then removed from the knowledge base, i.e. from the TBox.

This general approach can be used with different incision functions. The call to the CUT function can be replaced with a call to another implementation of the incision function. The first implementation is the most straightforward one, where it removes a random belief from every kernel. This is given in Algorithm 2.

Algorithm 2 Random removal

```
1: function RANDOMREMOVE(kernelset)
2: giveUpSet = {}
3: for kernel \in kernelset do
4: choose a random belief \alpha from kernel
5: giveUpSet = giveUpSet \cup \{\alpha\}
6: end for
7: return giveUpSet
8: end function
```

The time complexity of RandomRemove function is polynomial: O(m), where m is the number of kernels (size of the kernelset), assuming that the random selection takes constant time. However, this is clearly not a good algorithm. In this example:

kernelset =
$$\{\{a, c\}, \{b, c\}\},\$$

one of the possible outcomes of the RandomRemove algorithm is:

$$give UpSet = \{c, b\}$$

which unnecessarily removes b. This could happen when the algorithm selects c from the first kernel and b from the second kernel. This solution:

$$giveUpSet = \{c\}$$

seems more concise and removes less beliefs. The second solution could be obtained if the algorithm is smart enough to check if the second kernel has already been incised.

The next algorithm (Algorithm 3) does this. It goes over all kernels, and randomly selects a formula from every kernel to be removed. Every time it selects a belief, it marks all kernels that contain that belief so that they are not considered for belief removal in the following iterations.

Algorithm 3 has a worst-case time complexity of $O(n^2)$, where n is the size of the kernel set. This is because the second main loop has another nested loop inside it, and neither of them takes more than n steps to finish.

Algorithm 3 Random removal with exclusion

```
1: function RANDOMREMOVEANDEXCLUDE(kernelset)
       giveUpSet = \{\}
 2:
 3:
       for i=0 to size(kernelset)-1 do
           kernelset[i].valid = true
 4:
       end for
 5:
 6:
       for i=0 to size(kernelset)-1 do
 7:
 8:
           if kernelset[i].valid == true then
              choose a random belief \alpha from kernelset[i]
 9:
              giveUpSet = giveUpSet \cup \{\alpha\}
10:
              for j=i to size(kernelset)-1 do
11:
                  if kernelset[j].contains(\alpha) then
12:
                      kernelset[j].valid = false
13:
                  end if
14:
              end for
15:
16:
           end if
       end for
17:
       return giveUpSet
18:
19: end function
```

3.3.1 Greedy Contraction

The greedy approach works by selecting the beliefs that appear in more kernels to be removed before those that appear in less kernels. At each step, the algorithm finds which belief appears the most in kernels and removes it, and then it forgets about those kernels (it does not consider those kernels that contain the removed belief for the following steps). This means that if a belief is selected for removal, all the kernels that contain it are already incised and no more beliefs need to be removed from these kernels.

Algorithm 4 Greedy Selection

```
1: function GREEDYCONTRACT(kernelset, beliefs)
2: initializeValidKernels(kernelset)
3: initializeBeliefOccurrences(beliefs)
4: computeOccurrences(kernelset, beliefs)
5: return selectMaxBeliefs(kernelset, beliefs)
6: end function
```

Algorithm 4 explains how the greedy approach for contraction works. Given a set of kernels and the set of all beliefs (the knowledge base), it proceeds as follows: it computes the number of occurrences of every belief in the kernels given, then it removes them one by one starting from the beliefs with maximum number of occurrences, until all kernels are incised. It does not actually remove the beliefs, but it collects them in the *giveUpSet* that will be returned to the main contraction, which in turn will remove them from the

```
1: // Mark all kernels as not cut to keep track of the remaining kernels to be cut
2: function InitializeValidKernels(kernelset)
      validKernels = 0 //global variable
      for i = 1 to size(kernelset) do
4:
         validKernel[i] = 1 //global array
5:
         validKernels = validKernels + 1
6:
7:
      end for
8: end function
1: // Create an array to count the number of kernels that include each belief
2: function InitializeBeliefOccurrences(beliefs)
      for i = 1 to size(beliefs) do
         occurs[i] = 0 //global \ array
4:
      end for
5:
6: end function
```

```
1: // For each belief, compute the number of kernels that include it
 2: function COMPUTEOCCURRENCES(kernelset, beliefs)
       for i = 1 to size(beliefs) do
 3:
          for j = 1 to size(kernelset) do
 4:
             if kernelset[j].contains(beliefs[i]) then
 5:
                 occurs[i] = occurs[i] + 1
 6:
             end if
 7:
 8:
          end for
       end for
10: end function
```

```
1: // Select the belief that appears in biggest number of kernels, and add to giveUpSet
2: function SELECTMAXBELIEFS(kernelset, beliefs)
       giveUpSet = \{\}
3:
       while validKernels > 0 do
 4:
          max = 0
 5:
          maxIndex = 0
 6:
          // search for the belief with max number of occurrences
 7:
          for i = 1 to size(beliefs) do
 8:
             if occurs[i] > max then
9:
10:
                 max = occurs[i]
                 maxIndex = i
11:
             end if
12:
          end for
13:
          // add the selected belief from the containing kernels to giveUpSet and update
14:
   count
15:
          for i = 1 to size(kernelset) do
16:
             if kernelset[i].contains(beliefs[maxIndex]) then
                 validKernel[i] = 0
17:
                 validKernels = validKernels - 1
18:
                 updateOccurrences(kernelset[i], beliefs)
19:
             end if
20:
          end for
21:
          occurs[maxIndex] = 0
22:
          giveUpSet = giveUpSet \cup \{beliefs[maxIndex]\}
23:
       end while
24:
       return giveUpSet
25:
26: end function
```

```
1: // disregard the given kernel from the kernels that include each belief
2: function UPDATEOCCURRENCES(kernel, beliefs)
     for i = 1 to size(kernel) do
3:
         removeOccurances(beliefs, kernel[i])
4:
      end for
5:
6: end function
1: // decrement the number of kernels that include the given belief
2: function REMOVEOCCURRENCES(beliefs, belief)
3:
      for i = 1 to size(beliefs) do
         if beliefs[i] == belief then
4:
            occurs[i] = occurs[i] - 1
5:
         end if
6:
7:
      end for
8: end function
```

knowledge base. So the greedy approach can be embedded in the main algorithm as a way to choosing the beliefs that need to be removed.

The worst-case time complexity of the greedy algorithm is lower-bounded by the complexity of the "while" loop in the selectMaxBeliefs() function. If the size of beliefs is m, the size of kernelset is n, and the size of the biggest kernel is k, then the worst-case time complexity is $O(n^2 \cdot k \cdot m)$. This could be significantly enhanced by using more clever data structures to make the in-validation of the kernels (the step where we decrement the number of valid kernels and update the occurrences) faster. But this is not very important now, because the greedy algorithm does not add to the optimality we are targeting in this chapter. As said before, the greedy algorithm only targets the minimality.

3.4 Minimal change

According to the information economy principal (minimality), in every change of the epistemic state, loss of information should be minimum[13]. This means that a system should choose an epistemic change outcome that minimizes loss of information. To satisfy the requirement of the minimum change we need an algorithm that removes the least number of GCIs while hitting all the kernels; but the kernels are nothing but sets of beliefs (GCIs). Luckily, this is exactly the *minimal hitting set problem*, which already has some relatively efficient algorithms that we can use here.

3.4.1 Hitting set approach

Another (more efficient) approach to cutting kernels to perform contraction is the hitting set algorithm. The minimal hitting set problem is defined as follows:

Definition 3.4. [1] Given a set $S = \{s_1, s_2, ..., s_n\}$ of n non-empty sets, a minimal hitting set d is a set such that:

$$\forall_{s_i \in S} [s_i \cap d \neq \emptyset] \land \nexists_{d' \subset d} [\forall_{s_i \in S} (s_i \cap d' \neq \emptyset)]$$

Thus, d is a minimal hitting set if and only if it contains at least an element from every set, and no proper subset of it is a hitting set. In the context of kernel contraction in \mathcal{EL} , we can define the minimal hitting set contraction as:

Definition 3.5. (Minimal hitting set contraction) Given a kernelset $K = \{k_1, k_2, ..., k_n\}$ of n kernels, a minimal hitting set $giveUpSet \subset K$ is a set such that:

$$\forall_{k_i \in K} [k_i \cap giveUpSet \neq \emptyset] \land \nexists_{giveUpSet' \subset giveUpSet} [\forall_{k_i \in K} (k_i \cap giveUpSet' \neq \emptyset)]$$

Since kernels are subsets of the TBox, and our goal to find a giveUpSet that hits all kernels, we can use approaches to the minimal hitting set problem to solve it. There are

good approximation algorithms, such as the one introduced in [1], for the minimal hitting set problem, that are feasible in terms of running time.

The notion of *minimality* can be interpreted in terms of set-containment or cardinality. We call a hitting set minimal in terms of caridnality if there is no smaller-sized set that is a hitting set. However, the interpretation we use here is minimality according to set-containment; this means a minimal hitting set is a hitting set that has no proper subset that is a hitting set. So there might exist different minimal hitting sets with different cardinalities, but none of them is subsumed by a proper subset that is a hitting set[1]. But our goal was to satisfy the information economy principal by removing hitting sets with minimum cardinality.

For that reason, after getting all minimum hitting sets, we need to consider only the ones that are smallest in size. For the kernel set:

$$kernelset = \{\{a, b\}, \{a, c\}\},\$$

there are two minimal hitting sets:

$$\{a\}$$
 and $\{b,c\}$

and they are of different sizes. The following step then is to determine that the smallest of them should be selected for contraction, which is $\{a\}$.

We can now use one of the minimal hitting set algorithms combined with the cardinality selection step to implement contraction for a TBox, by implementing the minimal incision function that adopts them, to achieve minimum change (which we can call then, a *minimal incision function*). We are not going to implement such a function in this study, but for now, we will assume that there is a function:

min-hit-CUT(kernelSet)

that takes a set of kernels, and returns a minimal hitting set (minimal in both set-containment and cardinality) of sentences to give up. We can use this function, as if it is implemented, and may implement it in some future work.

3.4.2 Graph approach

Now, we introduce another technique for kernel contraction in \mathcal{EL} that is based on graphs. The reason why following graph approach is useful in solving contraction problem is that we are performing contraction of TBoxes, and they have an implicit graph-like hierarchy.

An \mathcal{EL} Tbox consists of GCIs (we can transform all definition formulas to GCIs), that can be seen as nodes and edges. A GCI can be thought of as a directed edge between two nodes representing concepts on the two sides of the subsumption symbol (\sqsubseteq). So reasoning with TBoxes is similar to reasoning with directed graphs. We can reduce TBox reasoning problems to graph problems and use graph algorithms to solve them. The subsumption

relationship that forms a TBox seem to have an intuitive interpretation as a directed graph; and that's why we might use the word "subsumption hierarchy" later in this study to denote the relationship between concepts in the TBox.

The \mathcal{EL} subsumption algorithm described in [5] uses a graph approach, perhaps because it is intuitive to think of TBoxes as graphs. A lot of work has been done in the area of Graph Theory, which makes it easy to use the already existing algorithms to solve some graph problems. Also, graphs are easy to imagine and work with.

Our goal is to reduce the kernel contraction problem to a graph problem, and, using some efficient graph algorithms to perform kernel contraction. The algorithm we build here starts by transforming the TBox into a graph. Then, it generates all kernels by finding all paths that imply the formula we are contracting. After that, it removes one formula from each kernel by removing an edge from each path, since paths here represent kernels (this will all be explained shortly). Finally it transforms the graph back into a TBox.

We will start by describing the algorithm in details, and then explain each of its main steps.

Main algorithm

Given an $\mathcal{EL}\ TBox\ T$, and a GCI \mathbb{A} (where \mathbb{A} is in the form of $C \sqsubseteq D$, such that C and D are arbitrary concept expressions), we contract \mathcal{T} by A using Algorithm 5:

Algorithm 5 Contraction using graph approach

```
    function GRAPHCONTRACT($\mathcal{T}$, $\mathbb{A}$)
    complete($\mathcal{T}$)
    graph = transform($\mathcal{T}$)
    C = arg_{left}($\mathbb{A}$) // gets the concept expression on the left side of the GCI
    D = arg_{right}($\mathbb{A}$) // gets the right-side concept expression
    paths = getPaths(graph, C, D)
    cutPaths(graph, paths)
    $\mathcal{T}$ = de-transform(graph)
    end function
```

The complete function in step 2 uses the \mathcal{EL} subsumption algorithm to compute all subsumptions of \mathcal{T} . All the subsumptions computed are added to \mathcal{T} . The subsumption algorithm proceeds in four steps:[3]

- 1. Normalize the TBox.
- 2. Translate the normalized TBox into a graph.
- 3. Complete the graph using completion rules.
- 4. Read off the subsumption relationship from the normalized graph.

The algorithm is explained in full details in [3]. We now need to explain the transformation of the $TBox \mathcal{T}$ into a graph.

Transforming the TBox into a graph

Assuming the $TBox \mathcal{T}$ contains GCIs of the form $C \sqsubseteq D$ (where D is an arbitrary concept, and C is a concept expression of the form $c_1 \sqcap ... \sqcap c_n$ such that $n \geq 1$), we construct a graph graph = (V, E), where V is a set of nodes and E is a set of directed edges. Starting with an empty V and E, for every GCI $C \sqsubseteq D$, add C and E to E and E to E and E to E and E to E represents a concept expression, and every pair E represents the subsumption relation E is a set of directed edges.

Algorithm 6 Transforming a *TBox* to a graph

```
1: function TRANSFORM(\mathcal{T})
2: result = new Graph(V, E)
3: for every C \sqsubseteq D in \mathcal{T} do
4: V = V \cup \{C, D\}
5: E = E \cup \{(C, D)\}
6: end for
7: return result
8: end function
```

Transformation analysis

It is worth mentioning that even though subsumption can be decided in polynomial time with respect to the size of the TBox (using the algorithm in [5]), computing all subsumptions can be exponential in the number of concepts. For example, with:

```
\mathcal{T} = \{Human \subseteq Mammal, Mammal \subseteq Animal\},\
```

the new \mathcal{T} after adding all subsumptions would be:

```
\{Human \sqsubseteq Human, Human \sqsubseteq Mammal, Human \sqsubseteq Animal, Mammal \sqsubseteq Mammal, Mammal \sqsubseteq Animal, Animal \sqsubseteq Animal\}.
```

The new TBox contains a lot of unneeded GCIs. This can be avoided by adding a small step to the subsumption algorithm in [5], such that after completing the subsumption graph using the completion rules, we can remove all subsumptions of the form $X \subseteq X$ (e.g. $Human \subseteq Human$); also we need to remove all subsumptions $C \subseteq D$ if $C \subseteq X$ and $X \subseteq D$ (e.g. exclude $Human \subseteq Animal$ if $Human \subseteq Mammal$, and $Mammal \subseteq Animal$ are in the TBox) are already there.

Now we have a graph graph that represents the $TBox \mathcal{T}$. The next step is to compute the paths (using getPaths() function) from C to D (where C and D are graph nodes)

using depth-first search. Obviously, computing the paths using depth-first search can take polynomial time. For simplicity, we assume the TBox is acyclic. This means we don't allow the following situation:

$$\{C \sqsubseteq D, D \sqsubseteq E, E \sqsubseteq C\}.$$

Thus, the graph must be acyclic too. The algorithm can still be generalized to account for cycles. Computing all paths can be done as in Algorithm 7.

Algorithm 7 Computing all paths between two nodes

```
1: function GETPATHS(graph, C, D)
       result = \{\}
 2:
 3:
       Stack path = new empty Stack
       computeAllPaths(graph, C, D, path, result)
 4:
       return result
 5:
 6: end function
 1: function COMPUTEALLPATHS(graph, C, D, path, paths)
 2:
       graph = (V, E)
       for every (C, X) \in E do
3:
          if X = D then
 4:
              Stack\ temp = new\ empty\ Stack
 5:
              temp.pushAll(path) // adds all edges without changing path
 6:
              temp.push((C,X))
 7:
              paths = paths \cup \{temp\} // adding the path to the list of paths
 8:
9:
              \operatorname{path.push}((C,X))
10:
              computeAllPaths(graph, C, X, path, paths)
11:
              path.pop()
12:
          end if
13:
       end for
14:
15: end function
```

This can also be done if the graph contains cycles. It would require using a more complicated algorithm that keeps track of the number of edges and the nodes visited during execution. Finding paths in cyclic graphs in explained in details in [10].

Graph kernel contraction

The function cutPath is the contraction function. Every path from C to D is actually a subsumption path that entails $C \sqsubseteq D$. Removing an edge from such path would break the subsumption between C and D through this path. So, in order to contract $C \sqsubseteq D$ it is enough to remove one edge from each of the paths from C to D; as each path represents a kernel of $C \sqsubseteq D$ and breaking them is sufficient to give up the subsumption.

So, a simple and very easy implementation (though inefficient) for the contraction function is to go over the set of paths, and remove a random edge from every path. Suppose that

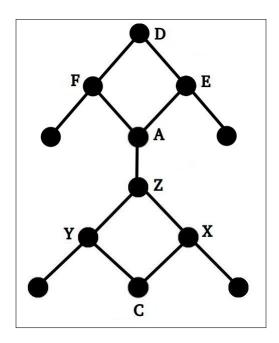


Figure 3.1: Four different paths between C and D, that share one edge.

the graph in Figure 3.1 is interpreted as C is the most specific concept that is subsumed by A $(C \sqsubseteq A)$, and as we climb the graph up, the concepts get more general. Here there are four C-D paths: (C, X, Z, A, E, D), (C, X, Z, A, F, D), (C, Y, Z, A, E, D), and (C, Y, Z, A, F, D). A function that removes random edges from each path might remove (C, X), (Y, Z), (E, D), and (F, D), which will actually remove the subsumption between C and D. However, it would be better (in terms of minimal change) to remove only (Z, A), which will also guarantee that subsumption between C and D is removed.

Sometimes we prefer to remove GCIs that involve most specific concepts during contraction (this will be discussed in the coming chapter). To contract such GCIs, we can just remove, from each C-D path, the edges going into C. In Figure 3.1, it would mean removing two edges: (C, X) and (C, Y). But removing such edges does not guarantee the minimum change. So we might end up having to choose which strategy is more preferred: minimal change or change with most specific concepts. The minimal hitting set approach that was mentioned earlier might not always satisfy specificity. So the user might have to choose which one to apply first, and which one to use as a tie breaker.

Removing the edges that involve nodes representing most specific concepts is straightforward; remove the edges that go into the most specific concept's node (C in our example). But it is not clear if one chooses to remove the least number of edges, instead, how this can be done. For this, we introduce an approach that uses the Minimum Cut algorithm to determine the minimum number of edges that need to be removed and identify them.

Reduction to network flow problem

As explained in [10], the network flow problem is the problem of computing the maximum flow possible in a network (represented as a graph) by finding the minimum cut of the network. The input to the Ford-Fulkerson algorithm, that solves this problem, would be:

- A graph G = (V, E).
- A source $s \in V$.
- A sink $t \in V$.
- Capacity function $C: E \to \mathbb{N}$ representing the maximum capacity of each edge.

To contract $C \sqsubseteq D$ given the TBox graph G, we choose C to be the source, D to be the sink, and we assume that the capacities of all edges are the same, equal to 1. The algorithm will find the maximum flow from C to D, which is equal to the capacity of the minimum cut (the sum of capacities of the cut edges); we can then extract the edges that form that minimum cut and remove them.

The approach of removing the minimum cut edges of the graph is equivalent to the approach of removing the minimal hitting set formulas of the kernels. The minimal hitting set is the set containing the minimum number of elements that hit all sets, while the minimum cut of the graph is the minimum number of edges (since they all have the same capacity) that cover all paths from C to D (where edges represent GCIs of the TBox and paths represent kernels.) So using the minimum cut approach should guarantee the minimum change for kernel contraction, as the minimal hitting set approach does.

Assuming we have a function "Ford-Fulkerson(graph, s, t)" that computes the maximum flow in the network (or graph) from source node "s" to a sink "t" with edge-capacities "1", and returns the set of edges of the minimum cut, we can modify the contraction algorithm to adopt the minimum cut approach as in Algorithm 8.

Algorithm 8 Another version of contraction algorithm

```
    function GRAPHCONTRACTUSINGMINCUT(T, A)
    complete(T)
    graph = transform(T)
    C = arg<sub>left</sub>(A) //arg<sub>left</sub> gets the concept expression on the left side of the GCI
    D = arg<sub>right</sub>(A) //arg<sub>right</sub> gets the right-side concept expression
    min-cut = Ford-Fulkerson(graph, C, D)
    remove min-cut edges from graph
    T = de-transform(graph)
    end function
```

For special cases such as contracting $C \sqsubseteq D1 \sqcap D2$, it is sufficient to contract $C \sqsubseteq D1$ first, and then contract $C \sqsubseteq D2$. But since the graph is already normalized (using

complete() function), rules of the form $C \sqsubseteq D1 \sqcap D2$ are already broken down into two: $C \sqsubseteq D1$, and $C \sqsubseteq D2$. Therefore, the conjunction \sqcap would only appear on the left hand side of a GCI in a normalized TBox (e.g. $A1 \sqcap A2 \sqsubseteq B$). In that case, for contracting $A1 \sqcap A2 \sqsubseteq B$, there will be a node $A1 \sqcap A2$, which we will use as a source node; the sink would be the node representing B.

As in [10], the minimum cut algorithm runs in polynomial time. So using it in the contraction algorithm will not have a significant effect on the complexity of the main algorithm (will not elevate the complexity from being polynomial to being exponential).

In some applications, the decision about which strategy to follow for choosing the edges to remove might vary depending on the situation. So the user can be asked in such case about which strategy to follow – specificity or minimality.

The last step of the algorithm is to transform the graph back to \mathcal{EL} . This can be done as follows: starting with an empty $TBox \, \mathcal{T}'$, for every edge (X,Y), add $X \sqsubseteq Y$ to \mathcal{T}' . The resulting TBox would be the result of contracting \mathcal{T} by \mathbb{A} . This is shown in Algorithm 9.

Algorithm 9 Transforming a graph back to a *TBox*

```
1: function DE-TRANSFORM(graph)
2: result = \{\}
3: graph = (V, E)
4: for every (X, Y) \in E do
5: result = result \cup \{X \sqsubseteq Y\}
6: end for
7: return result
8: end function
```

Since the running time of every step of the contraction algorithm starting from "transform(\mathcal{T})" until the last step is polynomial in the size of the TBox, the complexity of the algorithm will depend on the complexity of the first step (the complete() function). If building the subsumption hierarchy by generating all subsumptions of an $\mathcal{EL}\ TBox$ can be done in polynomial time, then the contraction algorithm will in turn take polynomial time. But if generating all subsumptions takes exponential time, then the algorithm will take exponential running time as well.

A Sample Run

Suppose we have a TBox \mathcal{T} :

$$\mathcal{T} = \{ A \sqsubseteq B, B \sqsubseteq C \}$$

which implies $A \sqsubseteq C$. Trying to contract \mathcal{T} by $A \sqsubseteq C$ using the network flow approach would work as follows:

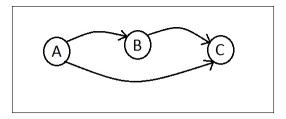


Figure 3.2: Graph representing TBox $\mathcal{T} = \{A \subseteq B, B \subseteq C, A \subseteq C\}$

- 1. complete(\mathcal{T}). The TBox is already normalized. Translating it into a graph and applying the completion rules will introduce $A \sqsubseteq C$ and will be added explicitly to the TBox.
- 2. $transform(\mathcal{T})$. After transforming the TBox into a graph, it will look like the graph in Figure 3.2.
- 3. getPaths(graph, A, C) will get all paths between A and C. There are only two paths: A-B-C and A-C.
- 4. Cutting the two paths A-B-C and A-C will be done by removing the edge between A and C, as well as one of the two edges A-B and B-C. So the resulting graph will only have one edge: A-B or B-C.
- 5. After transforming the graph again into a TBox, the result will be:

$$\mathcal{T} = \{A \sqsubseteq B\}$$

or

$$\mathcal{T} = \{B \sqsubseteq C\}.$$

The limitations of the algorithm

The previous example shows how the algorithm succeeds in finding the set of kernels by finding the paths from the source to the sink (where source and sink represent the two sides of the GCI that we need to remove). The example we will discuss now shows how the conjunction symbol (\sqcap) might introduces further complexity that the algorithm will not overcome. Given the TBox \mathcal{T} :

$$\mathcal{T} = \{A \sqsubseteq B, A \sqsubseteq C\}$$

we can see that it entails

$$A \sqsubseteq B \sqcap C$$
.

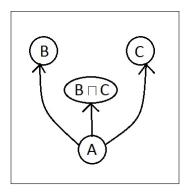


Figure 3.3: Graph representing TBox $\mathcal{T} = \{A \sqsubseteq B, A \sqsubseteq C, A \sqsubseteq B \sqcap C\}$

If we want to contract the TBox by $A \sqsubseteq B \sqcap C$, we would look for its kernels and remove a statement from each. In this example we have only one kernel:

$${A \sqsubseteq B, A \sqsubseteq C}.$$

So removing one of the two GCIs is enough to give up $A \sqsubseteq B \sqcap C$. Contracting it using our graph approach works as follows:

- 1. complete(\mathcal{T}). The TBox is already normalized. Translating it into a graph and applying the completion rules will introduce $A \sqsubseteq B \sqcap C$ and will be added explicitly to the TBox.
- 2. $transform(\mathcal{T})$. After transforming the TBox into a graph, it will look like the graph in Figure 3.3.
- 3. getPaths(graph, A, $B \sqcap C$) will get all paths between A and $B \sqcap C$. There is only one such path, which is the edge between them. So cutting this edge would be the only possible outcome of the algorithm.

This last step is where the algorithm fails. It finds only one of the kernels because it does not recognize that the two parallel edges from A to B and C actually form another kernel. This is because a kernel in our graph approach can only be represented as a path. The conjunction symbol \sqcap has a different meaning, and is not analogous to the path notion. However, the algorithm worked with the previous example because the GCI we tried to remove was the result of applying the transitivity property of the subsumption symbol (\sqsubseteq) to two other GCIs and they together were represented by a path in the graph representation.

Similarly, we can argue that the algorithm also fails when the existential quantification symbol (\exists) is used. For example, the following TBox:

$$\mathcal{T} = \{ A \sqsubseteq \exists r.B, B \sqsubseteq C \}$$

implies the following expression:

$$A \sqsubseteq \exists r.C$$

The graph approach will not get the kernels of that expression using the getPaths() function. So, it will also fail. But this is only because of the subsumption of B by C, where B is included in the role expression $\exists r.B$. If the existential quantification symbol is used without changing the symbols involved in the roles (such as B), the algorithm will work fine, as it will only be depending on the subsumption relations between concept expressions.

So it seems that the limitations of this algorithm are only due to the difference between the inference using paths between nodes and inference the introduces conjunction or existential quantification. However, the algorithm does not always fail when \sqcap or \exists is involved. In Figure 3.2, if we replace B by $(X \sqcap Y)$, or by $(\exists r.B)$, the algorithm will succeed. That is because the nodes do not change after the inference; all that is added by the subsumption algorithm are new edges.

Analysis of the graph algorithm

It is important to be able to measure the correctness of the solution we get by running the algorithm. The AGM framework provides a set of rationality postulates for contraction that we use for this purpose. Let's see whether the algorithm satisfies each of those postulates.

(K-1) For any sentence A and any belief set K, K_A^- is a belief set.

Since the algorithm only removes edges (or GCIs), the remaining set of edges still represent a set of GCIs, which is a belief set.

$$(\mathbf{K} - \mathbf{2}) \ K_A^- \subseteq K.$$

This is also satisfied as the new set is obtained by only removing GCIs.

(**K** - 3) If
$$A \notin K$$
, then $K_A^- = K$.

If the algorithm is given a GCI that is not in the set, the source and sink nodes will not be in the graph. In this case, the minimum cut algorithm will not have any flow and will not remove any edges. So, the set will remain the same after contraction.

(K – **4)** If not
$$\vdash A$$
, then $A \notin K_A^-$.

If a GCI is given the algorithm, the remaining set after contraction will not contain this GCI, because the edge between both sides of the GCI will be removed. By having the edge removed, the transformation of the graph into a TBox will not create such GCI.

(K – **5)** If
$$A \in K$$
, then $K \subseteq (K_A^-)_A^+$.

This is called the recovery postulate. This is not satisfied by the algorithm. The reason for this is that if a GCI is contracted by the algorithm, this can sometimes involve removing some edges along the path from the source node to the sink node. Adding the GCI back to the new set will not trigger the introduction of those other edges that were removed.

(K - 6) If
$$\vdash A \leftrightarrow B$$
, then $K_A^- = K_B^-$.

This is trivially satisfied, because in \mathcal{EL} two GCIs are equivalent if and only if they are exactly the same.

$$(\mathbf{K} - \mathbf{7}) K_A^- \cap K_B^- \subseteq K_{A\&B}^-.$$

The algorithm contracts only one belief at a time. If we would like to contract something like (A&B), we would do it by either contracting A or contracting B (contracting (A&B)) is the same as contracting one of them). And since contraction is monotonic (it only removes GCIs), the set resulting from contracting A cannot be bigger than the original, and same with B. Thus, the intersection of both of them cannot be bigger than one of them (as no other GCIs were introduced).

$$(K - 8)$$
 If $A \notin K_{A\&B}^-$, then $K_{A\&B}^- \subseteq K_A^-$.

If A is not present after contracting (A&B), it means that contraction of (A&B) was done by contracting A. In this case, the remaining set is the same for both contractions.

3.4.3 How to generate kernels?

So far, we have discussed two ways for generating all kernels:

- 1. Using the pinpointing algorithm.
- 2. Using the graph approach.

Computing kernels using the graph approach finds all kernels by performing depth-first search on the graph representation of the TBox. It finds all kernels in O(E), where E is the number of edges in the graph (or the number of GCIs in the TBox). However, it only works with subsumptions that do not include the conjunction symbol (\square). So, in cases where the conjunction is not used in the TBox, the graph approach will be at least as efficient as the pinpointing approach, if not better (the axiom pinpointing algorithm introduced in [6] finds one kernel in polynomial time).

3.4.4 What incision function to use?

If we care about selecting minimum number of beliefs to remove, then there are two incision approaches we discussed so far:

- 1. The minimum hitting set algorithm.
- 2. The minimum cut algorithm for the network flow problem.

The minimum hitting set problem is NP-hard. However, the algorithm we discussed in this study is a relatively efficient approximation algorithm. However, in cases where the minimum cut approach can be used, the minimum cut algorithm is probably the best and most efficient. The algorithm works by finding the max flow possible through the network build from the TBox, and then uses it to find the minimum cut.

The Ford-Fulkerson maximum flow algorithm typically takes O(E.C) steps to find the maximum flow, where E is the number of edges and C is the sum of all capacities. Since all edges in our approach has capacity 1, C is equal to E. So the complexity of the algorithm is $O(E^2)$. After finding the maximum flow, we can find the minimum cut in O(E). So the overall complexity of the algorithm is $O(E^2)$.

In other cases where the graph approach is not applicable, the minimum hitting set algorithm is a good candidate. In the next chapter, we will also give some techniques for implementing the incision function. Some of them might be even more suitable than the minimum hitting set approach, depending on the domain and the preferences of the user.

Chapter 4

Heuristics for contraction

In this study we care about approaching optimal contraction. In the context of contraction, we can define optimality to be removing the smallest number of beliefs, or removing beliefs that make more sense to be removed together. In the last chapter, we discussed contraction algorithms that attempt to achieve optimality by trying to remove the smallest number of beliefs, which we called minimum change. However, we haven't considered the significance of the subsumption hierarchy of \mathcal{EL} in choosing the beliefs to be removed from a TBox. In this chapter, we look at heuristics that are motivated by the subsumption relationship between formulas. We will discuss ways of determining a preferred set of beliefs to be removed, based on the subsumption relationship of \mathcal{EL} represented by the symbol \sqsubseteq .

According to localization and specificity heuristics¹, we can prefer to remove a certain set of beliefs to another, even if it is of bigger or equal size. We can say that it is "semantically better" to remove beliefs that are related than to remove unrelated beliefs; by "semantically" we mean according to the meaning of the language's logical connectives. Also, it is better to remove beliefs about specific concepts than general ones. These two selection heuristics, localization and specificity, can be used as tie-breakers when we have multiple giveUpSets of the same size and we need to select only one. They can also be used on their own regardless of the size of the giveUpSets.

4.1 Localization

Giving up beliefs about a certain concept means being in doubt of this concept. The more beliefs we remove, the more concepts we become in doubt of. This can lead us to consider removing beliefs about a certain concept rather than beliefs about many different concepts. As discussed in [11], it is more efficient to remove beliefs that share concepts or roles in a TBox as it means being in doubt of less number of concepts.

¹These heuristics will be explained shortly.

This seems a reasonable approach as humans tend to deal with related beliefs than unrelated ones. If we try to revise our knowledge, we will probably find ourselves leaning towards focusing on beliefs that share common concepts.

The following \mathcal{EL} example:

- (1) $AspergillusFumigatus \sqsubseteq Multicellular$
 - (2) $Multicellular \sqsubseteq NotBacteria$
- (3) $AspergillusFumigatus \sqsubseteq Eukaryotes$
 - (4) $Eukaryotes \sqsubseteq NotBacteria$

implies:

(5) $AspergillusFumigatus \sqsubseteq NotBacteria$,

and we can see that there are two (5)-kernels:

$$\{1, 2\}$$
 and $\{3, 4\}$

So, to contract the knowledge base by (5), we need to remove a belief from each kernel. However, removing (1) and (4) seems unreasonable because it means that we are in doubt of our knowledge of AspergillusFumigatus, Multicellular beings, Eukaryotes, and beings that are NotBacteria, while removing (1) and (3), on the other hand, makes us in doubt only of our knowledge of AspergillusFumigatus, Multicellular beings, and Eukaryotes. Not only that, but removing (1) and (3) actually means that our knowledge of AspergillusFumigatus is wrong; Our knowledge of Multicellular beings and Eukaryotes is not necessarily false. Same with removing (2) and (4), we suspect the soundness of our beliefs about beings that are NotBacteria.

Localization is implemented in [11] using a graph-like approach. The algorithm operates by defining edge-like relationship between formulas and using graph connectivity and graph clustering rules to determine localized sets.

Definition 4.1. For every two GCIs X and Y, we say that there is an edge between them if they share a concept or role.

After applying that, we will end up with a graph-like structure, where some GCIs of the knowledge base are connected by edges. The most basic path in a graph is an edge, but a path can be composed of more than one edge. Generally, a path is a sequence of edges.

Definition 4.2. (Connected GCIs) Two GCIs \mathbb{A} and \mathbb{B} are connected if and only if there is a path from one of them to the other, i.e. if there is a sequence of edges and GCIs $Ae_1G_1e_2...G_{n-1}e_nB$, where G_i is a GCI, e_i is an edge, every triple $G_ie_{i+1}G_{i+1}$ means G_i and G_{i+1} are connected by an edge, and $n \geq 0$.

In the last \mathcal{EL} example, we can remove one of the following sets to contract (5):

- {1, 3} (a)
- {1, 4}. (b)

There are more possible *giveUpSets*, but (a) and (b) are sufficient to show how the definition of connected GCIs can be applied.

For the set (a), there is an edge between (1) and (3) because they share the same concept AspergillusFumigatus. So, according to the definition, (1) and (3) are connected.

For the set (b), there is no edge between (1) and (4) because they don't share any concept or role. So according to the definition, (1) and (4) are not connected.

Definition 4.3. (Clusters) A GCI \mathbb{X} belongs to a cluster \mathfrak{C} if and only if \mathbb{X} is connected to every GCI in \mathfrak{C} .

Thus, according to the definitions, the set (a) has only one cluster, as (1) and (3) are connected and belong to the same cluster. However, the set (b) has two clusters, because (1) and (4) are not connected in (b), so they belong to two different clusters.

The algorithm uses the notion of clusters to identify the best giveUpSet. A GCI belong to a cluster if it shares concepts or roles with other GCIs in the giveUpSet. If a giveUpSet contains a lower number of clusters than another set, then it is better to remove it during contraction. This is because the more clusters there is in a set the more sparse the graph is, hence the less localized the contraction is.

The algorithm works on each possible give UpSet separately. It builds the graph-like relation between formulas in the set according to the definitions we gave. It is important that the algorithm computes the connectivity in each set separately, because two formulas might have a path between them in one set but not the other; so they would be considered connected in a set and disconnected in the other. The algorithm then counts the number of clusters. It does that for all the sets, then chooses the set with least number of clusters to be removed.

Algorithm 10 selects the giveUpSet with least number of clusters by first computing the number of clusters in each giveUpSet and then choosing the one with the smallest number. The time complexity of the algorithm is $O(n^3m)$, where m is the number of kernels, n is the size of the biggest kernel. The loop in the main function goes over the whole list of kernels, which is of size m, and the second function has a 3-level nested loop that goes over the kernel whose size is at most n in two levels. The third level checks the connectivity of two nodes, which can take at most n if it has to go over all the elements in the kernel.

4.2 Specificity

The subsumption hierarchy enforced by the subsumption relation in \mathcal{EL} categorizes concepts into different levels of generality and specificity. Consider the kernel:

Algorithm 10 Computing localized hit

```
1: // Compute the number of clusters in each set and choose the least
 2: function GETLOCALIZEDHIT(giveUpSets)
 3:
       setWithLeastClusters = null
 4:
       smallestNumber = \infty
       for giveUpSet \in giveUpSets do
 5:
          numOfClusters = qetNumberOfClusters(qiveUpSet)
 6:
          if numOfClusters < smallestNumber then
 7:
             setWithLeastClusters = qiveUpSet
 8:
             smallestNumber = numOfClusters
 9:
          end if
10:
       end for
11:
       {f return}\ set With Least Clusters
12:
13: end function
 1: // Compute the number of clusters
 2: function GETNUMBEROFCLUSTERS(giveUpSet)
       for i = 1 to size(giveUpSet) do
          label[i] = i
 4:
       end for
 5:
       for i = 1 to size(giveUpSet) do
 6:
          cluster[i] = 0
 7:
       end for
 8:
      // Label connected nodes (beliefs) similarly
 9:
      for i = 1 to size(qiveUpSet) do
10:
          for j = i + 1 to size(giveUpSet) do
11:
             if giveUpSet[i] and giveUpSet[j] are connected then
12:
13:
                 label[j] = label[i]
             end if
14:
          end for
15:
      end for
16:
       // Mark clusters based on used labels. Each label now represents a cluster
17:
      for i = 1 to size(giveUpSet) do
18:
          cluster[label[i]] = 1
19:
      end for
20:
21:
      // Calculate the number of clusters by counting the used labels
       for i = 1 to size(giveUpSet) do
22:
          numberOfClusters = numberOfClusters + cluster[i]
23:
       end for
24:
       return numberOfClusters
25:
26: end function
```

 $Lion \sqsubseteq Mammal$ $Mammal \sqsubseteq Vertebrate$ $Vertebrate \sqsubseteq Animal$

Figure 4.1: Simple Animal kernel

which implicitly entails

 $Lion \sqsubseteq Animal$

And suppose we would like to contract the TBox by $Lion \sqsubseteq Animal$. We have three options: we can remove $Lion \sqsubseteq Mammal$, $Mammal \sqsubseteq Vertebrate$, or $Vertebrate \sqsubseteq Animal$. Removing any of the three GCIs would guarantee minimum change. We can say that Lion is more specific than Mammal and Mammal is more specific than Vertebrate. Removing $Lion \sqsubseteq Mammal$, in this case, is more reasonable than removing $Vertebrate \sqsubseteq Animal$, because $Lion \sqsubseteq Mammal$ involves concepts that are more specific than the ones involved in $Vertebrate \sqsubseteq Animal$. Removing $Vertebrate \sqsubseteq Animal$ may affect more concepts in the subsumption hierarchy (in worst case) than those affected by removing $Lion \sqsubseteq Mammal$. This means that removing $Vertebrate \sqsubseteq Animal$ means that all individuals that are Vertebrates are no longer believed to be Animals, while removing $Lion \sqsubseteq Mammal$ only affects a subset of Vertebrates which is Lions.

So, it is preferable to remove GCIs that involve specific concepts rather than removing GCIs that involve more general ones. To account for specificity, we assign a label to each of the concepts representing its level in the subsumption hierarchy, where levels increase with generality. And during contraction, we consider contracting GCIs that involve concepts at lower level (more specific) before considering GCIs that involve concepts at higher level.

Before explaining how the algorithm works, we need to explain the notion of *children* that we use in the context of subsumption hierarchy. We use this notion to define the relationship between concepts that reflects a tree-like relationship between concepts. This will help us determine and compute specificity for each GCI.

Similar to the hierarchy of the subsumption graph of an \mathcal{EL} TBox, we give a definition to parent and child concepts as follows:

Definition 4.4. For the GCI $A \subseteq B$:

- A is a *child* concept relative to B, and
- B is a parent concept relative to A.

So by looking again at Figure 4.1, we can now say that:

• Mammal is a parent of Lion, and Lion is a child of Mammal

- Vertebrate is a parent of Mammal, and Mammal is a child of Vertebrate
- Animal is a parent of Vertebrate, and Vertebrate is a child of Animal
- Animal is the most *general* concept, i.e. Animal subsumes all other concepts
- Lion is the most *specific* concept (least *general*), i.e. it is subsumed by all other concepts

The approach we follow in this study for adopting the preference of removing sentences that contain more specific concepts is achieved by assigning weights to every GCI that reflects its generality. Then we select the kernels with minimal overall weight. Every GCI gets a numeric weight depending on the level of generality of the concepts involved in it (depending on their position in the subsumption graph). A GCI $A \sqsubseteq B$ gets weight '0' if A has no children, i.e. if there is no rule of the form $X \sqsubseteq A$, where X is an arbitrary concept; and the GCI involving its parent (A GCI of the form $B \sqsubseteq C$) will have weight '1' in that case. So the weight somehow represents the level in the subsumption graph, starting from level '0' at the trees' leaves (which is A in this example.) For simplicity, we assume that the TBox is acyclic – so we avoid the problems that will be caused by loops.

Now we can compute the weights for the last TBox:

- $Lion \sqsubseteq Mammal$ has weight = 0
- $Mammal \subseteq Vertebrate$ has weight = 1
- $Vertebrate \sqsubseteq Animal \text{ has weight} = 2$

The algorithm we introduce now removes kernels that contain GCIs involving most specific concepts by first computing the weights of every kernel based on the weights of the GCIs inside it. Following Definition 4.4, we start the algorithm by building the *children* labels for all the concepts in the TBox that we will use to assign weights to the GCIs. Given a TBox T, the algorithm proceeds as in Algorithm 11.

Algorithm 11 Building the children graph

```
1: function GRAPHBUILD(T)

2: children(X) = \emptyset, for every concept X in T

3: for every A \sqsubseteq B \in T do

4: children(B) = children(B) \cup \{A\}

5: end for

6: end function
```

Now we have a graph composed of nodes and their *child* relationship; the nodes represent concepts and the edges (or the child relationship) are the subsumption relationship between them. So GCIs are represented by edges in the graph. The roots are now nodes that are not

children of any node, and the leaves are nodes with no children. The edges involving leaves are assigned weight '0' and it increases by '1' every step towards the roots. So, the roots are most general concepts, and the leaves are most specific ones. If a GCI contains concepts $\alpha \sqsubseteq \beta$, where α has no children and β has no parents, then it gets weight '0' according to Algorithm 12. The weights are assigned to edges, not nodes. Given an edge e that connects node a to its child b, if b has more than one child, the weight of e is determined by the longest sequence of edges to a leaf starting at any of the children of b. So the GetMaxWeight function traverses a tree to all of its leaves computing the weight of each edge and choosing the weight according to the longest path to a leaf.

Algorithm 12 Assigning Weights

```
1: function AssignWeights(T)
2:
       for every A \sqsubseteq B \in T do
           weight(A \sqsubseteq B) = GetWeight(A \sqsubseteq B)
 3:
       end for
 5: end function
 1: function GETWEIGHT(X \sqsubseteq Y)
       if weight(X \sqsubseteq Y) \neq NULL then
 2:
           return weight(X \sqsubseteq Y)
3:
       end if
 4:
       if children(X) = \emptyset then
 5:
           return 0
 6:
 7:
       else
           return 1 + GetMaxWeight(X)
 8:
9:
       end if
10: end function
 1: function GETMAXWEIGHT(X)
       \max = -1
2:
3:
       for every Z in children(X) do
           w = GetWeight(Z \sqsubseteq X)
 4:
           if w > max then
 5:
               \max = w
 6:
           end if
 7:
 8:
       end for
       return max
9:
10: end function
```

Now we have every GCI in T assigned a weight relative to its level of generality. So, in every kernel we have a preference level of what to give up. Based on the weights, we will choose the sentences with less weight over the ones with more weight to remove. This preference can be used as a tie-breaker after we apply the minimal hitting set algorithm and end up with more than one minimal set. If we arrive at two minimal hitting sets for the kernels, we can compute the overall weight of each of the hitting sets and remove the GCIs in the set with less overall weight. This is shown in Algorithm 13. It uses the min-hit-CUT

algorithm that gets the minimum hitting sets given a kernel set. There could be more than one minimum hitting set.

Algorithm 13 Removing specific hitting set

```
1: function GETMOSTSPECIFICHIT(kernelset)
       min-hit-sets = min-hit-CUT(kernelSet)
 2:
 3:
       min = \infty
       hit = NULL
 4:
       w(s) = 0, for every s \in min-hit-sets
 5:
       for every s in min-hit-sets do
 6:
 7:
           for every A \sqsubseteq B in s do
               w(s) = w(s) + weight(A \square B)
 8:
 9:
           end for
           \mathbf{if}\ w(s) < \min\ \mathbf{then}
10:
               \min = w(s)
11:
               hit = s
12:
           end if
13:
       end for
14:
       return hit
16: end function
```

Now the CUT method in Algorithm 1 can be re-implemented using the algorithms introduced here to account for specificity heuristics. Given a TBox T and a GCI A, the contraction algorithm is given in Algorithm 14.

Algorithm 14 Contraction algorithm – modified

```
1: procedure CONTRACT(T, A)
2: kernelset = PINPOINT(T, A)
3: graphBuild(T)
4: assignWeights(T)
5: giveUpSet = GETMOSTSPECIFICHIT(kernelset)
6: T = T / giveUpSet
7: end procedure
```

The contraction algorithms in this version guarantees the minimality, then specificity, of the removed beliefs. It guarantees the minimality by reducing the selection of beliefs from the kernelset to the minimal hitting set problem. The better the performance and the optimality of the hitting set algorithms, the efficient and minimal the contraction algorithm is. If the minimal set algorithm produces more than one solution of the same size, the specificity heuristic is used as a tie-breaker. It selects the solution with more specific beliefs to be removed.

The time complexity of Algorithm 12 is polynomial. The worst-case time complexity is O(nl), where n is the size of the TBox T and l is the depth of the subsumption graph. Since the largest value for l is n, we can consider the worst-case time complexity to be $O(n^2)$. The

algorithm works as depth-first search because of the recursive call in getWeight function. The graphBuild function takes O(n) steps to build the parent-child relationship graph, where n is the size of the TBox T. Finally, the getMostSpecificHit function takes O(nm) steps in the worst case, where n is the size of the TBox T, and m is the number of kernel sets. So the contraction algorithm takes polynomial time if the minimum hitting set algorithm runs in polynomial time. But, if the minimum hitting set algorithm is not polynomial-time, then the contraction algorithm will not be. Thus, it is safe to say that the time complexity of the contraction algorithm is lower-bounded by the time complexity of the minimum hitting set algorithm's time complexity.

Chapter 5

Conclusion

We have discussed that the AGM framework introduces a reasonable set of rules that can be followed to reach rationality in building belief systems. However, we used a different formalism to express knowledge and perform contraction throughout this study than the one the AGM framework uses. This formalism, which is \mathcal{EL} , belongs to a family called Description Logic (DL). \mathcal{EL} is one of the simple members of the DL family, and, as the rest of the DLs, uses concepts and roles as the building blocks of the knowledge base. We only showed how to perform contraction on TBoxes, and not ABoxes. We then discussed some basic approaches to contraction and introduced a general algorithm that can accommodate the use of different heuristics to seek optimality.

We introduced a restricted contraction algorithm that uses graphs and solves contraction as a network flow problem. The graph approach is restricted in the sense that it can be used only in certain cases where inference in the TBox involves only the transitive property of the subsumption relation. We showed how the approach is sound and complete (only in the restricted cases) by showing that it follows most of the AGM postulates for contraction. We also discussed three heuristics, two of which are based on the semantics of the subsumption hierarchy of \mathcal{EL} , which are Localization and Specificity. We showed also how the greedy approach for contraction can be used.

The bottleneck for the contraction algorithms we discussed is actually generating all the kernels of a specific belief. The pinpointing algorithm we use for this purpose can sometimes take exponential number of steps with respect to the size of the TBox. Other than that, all the algorithms we discussed take polynomial time to run.

What can be done in the future is to give an algorithm that contracts explicit knowledge in the ABox we well. Also, the use of \mathcal{EL} was very important to finding polynomial time contraction algorithms. Would using other more expressive versions of DL result in higher order contraction than polynomial? What advantage would using a more expressive DL add and will it be worth it if contraction algorithms get harder and more complicated?

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