

LEAN-aided verification of datalog reasoning results

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March 29, 2024

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Acknowledgements

I would like to thank my advisors Lukas Gerlach and Dr. Stephan Mennicke for introducing Lean and the problem to me and their helpful comments during the creation of this thesis. Lukas also improved my previous graph model in section 5 by adding the hash map implementation.

I would like to thank Prof. Markus Krötzsch for the formal supervision and his helpful comments during the writing. He proposed amongst other things example 9.

1 Introduction

One of the most important problems in software engineering is writing bug-free software. Bugs in software may endanger the well-being of property and humans.

A recent example of the consequences of buggy software was the Horizon software used by the British Royal Mail [10]. It was supposed to be an accounting software used in the accounting of the individual post offices. Horizon wrongly calculated the balance which led to the individual subpostmasters being accused of theft or fraud. Over 900 people were sentenced to the repayment of alleged damages or prison sentences. These people still suffer financial, social or health consequences despite being innocent and multiple persons committed suicide.

The Royal Mail trusted their program too much to consider that it contained errors. But what methods exist to increase the confidence in a program? Firstly, we can prove the correctness of a program, which is done in the field of formal verification. These proofs are complicated and often very large so they need computer assistance. Due to time constraints or unfamiliarity with the methods this is only very rarely done in use cases that require specific security guarantees. An example is the verified C compiler CompCert[19].

In practice, software developers mostly employ a test-driven approach to software correctness. They design specific use cases and specify what the program is supposed to do in this case. Designing such cases is difficult and cannot cover every scenario so we can only see that some scenarios are bug-free.

An interesting development in algorithms research are certifying algorithms [21]. These algorithms do not only return a solution but also a certificate as an explanation why the returned value is the correct answer. In the case of SAT-solving, they would for example return a possible model as well as stating that a formula is satisfiable. The user can then verify on their own that the result is correct without inspecting the code's inner workings. This can also be done by programs called checkers. These are usually smaller and simpler so that they can be formally verified.

Our interest concerns computing the results of datalog programs. Datalog is a logic programming and query language that allows to write rules in the following way

$$\begin{aligned} T(?x, ?y) &\leftarrow E(?x, ?y). \\ T(?x, ?z) &\leftarrow E(?x, ?y), T(?y, ?z). \end{aligned} \tag{1}$$

The rules can be interpreted as formulas from first-order logic that are universally quantified and use the implication. There are multiple equivalent semantics of datalog that deal with the question of what follows from a datalog program and a given set of facts often known as the database.

This task can be solved in polynomial time which allows using datalog in multiple interesting applications from data analysis to data integration.

Modern datalog engines like Nemo[16] or Souffle[18] can work with millions of database elements and use many optimization techniques to achieve this. Unfortunately, the correctness may suffer from this and formally verifying an engine is out of reach. They offer proof trees as an explanation why a certain element was derived so that we may view them as an instance of certifying algorithms. In this style, we want to implement a checker that takes these certificates and tells us whether a datalog reasoning result is correct.

With the goal of improving the confidence in the correctness of our checker, we will formally verify it using the proof assistant Lean[13]. A proof assistant allows users to define functions and objects and prove results about the objects in machine-readable style. Therefore the proof assistant can verify that the proof is indeed correct or if not raise errors.

After introducing the basics about datalog, certifying algorithms and Lean in section 2, we will formalize datalog in Lean in section 3. After that, we provide two methods to verify a datalog reasoning result: a classical result based on proof trees in section 4 and another result based on graphs we see as more promising in practice in section 5. These methods however can only tell us that all atoms present in the result are correctly derived, i.e. soundness. They do not tell us whether more could be derived. A method to solve this question is presented in section 6.

Finally, we show in section 7 that our results are not only formally correct but can also be used in practice.

2 Preliminaries

In this section, we introduce concepts that we are going to use in future sections. We start by defining datalog, which we will formalize later. After that, we review the concept of certifying algorithms and finally, we introduce Lean.

2.1 Datalog

Datalog is a logic programming language and can be syntactically considered as a subset of Prolog. An introduction to datalog and other query languages can be found in [1], which we recall next. Some knowledge about first-order logic may be beneficial, which be found for example in [15].

In order to write datalog rules, we require the existence of three at most countable sets V, C and P , where V represents variables, C represents constants and P represents predicate symbols. Every predicate symbol p in P has an arity $ar(p) \in \mathbb{N}$, where $0 \in \mathbb{N}$.

Example 1. *In general, we consider in this work R as strings that start with a capital letter, V as strings that start with a question mark and C as all other non-empty strings.*

Example 2. *In a more concrete example we have the variables $V = \{?x, ?y, ?z\}$ and constants $C = 1, \dots, 7$. The predicate symbols are the binary predicates (i.e. they have an arity of two) R and E and the nullary (i.e. arity zero) predicate Q .*

In the language of first-order logic as in [15], C and P build the signature of the language we are going to build. We note that this fragment of first-order logic does not use function terms.

The following definitions are adapted from logic for the signature above.

A *term* is either a variable or a constant. In order to differentiate meaningfully between them, we require that $C \cap V = \emptyset$ so that no symbol is both a variable and a constant. As long as both C and V are finite, we will only have a finite amount of terms. This is in contrast to general first-order logic or logic programming where function symbols can lead to an infinite amount of terms even if the signature is finite.

Example 2 (continued). *Examples of terms in our language are $?z$, 2, 5 or $?x$.*

A counterexample would be word or A as those elements do not occur in either C or V .

An *atom* is an expression of the form $A(t_1, \dots, t_n)$ for $n \in \mathbb{N}$ where A is a predicate symbol ($A \in P$) and t_1, \dots, t_n are terms. Additionally, we require that n is the arity of A , i.e. $ar(A) = n$.

Example 2 (continued). *Examples for atoms are here: $Q()$, $R(?x, ?y)$, $E(2, 3)$ or $E(?x, 7)$. In every atom, the number of terms matches the arity of the predicate symbol. We note that it is allowed to mix variables and constants in the terms of an atom.*

A first counterexample would be $Q(a)$ because the arity does not match. Another counterexample would be $R(E(2,3),4)$ because $E(2,3)$ is not a term.

In the example, some atoms used only constants in their terms, while others included variables. This is expressed by $Vars(A(t_1, \dots, t_n)) = \{t_i \mid t_i \in C\}$. If for some atom a $Vars(a)$ is empty, we call this a *ground atom*.

Example 2 (continued). In the previous example $Q()$ and $E(2,3)$ are ground atoms. The first does not have any terms at all, whereas the other only uses constants.

For the other atoms we have $Vars(R(?x, ?y)) = \{?x, ?y\}$ and $Vars(E(?x, 7)) = \{?x\}$, so that they are not ground atoms.

A *rule* is an expression of the form $H \leftarrow B_1 \dots B_n$ for atoms H and B_i for $n \in \mathbb{N}$. We call H the *head* of the rule and B_1, \dots, B_n the *body* of the rule and define the functions $head(r) = H$ and $body(r) = B_i$. We allow rules with an empty body. These rules are called *facts*.

We can generalize the $Vars$ function to rules and use this to define *ground rules*. For a rule $r = H \leftarrow B_1 \dots B_n$, we define

$$Vars(r) = Vars(H) \cup \bigcup_{i \in \{1, \dots, n\}} Vars(B_i)$$

and call r a ground rule if $Vars(r) = \emptyset$. This means that a rule is a ground rule if the head and every atom in the body is a ground atom.

Example 2 (continued). Examples for rules are $E(1,2) \leftarrow$, $Q() \leftarrow E(?x, ?y)$ or $T(?x, ?z) : \neg E(?x, ?y), T(?y, ?z)$.

$E(1,2) \leftarrow$ is both a fact and a ground rule, but not every fact is a ground rule as for example $E(?x, ?x) \leftarrow$

A *program* P is a finite set of rules. We only consider finite sets as infinite sets are difficult to express in practice.

Example 2 (continued). We consider the program $P = \{$

$$\begin{aligned} &E(1,2) \leftarrow \\ &E(1,3) \leftarrow \\ &E(3,5) \leftarrow \\ &E(4,6) \leftarrow \\ &E(4,7) \leftarrow \\ &Q() \leftarrow \\ &T(?x, ?y) \leftarrow E(?x, ?y) \\ &T(?x, ?x) \leftarrow \\ &T(?x, ?z) \leftarrow T(?x, ?y)T(?y, ?z) \end{aligned} \tag{2}$$

$\}$, which contains both facts and other rules.

With this, we have introduced all the needed elements for the syntax of datalog, but we have not yet spoken of the semantics of datalog. What is the program P supposed to express?

We have noted the similarities between datalog and first-order logic when defining the syntax of datalog. Nonetheless, neither rules nor programs are direct elements of first-order logic. It is however not difficult to bring them into a form, where we can interpret them as elements of first-order logic and use the semantics of first-order logic for it.

The symbol \leftarrow used in the definition of rules appears to be very similar to the implication \rightarrow from first-order logic so we can use this fact to transform any rule into a first-order formula. This formula has still free variables so that we universally quantify over all remaining variables to gain a sentence. After doing this we can consider any program as a finite first-order theory.

Example 2 (continued). *The equivalent sentence for the fact $E(1,2) \leftarrow$ is*

$$E(1,2)$$

and for the rule $T(?x, ?z) \leftarrow T(?x, ?y)T(?y, ?z)$ is

$$\forall ?x, ?y, ?z. T(?x, ?y) \wedge T(?y, ?z) \rightarrow T(?x, ?z)$$

The semantics of a first-order theory or sentence are its models. Unfortunately, a theory may have multiple models as the following example shows for the theory for P .

Example 2 (continued). *Any fact must be true in a model, but the implications allow us more freedom. As both E and T are binary predicates, we can represent them as a graph. We use blue, when both E and T are present, green for only T and orange for E*

The first model expresses T as the reflexive transitive closure of the input E , whereas the second model expands the first model and adds more additional atoms to a model.

Such an ambiguity is not ideal when we want to compute the query results of datalog. Some fragments of logic programming allow multiple models such as answer-set programming, but for datalog, we want a unique model. This is known as the least model. For this, we need to define what least means and show that it exists.

We call a set of ground atoms an *interpretation*. An interpretation represents all true atoms. We want to define the model property from first-order logic for this definition and show then that there exists a least model according to the subset relation.

Recall from first-order logic that $\forall x.\phi(x)$ holds in a structure \mathcal{A} if any element $a \in \mathcal{A}$ we have that $\phi(a)$ holds. As we do not have any function symbols all elements in the universe are constants, so instead of using the universal quantifier we could just create rules by replacing variables with all possible constants. Formally, this is defined using *groundings* or *instantions* which are functions

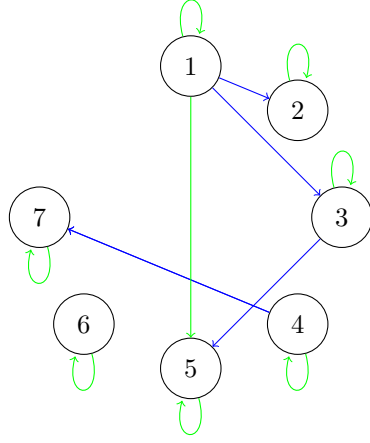


Figure 1: First model

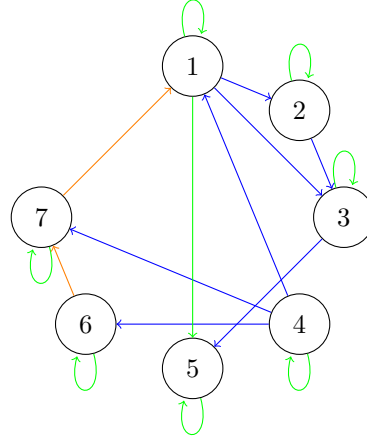


Figure 2: Second model

that map variables to constants. We can apply a grounding g to an atom by replacing every variable v in the terms by $g(v)$ and apply g to a rule by applying it to the head and every atom in the body. At the end of this, we have replaced every variable by a constant and have gained a ground rule. The ground program $\text{ground}(P)$ of a program P is the set of all ground rules that are the result of applying some grounding to a rule from P

We call a ground rule r *true* in an interpretation I if whenever $\text{Body}(r)$ is a subset of I , then also $\text{Head}(r)$ is in I . We call an interpretation I a *model* of a program P if every rule of $\text{ground}(P)$ is true in I .

So now we have defined models and can define the least model as well. We still need to show that the least model exists for which the following lemma is helpful.

Fact 1. *Let M_1, M_2 be two models of a program P . Then also $M_1 \cap M_2$ is a model of P .*

Therefore the intersection of all models is a model as well and due to the properties of the intersection it is least according to the subset relation.

In total, we call

$$\bigcap_{M \text{ is model of } P} M$$

the *least model* of P and refer to this characterization as the *model-theoretic semantics* of P .

Example 2 (continued). *In this example, the first model is actually the least model and the second model is just some other model.*

Therefore the semantic of this program is whenever $Q()$ is given, the reflexive-transitive closure of E . If $Q()$ is not given then it is only the reflexive closure.

Additionally, we see that the rules for T encode general rules whereas the other rules are more a specific input. We might want to reuse these rules for many different instances of E , but so far we also need to write a new program.

The example raises questions about the reusability of a program. Additionally, we talked in the introduction about database queries but never talked about databases.

We consider a *database* as a set of ground atoms similar to an interpretation. What is now the semantics of a program P and a database d ? We simply add every element of d as a fact to P and reuse the previous semantics. Additionally, we can also move all the specific facts from the program into the database and gain a reusable program. An alternative model definition for a pair of P and d is therefore: An interpretation I is a model for P and d if every ground atom from d is in I and every rule from $\text{ground}(P)$ is true in I . fact 1 holds again and we reuse the definition for the database and program case.

Both views are equivalent. We have shown how we can simulate a database by a program. We can simulate the program case with the program and database case by simply using an empty database.

We have now defined the semantics and shown a connection with databases. But this definition is not exactly ideal. In order to find the semantics of a program we would need to check all interpretations and then intersect them all. This is computationally expensive. Is there a simpler way?

We can define a function that computes the least model in a more direct way. For this, we consider the case where only the program P is present.

Consider an interpretation I . We call a ground atom a an *immediate consequence* of I if there exists a ground rule r in $\text{ground}(P)$ with the head a and $\text{Body}(r) \subseteq I$.

The immediate consequence operator T_P adds all immediate consequences to a model.

$$T_P(I) = \{a \mid a \text{ is an immediate consequence of } I\}$$

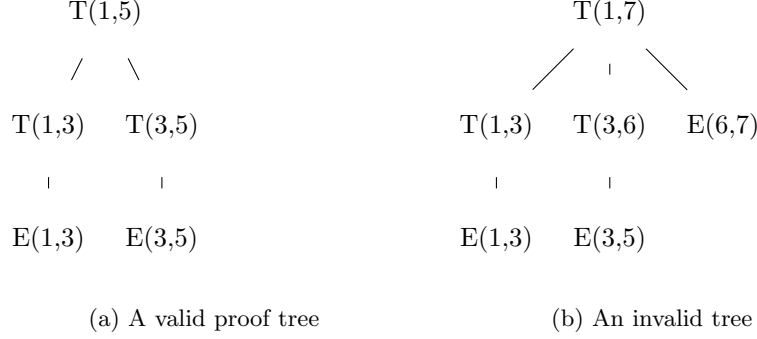
A fixed point of a function f is an element k such that $f(k) = k$. The repeated application of T_P starting from \emptyset yields a fixed point that is equal to the model-theoretic semantics of a program P .

Example 2 (continued). *We consider again the program P .*

Due to the fact $E(1, 3) \leftarrow$, $E(1, 3)$ is an immediate consequence of \emptyset . Applying T_P again, we have that $T(1, 3)$ is an immediate consequence of $\{E(1, 3)\}$ due to the rule $T(?x, ?y) \leftarrow E(?x, ?y)$ with the grounding

$$g(v) = \begin{cases} 1 & \text{if } v = ?x \\ 3 & \text{else} \end{cases}$$

Applying this to every fact, we gain an interpretation I that contains $T(1, 3)$ and $T(3, 5)$. Then $T(1, 5)$ is an immediate consequence of I due to the rule $T(?x, ?z) \leftarrow T(?x, ?y)T(?y, ?z)$ and the grounding



$$g'(v) = \begin{cases} 1 & \text{if } v = ?x \\ 3 & \text{if } v = ?y \\ 5 & \text{else} \end{cases}$$

The least fixed-point of T_P is called the *fixed-point semantics* of P and is the basis for most implementations of datalog reasoners.

Our goal is to explain why an atom is in the datalog result. The third important semantic of datalog helps here.

A tree t of ground atoms is a proof tree for a ground atom a in a program P and database d if the following three conditions hold:

1. the root of t is a
2. for every node n and its children l in t one of the following two conditions holds:
 - (a) $n \leftarrow l_1 \wedge \dots \wedge l_n$ for $l_1, \dots, l_n \in l$ is a ground rule from $\text{ground}(P)$, or
 - (b) n is a leaf and n is in the database.

Example 2 (continued). *The tree in section 2.1 is valid. The leaves of this tree are facts and all other nodes represent ground rules from $\text{ground}(P)$ in the previous step.*

The tree in fig. 3b is not valid. $E(6,7)$ is neither a fact nor in the database. Additionally there is no rule in P that can result into $T(1,7) \leftarrow T(1,3) \wedge T(3,6) \wedge E(6,7)$.

The proof-theoretic semantics of a program P and database d is the set of ground atoms that have a valid proof tree. Again this can be shown to be equal to the other semantic definitions. We will formally prove the equality of the proof-theoretic and the model-theoretic semantics of datalog in this work.

In the rules of the program P we note a special rule of the form $T(?x, ?x) \leftarrow$. It is a fact, but not a ground rule and the only rule where this is the case. We



Figure 4: A conventional algorithm

call a rule $H \leftarrow B_1 \wedge \dots \wedge B_n$ *safe*, if every variable that occurs in the head also occurs in the body, i.e.

$$\text{Vars}(H) \subseteq \bigcup_{i \in \{1, \dots, n\}} \text{Vars}(B_i)$$

We call a program safe if every rule in the program is safe. Safe programs are considered better because their result does not depend on the set of constants C which is in practice often either not given or infinite, but only on the specific database and/or the facts in the program.

Example 2 (continued). *We can transform P into the safe program P' by adding a new unary predicate N to the predicate symbols. P' is then the union of $P \setminus \{T(?x, ?x) \leftarrow\}$ and the following rules:*

$$\begin{aligned} N(?x) &\leftarrow E(?x, ?y) \\ N(?y) &\leftarrow E(?x, ?y) \\ T(?x, ?x) &\leftarrow N(?x) \end{aligned} \tag{3}$$

The new predicate N (for nodes) represents any elements that occur in the E relation and is used in the body for the reflexive rule.

If some constants are desired that do not occur in any E relation, one can also directly encode that into the database.

What are the semantics of this newly created safe program P' ? It turns out, that it is equal to the semantics for P for all original predicates, i.e. if we remove all ground atoms that use N we gain the same result. Therefore we state that any program can be transformed into an equivalent safe program.

2.2 Certifying algorithms

Most algorithms one encounters either in books like [12] or in real programs are comparable to fig. 4. We have an input, give this into an algorithm which is a black box for us and then we receive a result as an output. Without inspecting the implementation we have to trust that the results are correct. An alternative to this are *Certifying algorithms* depicted in fig. 5. They offer an explanation in addition to the result that can be checked independently by the checker or the user themselves. We follow the presentation in [21] which offers information far beyond this section as well.

Recall a definition of the class NP from complexity theory:

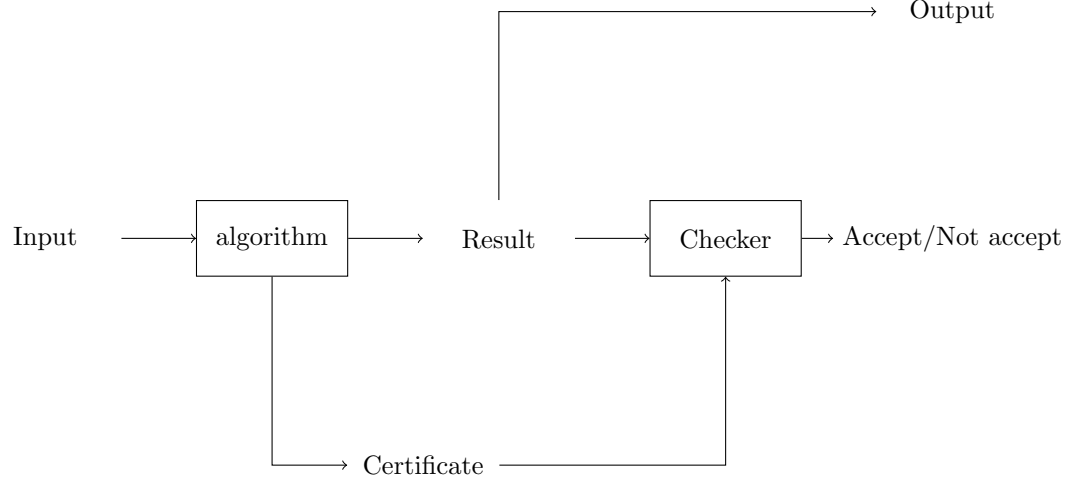


Figure 5: A certifying algorithm algorithm

Definition 1 ([3]). *A language $L \subseteq \{0, 1\}^*$ is in NP if there exists a polynomial $p : \mathbb{N} \rightarrow \mathbb{N}$ and a polynomial-time Turing machine M such that for all $x \in \{0, 1\}^*$:*

$$x \in L \Leftrightarrow \exists u \in \{0, 1\}^{p(|x|)} \text{ s.t. } M(x, u) = 1$$

If $x \in L$ and $u \in \{0, 1\}^{p(|x|)}$ satisfy $M(x, u) = 1$, then we call u a certificate for x

We can generalize this and require an algorithm not only to give us an output x but also a certificate u as a reason why x is the correct output. Then we can either check by ourselves that x is a correct solution according to u or use another program that checks this. As verifying a solution is never harder than computing it, the checker is usually simpler so that we either see directly that the checker is correct or we can formally verify it.

The formal framework of [21] defines certifying algorithms in the following way. We consider the set X of input values and the set Y of output values of a function f . A predicate $\phi : X \rightarrow \{\text{true}, \text{false}\}$ states a precondition for the inputs and another predicate the postcondition $\psi : X \times Y \rightarrow \{\text{true}, \text{false}\}$, ϕ allows us the algorithm to only accept part of the input space and $\psi(x, y)$ typically expresses that y is a valid output for the input x . In cases where x is not a valid input, we use the new symbol \perp to denote that the algorithm does not return anything and denote by $Y^\perp = Y \cup \{\perp\}$. The certificate or *witness* from a set W and its correctness is expressed by the predicate $\mathcal{W} : X \times Y^\perp \times W$ with the following properties:

1. **Strong witness property:** Consider a triple (x, y, w) that satisfies the witness predicate \mathcal{W} . If $y = \perp$, i.e. the input is not valid, we want w to

be a proof of this fact. Else we have that $y \in Y$ and we want w to be a proof that the postcondition is satisfied, i.e.

$$\forall x, y, w (y = \perp \wedge \mathcal{W}(x, y, z) \rightarrow \neg\phi(x)) \wedge (y \in Y \wedge \mathcal{W}(x, y, z) \rightarrow \psi(x, y))$$

2. **Simplicity:** This statement above has a simple proof
3. **Checkability:** It is possible to check efficiently if $\mathcal{W}(x, y, z)$ holds for a triple (x, y, z)

A *(strongly) certifying algorithm* is an algorithm that stops on all inputs $x \in X$ and returns a tuple $\langle y, w \rangle$ such that $\mathcal{W}(x, y, w)$ holds.

This is explained in the following examples. We start with an example from graph theory.

Example 3 ([21]). *Consider the problem of deciding whether a graph $G = (V; E)$ is bipartite, i.e. there exists a partition of V into V_1, V_2 with $V = V_1 \cup V_2$ and $V_1 \cap V_2 = \emptyset$, such that for all edges $e \in E$ we have that $e \cap V_1 \neq \emptyset$ and $e \cap V_2 \neq \emptyset$, so that all edges are only between vertices that are in the different partitions.*

The set of inputs X is the set of strings over $\{0, 1\}$ and $\phi(x)$ holds whenever x encodes a graph. The postcondition is the following: If $y = \text{true}$, then x encodes a bipartite graph. If $y = \text{false}$, then x does not encode a bipartite graph.

Adopting the well-known algorithm for checking whether a bipartite graph, we can construct the following certifying algorithm.

1. *Check if x encodes a binary graph. If not return \perp and a witness that describes the problem in the encoding.*
2. *Explore the graph in depth-first search and color vertices along the path by alternating colors. If we want to color a vertex by a color c and it already has the color c' , we continue exploring another path if $c = c'$, or stop if $c \neq c'$ and return false. If that is the case, we have a cycle of odd length in our current path and return this cycle as a witness.*
3. *If we reach this step, then all vertices are colored in a way that no neighboring vertices have the same color. Then we return true and the two sets of colored vertices as the witness.*

We know from graph theory that a graph is bipartite iff it has no cycle of odd length.

For any value of y a checker can efficiently check if the returned reason does indeed hold for x .

The above example illustrates the example well for a decision problem, but we want to decide whether a set of ground atoms is the semantics of a datalog program and database, i.e. a function problem. The next example illustrates a certifying algorithm for a function problem in number theory.

Example 4 ([21]). Let X be the set of pairs of natural numbers and Y the set of natural numbers. We want to compute the greatest common divisor, gcd , for two numbers a and b that are not equal to zero, i.e. the largest $g \in \mathbb{N}$ that is a divisor of both a and b . The precondition $\phi(\langle a, b \rangle)$ is that both a and b are different from zero and $\text{psi}(\langle a, b \rangle, g)$ shall be true, if g is $\text{gcd}(a, b)$.

The normal way of computing this involves the Euclidean algorithm, but this is not certifying. The extended Euclidean algorithm however returns in addition to the greatest common divisor g two numbers s and t such that $g = \text{gcd}(a, b) = as + bt$. In [21], the back direction is proven, i.e. that if g is a divisor of a and b and $g = as + bt$, then g is also the greatest common divisor of a and b .

Therefore a certifying algorithm in order to compute the gcd is the extended euclidean algorithm, that returns as y $\text{gcd}(a, b)$ and as the certificate the pair $\langle s, t \rangle$.

A checker then only has to check that $g = as + bt$ hold and that g divides both a and b .

For decision problems, we require an explanation when the input is in a language as well as an explanation when the input is not in the language. A common misconception is that therefore certifying algorithms are only possible for problems that are known to be in $NP \cap coNP$, which is presumed to not include the interesting NP -complete problems. This is not the case as we do not require the certificate to be polynomial in the size of the input. Indeed, SAT-solvers that decide the NP-complete satisfiability problem of propositional logic offer certificates in the DRAT proof format as a model or a proof of unsatisfiability[23].

In the datalog case, many modern reasoners such as Nemo or Soufflé already offer certificates for atoms in the form of the previously introduced proof trees. We therefore focus on implementing a checker for datalog.

In [2] an approach to develop checkers for certifying algorithms is outlined. There the checker is implemented in C because already the certifying algorithm itself is written in C++, so that both versions are similar. In order to verify the correctness of the checker they employ the proof assistant Isabelle. The C-code is exported by a tool into Isabelle so that the verification can take place. In Isabelle then the problem is defined and it is proven that the export does fulfill the desired properties.

In this workflow, we have to trust the correctness of the hardware, the C compiler, the tool that exports C code to Isabelle and of Isabelle itself. Additionally, we have to trust that we have correctly defined and specified the problem in Isabelle.

In this work, we modify this workflow a bit. We use a different proof assistant, Lean, instead of Isabelle. Lean not only a proof assistant but also a functional programming language. Therefore we can implement the checker directly in Lean and do not have to export a model of our code into the proof assistant. Our trust base is therefore smaller, as we no longer need an additional compiler for the language the checker is written in nor a tool to export this into the proof assistant.

2.3 Lean

We introduce in this section Lean, a programming language and proof assistant. As of writing, the current version is Lean 4. A more in-depth introduction can be found in [17] or [7].

Lean’s core is a small trusted kernel[14], that captures the most important functionalities and can be extended by the user. Since version 4 Lean is implemented in Lean itself[13]. The most important libraries for Lean that we will use are the standard library Std4[9] which contains important data structures and tactics and mathlib4[8], which contains definitions, tactics and theorem from diverse areas of mathematics and computer science.

As we have just heard, Lean can encode many mathematical results. The foundations of mathematics are often built on top of set theory (e.g. [15]) but most proof assistants instead use *type theory*. Type theory was introduced by Bertrand Russel as an alternative foundation during the foundational crisis of mathematics.

Any element, a *term*, has a *type* in type theory. This is denoted by `term: type`.

Example 5. *Different terms in their types are displayed below:*

```
42:nat
true:bool
sort([1,2,3]): List (nat)
nat: Type
Type: Type1
```

nat is a natural number, *true* is a boolean, the result of the application of *sort* to *[1,2,3]* has the type *List nat*. All previous types have a type as well, *Type*. *Type* itself also has a type *Type₁*. This forms an infinite sequence of type universes, that are non-cumulative, i.e. no term has multiple types.

In contrast to set theory, functions are a direct element of type theory and do not need a complicated encoding. For any types A and B, there exists the type of functions from A to B, $A \rightarrow B$.

We can define functions in Lean by the keyword `def` similar to other functional programming languages, such as the `id` function for natural numbers. In the parentheses, we denote an argument `x` with its type. Due to currying a function can have multiple arguments. After the colon, we denote the returned type of the function, which is here again a natural number. Therefore we have the following type $\mathbb{N} \rightarrow \mathbb{N}$, as the function takes a natural number as an input and returns a natural number as well. After the `:=` we denote the natural number this function returns.

```
def id (x: ℕ): ℕ := x
```

Everything we have seen so far holds in general in type theory. Lean itself is based however on a specific type theory, the Calculus of Inductive Constructions (CIC), a dependent type theory[13, 11].

The `id` function we defined above is correct but only works for natural numbers. We would have to implement another `id` function for booleans, string or any other type. This is rather cumbersome because we never actually use any properties of the specific type. We can generalize this function to help us with this problem.

```
def id (A: Type)(a: A): A := a
```

This `id` function has now two arguments. The first argument is the type we want to use it on, the second argument is an element of this type. `id N` works as previously and has the type $\mathbb{N} \rightarrow \mathbb{N}$. It seems less clear which type `id` itself has because this depends on the first argument. Such types are called *dependent types* and we would denote this specific type as $\prod (A: \text{Type}) A \rightarrow A$, known as a *Pi type*. Intuitively, a Pi type is a function that returns a type for any input value.

In practice, we often do not want to write down too much. We can write the function instead also as:

```
def id {A: Type}(a: A): A := a
```

Now the type `A` is an implicit argument, which is denoted by the braces, and Lean can fill it in for the user based on the input `a`. Therefore the user can write just `id 42` instead of `id N 42`.

So far we have discussed what types are and how we can define functions that are important for the programming aspect of lean, but this does not tell us much about the use as a proof assistant. Due to the Curry-Howard correspondence a proof of statement can be equivalently seen as a function that has the type of the statement. This type of statements is known in Lean as `Prop`. There exist functions like `Or: Prop → Prop → Prop` and the dependent types form the quantifiers. We view the Pi-type $\prod(a : A), \beta(a)$ as a function mapping `a` to the true statements of $\beta(a)$, which is equivalent to the universal quantor. `Prop` therefore has elements such as $\forall (x : \mathbb{N}), 0 \leq x$ or $\exists (x y : \text{Bool}), x = y \vee \neg x = y$.

Every statement defines its own type, whose elements are the proofs for it. In contrast to other types, all elements of these types are equivalent which is known as *proof irrelevance*, so that elements of these types only serve as a witness for their truth.

Proofs can be constructed in Lean in two ways. We start by writing `theorem` (or lemma, proposition etc. in `mathlib`) and denote it as we did for a function. The main difference is that the type is what we want to prove and that the function can not be executed.

```
theorem testTheorem {A: Type}:  $\forall (x y z: A), x = y \rightarrow \text{id } y = z \rightarrow$   

   $z = x$  := by  

  intro x y z h1 h2  

  unfold id at h2  

  rw [h1]  

  apply Eq.symm  

  exact h2
```

The proof is done in tactic mode. We open the tactic mode with the keyword **by**. At the start our context only contains that A is a type and the goal is $\forall(xyz : A), x = y \rightarrow idy = z \rightarrow z = x$, which is compactly written as

$$\{A : Type\} \vdash \forall(x\ y\ z : A), x = y \rightarrow id\ y = z \rightarrow z = x$$

We apply tactics to change the goal or a hypothesis in the context. These may introduce new goals or fulfill old goals. A proof is finished, when all goals are proven. We explain the proof now in more detail.

1. We start with the **intro** tactic. This moves elements that are universally quantified or the start of an implication from the goal into the context. Alternatively, one can use **revert** to move a hypothesis from the context back to the goal to prove a stronger statement.

At the end, the state is:

$$\{A : Type\}(x\ y\ z : A)(h1 : x = y)(h2 : id\ y = z) \vdash z = x$$

2. After that we use the **unfold** tactic at the hypothesis $h2$. This replace id with its definition in $h2$, which now looks like this: $(h2 : y = z)$
3. We use now the tactic **rw**, i.e. rewrite, that replaces in the goal x by y due to the hypothesis $h1$ so that the goal is now $z = y$
4. The goal is now almost the same as $h2$, but the order is wrong. Therefore we use the **apply** tactic with

$$\text{Eq.symm} := \forall\{A : Type\}(ab : A), a = b \rightarrow b = a \ .$$

Apply tries to match the statements consequent with our goal and opens new goals for the antecedents of this statement.

Our current proof state is this:

$$\{A : Type\}(x\ y\ z : A)(h1 : x = y)(h2 : y = z) \vdash y = z$$

5. Our goal is now exactly the same as the hypothesis $h2$. The **exact** tactic uses $h2$ to complete the goal.

Other important tactics not seen here are **by_contra** which allows a proof by contradiction, **by_cases** which allows a case distinction or **constructor** that splits an and statement into two goals. Lean offers also methods that try to prove goals themselves like **simp** that uses lemmas marked with **@simp** or **tauto**, that can recognize some tautologies. The set of tactics is however not fixed and the user can introduce new tactics to simplify the reasoning process.

The majority of formal proofs in this work are done in tactics mode, but we see above that describing them in natural language is quite verbose. Therefore we will only describe the idea we followed outside of this section.

These proofs are done backwards, i.e. starting from the goal towards the assumptions. There are also forward proofs that start from the assumptions by building hypotheses until we reach the goal or by composing theorems like functions. While this is possible, it was rarely done in this work because. A good source for this is either [17] or [4].

We can also use `def` to define new types. Sets of type `A` are in Lean implemented as functions from `A` to `Prop`. Functions to `Prop` are in general not computable so that the membership in a set is also not computable in contrast to for example lists.

Transform the tactic proof into the forward style.

```
def Set (α : Type) := α → Prop
```

This introduces a new type. If we want to just use functions that accept contain elements of type `Set`, using `def` would create a new type for which we would have to define again the set operations \cup or \cap . In this case, one can use the `abbrev` command which provides a simple alias.

```
abbrev NatSet := Set ℕ
```

The CIC and Lean allow the creation of types by induction. We can create our own implementation of the natural numbers with the `inductive` keyword. After that, we list the constructors. These can be constant constructors such as `zero` or function creators like `succ`. An inductive type may have many constructors of both types or even none as the empty type.

```
inductive myNat: Type
| zero: myNat
| succ: myNat → myNat
```

Our definition states that an element of `myNat` is either `zero` or the result of applying `succ` to some other element of `myNat`. Any element of `myNat` is only the result of one of these constructors and all elements of `myNat` are distinct.

A typical mathematical definition for this would be: `myNat` is the smallest set S of elements such that it contains `zero` and whenever an element a is in S , then also `succ(a)` is in S .

We can define for inductive types functions recursively like the add function.

```
def myNat.add (n m: myNat): myNat :=
match m with
| zero => n
| succ m' => succ (add n m')
```

If we want to prove a statement about elements on an inductive type, we can use the `induction` tactic to create proofs similar to those by (structural) induction in natural language proofs. If no induction hypothesis is needed one can also use the `cases` tactic that creates a goal for every constructor

```
theorem myNat.add_zero (n: myNat): myNat.add zero n = n := by
  induction n with
  | zero =>
```

```

      unfold add
      rfl
| succ n' ih =>
  unfold add
  rw [ih]

```

We start with the induction tactic to perform an induction over n . Each constructor is listed separately and after the arrow, we start the proof for this constructor. In the zero case, it is enough to unfold the definition of `add` to get `zero=zero` and use `rfl` which proves these equalities.

In the `succ` case, we have two values: the `myNat` element n' we use in the constructor of `succ` and also the induction hypothesis `ih: add zero n' = n'`. We can unfold the definition of `add` again and use `rw` to replace `add zero n'` by `n'` in the goal `succ (add zero n') = succ n'`. The `rw` tactic applies `rfl` at the end automatically to finish the proof.

After defining the natural numbers inductively, we can also define the even numbers by induction. We again have the zero constructor and a `succ` constructor that now shall represent the next natural number.

```

inductive evenNat: Type
| zero: evenNat
| evenSucc: evenNat → evenNat

```

While this works, we see no relation between the elements of `myNat` and `evenNat`. We can of course define a function that maps elements of `evenNat` to `myNat` and use this to transform any even number to a natural number.

```

def evenNatToMyNat (e: evenNat) :=
  match e with
  | zero => myNat.zero
  | succ e' => myNat.succ (myNat.succ (evenNatToMyNat e'))

```

This approach works but is a bit cumbersome. We always need to type this ourselves and if we forget it Lean will complain. There exists however the `class` `Coe`. A class describes an abstract property e.g. of a type and can be implemented by a specific type via `instance`.

The `Coe` looks (in a simplified version) like this. It takes two types and a function that maps from one type to the other. We can show that this property holds for `evenNat` and `myNat` by using `evenNatToMyNat` to define an instance of this class. Then we can use elements of `evenNat` in functions defined for `myNat` and Lean will automatically convert them using `evenNatToMyNat`.

```

class Coe (α : Type) (β : Type) where
  coe : α → β

```

```

instance coeEvenNatMyNat: Coe evenNat myNat := ⟨evenNatToMyNat⟩

```

An alternative way of defining types are structures that are comparable to classes in object-oriented languages. Structures have fields that have a type and

we start listing them one by one after the `where` statement. Structures are like inductive types with a single constructor. Therefore they are not allowed to be recursive, i.e. no structure A may have an element of type A , because there would be no base case. We can build an element of the type of the structure by passing the elements in braces and can use the name of the field to access an element.

```
structure player where
  (name:String)
  (numGoals: ℕ)
  (active: Bool)

def player1: player := {
  name := "Test",
  numGoals := 10,
  active := true
}

def player1Goals: ℕ := player1.numGoals
```

Returning to our example of `myNat`, we can define some more functions. In the previous section, we talked about the Euclidean algorithm. We can try to implement it ourselves in Lean. Firstly, we would require the modulo function for that. We will only define it for now to not lose focus and use the keyword `sorry` in the value. This closes in proofs any goal or can be used for incomplete functions, but will throw an error if executed.

```
def myNat.mod (n m: myNat): myNat := sorry
```

The Euclidean algorithm can be implemented in the following way:

```
def myNat.Euclid (a b: myNat): myNat :=
  if b = zero
  then a
  else Euclid b (myNat.mod a b)
```

Lean will not accept this function and instead complain in the line of the if that it failed to synthesize an instance of `DecidableEq` for `myNat`. `DecidableEq` allows us to use `=` for a type in a function. This is not a trivial requirement as for example the equality of real numbers is undecidable[22]. We need to provide a function to Lean that can be used to decide whether two elements of `myNat` are equal. During the definition of inductive types, we already noted, that all elements of an inductive type are different, which allows us to do this. Luckily, Lean can even do this alone if we tell Lean to do this using the `deriving` keyword. It can even derive multiple instances such as the `Inhabited` instance that states that there exists some default element of this type.

```
inductive myNat: Type
| zero: myNat
```

```
| succ: myNat → myNat
deriving DecidableEq, Inhabited
```

Now the error will disappear, but a new one will be introduced. Lean does not recognize the termination of `myNat.Euclid`. We have already defined recursive functions before and we did not encounter this problem. In `myNat.add` we however used the inductive schema and only did the recursion on n in the case of `succn`. This allowed Lean to conclude that we only call the function on smaller elements so that the function will terminate.

To show termination, one is required to define some *well-founded relation*, i.e. a relation that does not have any infinite descending chains. Such a relation would be the less-than relation of the natural numbers so it is often convenient to define some size measure that measures the size of our elements in the natural numbers and use this. Such a function exists automatically for any inductive type and is called `sizeOf`. We can provide this and are only left to prove that `sizeOf (mod a b) < sizeOf b`, which we will leave open here.

```
def myNat.Euclid (a b: myNat): myNat :=
  if b = zero
  then a
  else Euclid b (myNat.mod a b)
termination_by sizeOf b
decreasing_by
  simp_wf
  sorry
```

An alternative to this termination proof would be to mark this function using the `partial` keyword, which denotes that this function does not always terminate. This is however often not beneficial as we no longer can no longer use the `unfold` tactic to gain the definition of this function in a proof.

We want to take a look at another example of an inductive type, that we will often use namely lists. A list (here `List'` as `List` is already a standard definition) is either the empty list or the result of combining a list with a new element and we can define a get function

```
inductive List' (A: Type): Type
| nil : List' A
| cons : A → List' A → List' A
deriving DecidableEq

def getElement (A: Type) [Inhabited A] (l: List' A): A :=
match l with
| nil => Inhabited.default
| cons hd _ => hd
```

We see in this definition a new type of argument. We want to return an element of the list. If the list is empty, this proves to be difficult. In that case, we return the default element that exists because A is an instance of `Inhabited`

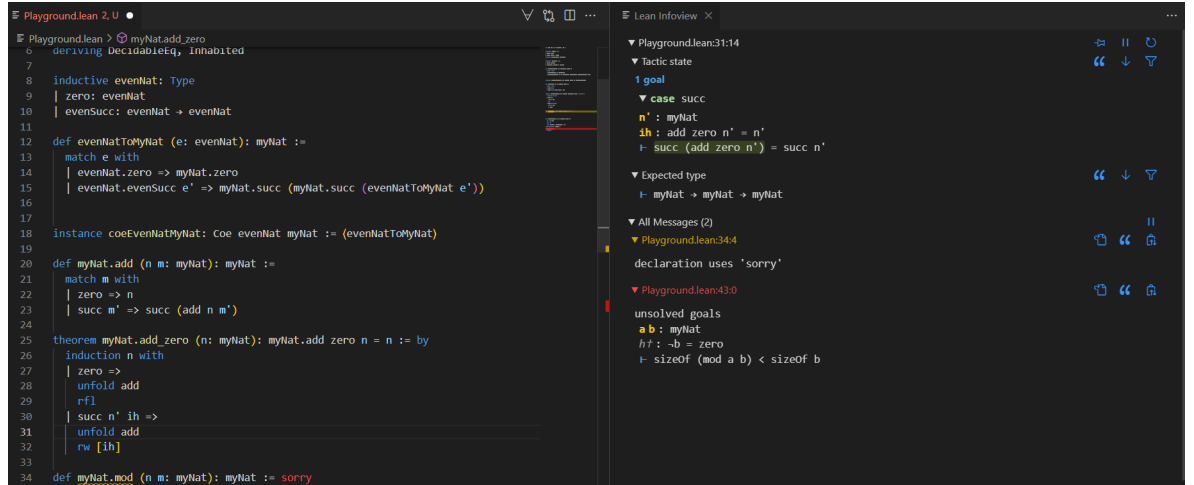


Figure 6: An example of Lean in practice. On the left side is the code seen and on the right side the context in a proof with the current goal

. Such a type-class parameter is given in brackets and Lean will look for the instance when using this function.

This may not always be the desired behaviour and maybe we want to return nothing if the list is empty. In this case, we can use the `Option` Type. It has two constructors, one that contains the element and the other symbolizes that there is nothing. Here α has the type `Type u` that symbolizes it works for any type universe.

```
inductive Option ( $\alpha$  : Type u) where
| none : Option  $\alpha$ 
| some (val :  $\alpha$ ) : Option  $\alpha$ 
```

We will also often use the Exception type. This uses two types and returns one in the error case and the other in the normal case.

```
inductive Except ( $\varepsilon$  : Type u) ( $\alpha$  : Type v) where
error :  $\varepsilon \rightarrow$  Except  $\varepsilon$   $\alpha$ 
ok :  $\alpha \rightarrow$  Except  $\varepsilon$   $\alpha$ 
```

3 Formalization of Datalog

In the previous section, we introduced datalog. Our goal is to check whether ground atoms are the result of correct datalog derivations. In the pursuit of a proof of correctness for our algorithms, we need to know what correct derivations are which we solve by formalizing the syntax and semantics of datalog in Lean. As of writing, there is to the best of our knowledge no formalization of datalog in Lean yet.

There does exist a formalization of datalog in Coq[5]. The formalization includes the syntax of datalog and the fixed point semantics of datalog with a certified datalog engine. Similarly, we will also formalize the syntax of datalog. After that, the paths will diverge as we are interested in the proof-theoretic semantics of datalog to check proof trees. Additionally, we will also formalize the model-theoretic semantics of datalog for completeness arguments and to have extra security that our formalization holds by proving that both semantics are equivalent.

3.1 Syntax

We recall from the preliminaries that an atom is of the form $A(t_1 \dots t_n)$ for sets of predicate symbols, variables and constants P, V and C and an arity function $ar : P \rightarrow \mathbb{N}$. If we were to directly formalize this an atom would be defined in the following way, where we use types instead of sets.

```
def atom (C: Type) (P: Type) (V: Type) (ar: P → ℕ): Type :=  
  sorry
```

Such a definition is rather verbose with already four arguments. Anything like the semantics that will use atoms will require even more inputs. To have a more compact representation we reuse the definition of a signature that shrinks the number of arguments to just one. We use types instead of sets as this is more natural in type theory. We can consider the type as a set. If we were to use sets we would have to decide already which type these sets should have which seems unclear for now. This allows us to instead define later the types we want to use.

The formalization showed us that the requirements of countability of the sets P , C and V were not required. Any result we wanted to prove holds already in this general case. Therefore we forego of modelling this assumption.

A `signature` is then a structure that has a type for constants, variables¹ and predicate symbols and an arity function.

```
structure signature where  
  (constants: Type)  
  (vars: Type)  
  (predicateSymbols: Type)  
  (predicateArity: predicateSymbols → ℕ)
```

¹We use vars instead because variable(s) is a keyword in Lean

In the future, unless denoted otherwise, we will always use a fixed signature τ and assume that all types have the `DecidableEq` and `Hashable` properties to use Lean automatic derivation for later uses in the program.

Another requirement was that the sets of constants and variables are distinct. This proved again to be unnecessary in our formalization because we define terms as an inductive type with a constructor for the constant and the variable case. Therefore constants and variables in an atom will always be distinct.

```
inductive term ( $\tau$ : signature): Type
| constant :  $\tau$ .constants  $\rightarrow$  term  $\tau$ 
| variableDL :  $\tau$ .vars  $\rightarrow$  term  $\tau$ 
deriving DecidableEq, Hashable
```

For an atom we have fields for the symbol, the list of terms and a proof that the length of the list matches the arity of the symbol.

```
structure atom where
  (symbol:  $\tau$ .predicateSymbols)
  (atom_terms: List (term  $\tau$ ))
  (term_length: atom_terms.length =  $\tau$ .predicateArity symbol)
deriving DecidableEq, Hashable
```

Two structures of the same type are equal if all their fields are equal. Due to proof irrelevance, we gain the following expected criteria for the equality of atoms.

Lemma 1 (`atomEquality`). *For all τ -atoms a_1, a_2 , we have $a_1 = a_2$ iff their symbols and `atom_terms` are equal.*

Rules and programs can be transformed in straightforward way. Here we use lists

```
structure rule where
  (head: atom  $\tau$ )
  (body: List (atom  $\tau$ ))
deriving DecidableEq

abbrev program := Finset (rule  $\tau$ )
```

Next, we want to define the ground versions of our previous definitions. Groundings are simply the functions from variables to constants.

```
def grounding ( $\tau$ : signature):=  $\tau$ .vars  $\rightarrow$   $\tau$ .constants
```

We see multiple ways to define ground atoms. Firstly, we can define them like we defined atoms but use constants instead of terms.

```
structure groundAtom ( $\tau$ : signature) where
  symbol:  $\tau$ .predicateSymbols
  atom_terms: List ( $\tau$ .constants)
  term_length: atom_terms.length =  $\tau$ .predicateArity symbol
deriving DecidableEq, Hashable
```

Secondly, we can define ground atoms as a special type of atom by constructing a new structure consisting of an atom and a proof that for all terms exists some constant that is equal to it.

```
structure groundAtom (τ: signature) where
  atom: atom τ
  ground: ∀ (t: term τ), t ∈ atom.atom_terms → ∃ (c: τ.
    constants), t = term.constant c
```

The second variant allows us an easy conversion from ground atoms to atoms by simply returning the atom element. Also, we can convert atoms easily to ground atoms as soon as we have the proof. These conversions have to be written by hand in the first variant.

The first variant on the other hand allows us to define functions that create ground atoms more directly. We can take a grounding and just map the term list using this function without having to provide a proof. In the second variant, we have to first define the function on the atom level and then have to prove that this operation does not create any variables. This may sound like extra security in case we mess things up, but when defining the terms as a list of constants the type checker of the kernel does the check for us.

The number of conversions is rather limited whereas an easier way to define functions may be useful more often. Therefore we chose the first variant.

We start by defining these conversions that we now have to do manually. We can convert a ground atom to an atom by mapping every constant to a term via `term.constant`. The map operation does not change the length of a list of atoms so that the term length property stays true(`listMapPreservesTermLength`).

```
def groundAtom.toAtom (ga: groundAtom τ): atom τ := {
  symbol:=ga.symbol,
  atom_terms:= List.map term.constant ga.atom_terms,term_length
    := listMapPreservesTermLength ga
}
```

For later proofs, it is interesting to know that this is an embedding of the ground atoms into the atoms, i.e. that if two ground atoms are different then the result of their `toAtom` functions is also different.

For this, we need the following result. If two lists are equal then mapping both lists by the same function f results in the same list. If the function is injective also the back direction holds, which we prove by induction(`listMapInjectiveEquality`).

Lemma 2 (`groundAtomToAtomEquality`). *Let a_1, a_2 be two ground atoms. Then a_1 is equal to a_2 iff the result of `groundAtom.toAtom` of both is equal.*

Proof. If $a_1 = a_2$, then also their result is the same.

For the back direction, we know that the results of `groundAtom.toAtom` are equal and want to show that they are equal. We use a similar lemma as lemma 1 for ground atoms(`groundAtomEquality`). Therefore we have to show that their symbols and terms are equal. As `toAtom` does not change the symbol, the first claim follows. For the second claim we employ the fact the

constructors of an inductive type like `term.constant` are injective functions² and `listMapInjectiveEquality` to conclude that the terms are equal. \square

We can therefore employ `groundAtom.toAtom` safely as the type coercion from ground atom to atom.

So that we can convert atoms to ground atoms, it would be enough to use a proposition that says that all elements are constants. In later uses such as the definition of safety for rules, it will be beneficial to have a function that computes all variables that occur in an atom.

After defining the conversions we define ground rules similar to rules.

```
structure groundRule ( $\tau$ : signature) where
  head: groundAtom  $\tau$ 
  body: List (groundAtom  $\tau$ )
deriving DecidableEq
```

Think of a way to present the back transformation in a good way.

We can apply a grounding to a term by replacing a variable by its grounding result and keeping the constant.

```
def applyGroundingTerm (g: grounding  $\tau$ ) (t: term  $\tau$ ): term  $\tau$  :=
  match t with
  | term.constant c => term.constant c
  | term.variableDL v => term.constant (g v)
```

Using this function we apply groundings also to atoms and rules.

```
def atomGrounding (g: grounding  $\tau$ ) (a: atom  $\tau$ ): groundAtom  $\tau$  := {
  symbol := a.symbol,
  atom_terms := List.map (applyGroundingTerm' g) a.atom_terms,
  term_length := applyGroundingTerm'PreservesLength g a
}

def ruleGrounding (r: rule  $\tau$ ) (g: grounding  $\tau$ ): groundRule  $\tau$  := {
  head:=atomGrounding g r.head,
  body:= List.map (atomGrounding g) r.body
}
```

The ground program of a program P is the set of all ground rules that are the result of the application of a grounding to a rule from P .

```
def groundProgram (P: program  $\tau$ ) :=
  {r: groundRule  $\tau$  |  $\exists$  (r': rule  $\tau$ ) (g: grounding  $\tau$ ), r'  $\in$  P  $\wedge$  r =
    ruleGrounding r' g}
```

3.2 Semantics

After finishing the definition of the syntax, we start formalizing the semantics. We discussed in the preliminaries two ways for the semantics. We decided to

²This can be shown with the `injection(s)` tactic.

formalize the semantics with the database as this is more general. It is simpler to pass an empty database into the checker than to write every fact from the database into the rule file.

In this section, we do not want to deviate too much from the path by implementing databases in a complicated way. For now, a database is simply something that has a contains function that returns true if an element is in the database. This class can be implemented in multiple ways in the algorithm later.

```
class database ( $\tau$ : signature) where
  (contains: groundAtom  $\tau \rightarrow$  Bool)
```

We call an interpretation a set of ground atoms as in the preliminaries.

```
abbrev interpretation ( $\tau$ : signature)
et (groundAtom  $\tau$ )
```

We start by defining the proof-theoretic semantics as proof trees are our certificates in the checker so this is the most important semantic for us.

We require the notion of a tree to formalize the proof-theoretic semantics. In mathlib exists a definition of a tree³, but this is in spite of the name only a binary tree. Then we could only represent rules in the proof tree whose body has at most length two. While any datalog program can be transformed into an equivalent program that only contains rules whose body has a length of at most two, this is not desirable as rules in practice are often longer. We could require the user to just transform the rules into this form, but this seems to hinder the acceptance of this tool and may mask errors that only occur in optimizations for longer rules.

Therefore we start by defining a tree as an element t with as a vertex a and a list of subtrees l . We call a the root of t and the root of the trees in l as the children of t . A leaf is an element whose subtree list is empty.

```
inductive tree (A: Type)
| node: A  $\rightarrow$  List (tree A)  $\rightarrow$  tree A
```

```
def root: tree A  $\rightarrow$  A
| tree.node a _ => a
```

```
def children: tree A  $\rightarrow$  List A
| tree.node _ l => List.map root l
```

An important measure for trees is the height defined as the longest path from the root to a leaf. An alternative recursive definition is that it is the maximum of the height of all subtrees plus one, which we can implement with the listMax function.

```
def listMax {A: Type} (f: A  $\rightarrow$   $\mathbb{N}$ ): List A  $\rightarrow$   $\mathbb{N}$ 
```

³see at https://leanprover-community.github.io/mathlib4_docs/Mathlib/Data/Tree.html

```

| [] => 0
| (hd::tl) => if f hd > listMax f tl then f hd else listMax f tl

def height (t:tree A): ℕ :=
  match t with
  | tree.node a l => 1 + listMax (fun ⟨x, _h⟩ => height x) l.
  attach
termination_by sizeOf t
decreasing_by
  simp_wf
  apply Nat.lt_trans (m:= sizeOf l)
  apply List.sizeOf_lt_of_mem _h
  simp

```

Our implementation looks a bit different compared to the text. This is because we are required to prove the termination of this function. We call the height function recursively, but with a list function so that we no longer follow the inductive schema directly. We use the `List.attach` function that takes a list l and maps any element a from l to an element of $\{ a \mid a \in l \}$, so a pair of the original element and a proof that it is a member of l .

Using the `sizeOf` function, we prove termination by showing that `sizeOf x < 1 + sizeOf a + sizeOf l`. By transitivity, it is sufficient to prove that `sizeOf x < sizeOf l` holds. Because we used `l.attach` instead of simply `l`, we have a proof of $x \in l$ available and can use that the size of any list member must be smaller than the size of the list itself, which finishes the proof.

We call a tree t_2 a **member** of a tree t_1 if it occurs in the subtrees of t_1 and can prove that any member has a smaller height than the original tree (`heightOfMemberIsSmaller`) because the result of f for some member a of l is less-equal to the value of `listMax f l` (`listMax_le_f_member`).

With this, we finish the general results about trees and can return to formalizing the proof-theoretic semantics. A proof tree is a tree whose vertices are ground atoms.

```
abbrev proofTree (τ: signature):= tree (groundAtom τ)
```

Some proof trees will not represent valid derivations by our definition. The next step is to define a predicate that shows the validity of a tree similar to our definition in the preliminaries.

Again, we want to design our checker with the best compatibility in mind. In some papers (e.g. [6]) the leaves of a valid proof tree have to be database elements. We leave this open to allow also facts from the program to serve as leaves for those programs that do not come with a database.

We express the validness of a proof tree with the root a and the list of subtrees l with respect to a program P and a database d by a disjunction with two disjuncts. We call the first the *rule case*. There we require a rule r and a grounding g , such that r is in P , the rule grounding of r with g yields the ground rule we gain from the root and its children. Additionally, all subtrees must be

valid, which we again express via `List.attach` for the termination proof. The second case is the *database case*. There *a* must be a leaf and contained in the database.

```
def isValid (P: program  $\tau$ ) (d: database  $\tau$ ) (t: proofTree  $\tau$ ): Prop
  :=
  match t with
  | tree.node a l => (  $\exists$ (r: rule  $\tau$ ) (g:grounding  $\tau$ ), r  $\in$  P  $\wedge$ 
    ruleGrounding r g = {head:= a, body:= (List.map root l)}  $\wedge$  l.
    attach.Forall (fun <x, _h> => isValid P d x))
   $\vee$  (l = []  $\wedge$  d.contains a)
termination_by sizeOf t
decreasing_by
  simp_wf
  decreasing_trivial
```

We see from this definition that any element that is contained in the database has a simple proof tree. The tree is just the element as the root without any subtrees (`databaseElementsHaveValidProofTree`).

We do not consider input or export predicates so that the database elements are always part of the semantics. This also simplifies the definition for `isValid` in contrast if we only want elements that are the result of some rule to be in our semantics.

The proof-theoretic semantics with respect to a program *P* and database *d* is the set of ground atoms that are the root of a valid proof tree. We avoided earlier defining the types of a signature as a finite type so that we cannot expect a finite set here. We will manage to prove the same results and have to avoid proving that there are only finitely many ground atoms.

```
def proofTheoreticSemantics (P: program  $\tau$ ) (d: database  $\tau$ ):
  interpretation  $\tau$  :=
  {a: groundAtom  $\tau$  |  $\exists$  (t: proofTree  $\tau$ ), root t = a  $\wedge$  isValid P d t}
```

The proof-theoretic semantics provides with proof trees a good explanation of why an element is part of the solution. We only have to verify that the proof tree is correct. This however tells us only that we have found a subset of the solution. It tells us not whether there are no derivations possible. For this, the other semantics are more equipped for. They describe the solution as the least element of some set. If we verify that the result is in this set (i.e. the set of models or the set of fixed points), then our result is a superset of the solution. If both criteria hold, then we have exactly the solution.

In addition to these algorithmic goals, formalizing another semantics strengthens our formalization because we might spot some wrong assumptions we did the definitions that way.

Both the model-theoretic and the fixed point semantics are defined to be the least object of something. We decided to formalize the model-theoretic semantics because we can directly give the model. For the fixed-point semantics, we would first need to prove that the fixed point even exists and may need some

theorems about fixed points, but such a formalization (in Coq) can be found in [5].

We start by formalizing the criteria for a rule being true. An interpretation is a set whereas a body is a list. So that we can compare them, we define the `groundRuleBodySet` of ground rule r as the conversion of the body to a finite set. This operation preserves the members so that a ground atom is in the `groundRuleBodySet` of r iff it is in the body of r .

```
(groundRuleBodySet_iff_groundRuleBody)

def groundRuleBodySet (r: groundRule  $\tau$ ): Finset (groundAtom  $\tau$ ) :=
  List.toFinset r.body
```

This allows us to define the criteria for a rule being true in a natural way. Whenever all elements of the body are in the interpretation, then the head must be a member of the interpretation as well.

```
def ruleTrue (r: groundRule  $\tau$ ) (i: interpretation  $\tau$ ): Prop :=
  (groundRuleBodySet r).toSet  $\subseteq$  i  $\rightarrow$  r.head  $\in$  i
```

An interpretation is a model, if all rules from the ground rule are fulfilled. Additionally, it also needs to contain the elements from the database so that we will be able to prove the equivalence of the semantics later because all elements in the database are already in the proof-theoretic semantics.

```
def model (P: program  $\tau$ ) (d: database  $\tau$ ) (i: interpretation  $\tau$ ) :
  Prop :=
  ( $\forall$  (r: groundRule  $\tau$ ), r  $\in$  groundProgram P  $\rightarrow$  ruleTrue r i)
 $\wedge \forall$  (a: groundAtom  $\tau$ ), d.contains a  $\rightarrow$  a  $\in$  i
```

Now we are equipped with the necessary tools to formalize the model-theoretic semantics. In the preliminaries, we defined the semantics as the intersection of all models, i.e.

$$\bigcap_{M \text{ is model of } P} M$$

The corresponding operation to this intersection in Lean is called `Set.iInter`. Therefore we would need to transform the set of interpretations to an indexed family.

```
def iInter (s :  $\iota \rightarrow$  Set  $\alpha$ ) : Set  $\alpha$ 
```

We can instead define it more directly. Sets are equal whenever they have the same members by the principle of extensionality. Therefore we also try to find a formula that is true for an element whenever it is in $\bigcap_{X:\phi(X)} X$. We know that an element is in $X \cap Y$ whenever it is in both X and Y . We can generalize this to conclude that an element is in $\bigcap_{X:\phi(X)} X$ if it is in all sets X that satisfy ϕ .

```
def modelTheoreticSemantics (P: program  $\tau$ ) (d: database  $\tau$ ):
  interpretation  $\tau$  :=
  {a: groundAtom  $\tau$  |  $\forall$  (i: interpretation  $\tau$ ), model P d i  $\rightarrow$  a  $\in$  i}
```

We now have to prove that this is actually the least model. We start by showing that it is a subset of every model. This follows basically from the definitions.

Lemma 3 ([leastModel](#)). *Let P be a program and d be a database. For all models M of P and d , the model-theoretic semantics of P and d is a subset of M*

Proving that our definition is a model takes a bit more work.

Lemma 4 ([modelTheoreticSemanticsIsModel](#)). *For all programs P and databases d the model-theoretic semantics of P and d is a model.*

Proof. Let M be the model-theoretic semantics of P and d . We have to show that it fulfills both model criteria. We start by showing that all rules are true in M . We assume for a contradiction that there is a rule r that is not true, i.e. that the body set is a subset of M , but the head of r is not in M . Then there must be a model M' of P and d such that the head of r is not in M' . By lemma 3 we know that all elements of M must be in the model of M' . Therefore the body of r is a subset of M' and since M' is a model the head of r must be in M' which violates our assumptions.

Now we show that all elements in d are in M as well. The proof works in the same way as before but is simpler. Suppose that this does not hold. Then there is a database element a which is not in M . By the definition of M there must exist a model M' such that a is not in M' . But any model must contain all database elements which yields the contradiction and finishes the proof. \square

The remainder of this chapter is spent proving the equivalence of both semantics, i.e. the following theorem.

theorem [SemanticsEquivalence](#) (P: program τ) (d: database τ):
proofTheoreticSemantics P d = modelTheoreticSemantics P d

By the anti-symmetry of the subset operation, it suffices to show that both semantics are a subset of each other.

We start by proving $\text{proofTheoreticSemantics } P \ d \subseteq \text{modelTheoreticSemantics } P \ d$. We actually prove a stronger statement by showing that all elements in the proof-theoretic semantics are in any model. By lemma 4 the model-theoretic semantics are a model so that the proof follows.

Lemma 5 ([proofTreeAtomsInEveryModel](#)). *Let P be a program and d be a database. Let a be a ground atom in the proof-theoretic semantics of P and d . Then we have $a \in M$ for all models M of P and d .*

Proof. An element is in the proof-theoretic semantics whenever it is the root of a valid proof tree t . We prove this by strong induction on the height of t for all trees t and ground atoms a .

There are two cases when t is valid. If it is valid and in the rule case, then there exists a rule r and grounding g such that r is in P and the grounding of r ,

which we call r' , has the head a and the body l . All elements of l are the root of valid proof trees as well and by the definition of the height function have a smaller height. By the induction hypothesis therefore all elements of l are in M . Since r' is the result of applying a grounding to a rule from P , r' must be true in M . Therefore a is also in M .

In the database case, the root is an element of the database. Any element of the database must be in any model by definition. \square

For the back direction, it suffices to show that the proof-theoretic semantics are a model as well by lemma 3.

Lemma 6 (`proofTheoreticSemanticsIsModel`). *Let P be a program and d be a database. Then the proof-theoretic semantics of P and d is a model for P and d .*

Proof. As any element from the database d has a valid proof tree, the database is contained in the proof-theoretic semantics.

We need to prove that all rules from $\text{ground}(P)$ are true in the proof-theoretic semantics. Let r be such a rule and suppose that every element of $\text{body}(r)$ is in the proof-theoretic semantics. Therefore there exists a list of trees $t_1 \dots t_n$ such that all trees are valid and the list $\text{root}(t_1) \dots \text{root}(t_n)$ equals the body of r . Then we can build a new tree with the root $\text{head}(r)$ and the children $t_1 \dots t_n$. This tree is valid because all subtrees are valid and the root and its children are the instance of a ground rule of P . Therefore also $\text{head}(r)$ is in the proof-theoretic semantics and r is true. \square

We expand a bit on the proof above. The lemma we use to get the valid proof tree for $\text{head}(r)$ is called `createProofTreeForRule`. We know that all elements in the body have valid proof trees and need a list of these proof trees. We show in the lemma `getTree_Helper` a more general property from which this follows.

Lemma 7 (`getTree_Helper`). *Let A and B be types. Let l be a list of type A and S be a finite set of type A . Let f be a function from B to A and valid a property of elements of type B . If all elements from l are in S and for any element a in S there exists an element b such that $f(b) = a$ and b is valid, then there exists a list l' such that mapping l' with f yields l and all elements of l' fulfill the `isValid` predicate.*

Before proving the statement, we explain how we want to use this. A is supposed to be the ground atoms and B is supposed to be the proof trees. The function $f : B \rightarrow A$ is the root function, l is the body of the rule r and S is the finite set of the body elements. If we want to show that the head is in the proof-theoretic semantics, we already know that every atom a in the body has a valid proof tree whose root is a . Then we get the list of proof trees we can use as the children for the rule creation.

Proof of lemma 7. We prove this by structural induction on l . If l is empty then we can simply use the empty list for l' that fulfills the desired properties.

If l has the shape $hd :: tl$, then we can get a hd_b element for hd since hd in S . By the properties of S , hd_b is valid and is mapped to a via f . Any element in tl is still in S so that the induction hypothesis provides us a list tl' that maps via f to tl with only valid members. Then the list $hd_b :: tl'$ is the witness for l' . \square

This concludes the back direction and we finish the proof of the equivalence of the semantics and conclude this chapter.

Theorem 1 ([SemanticsEquivalence](#)). *For any program P and database d the model-theoretic and the proof-theoretic semantics of P and d coincide.*

4 Validating proof trees

After introducing the problem and modeling datalog in Lean, we now describe the algorithm to verify a solution partially. We focus in this section on the soundness by verifying a given proof tree. In the previous section, we introduced the following characterization for valid proof trees:

```
def isValid(P: program  $\tau$ ) (d: database  $\tau$ ) (t: proofTree  $\tau$ ):
  Prop :=
  match t with
  | proofTree.node a l =>
  (  $\exists$ (r: rule  $\tau$ ) (g:grounding  $\tau$ ),
    r  $\in$  P  $\wedge$ 
    ruleGrounding r g = groundRuleFromAtoms a (List.map root l)
     $\wedge$  l.attach.All2 (fun <x, _h> => isValid P d x))
   $\vee$  (l = []  $\wedge$  d.contains a)
```

This is a disjunction so we need procedures to decide for both disjuncts if they are true. The second disjunct consists of a database check and a check of list emptiness. Both can be done straightforwardly.

The first part is more interesting. Since we use there existential quantifiers, this is not computable and have to implement a function for this. While we can simply iterate over the program to check for the existence of an r , the number of groundings might be exponential or even infinite. Instead of using groundings, we want to use some more sophisticated to check this which we introduce next.

4.1 Substitutions

A grounding is a function from variables to constants. This means we always need to specify for every variable a constant that it is mapped to. This was good in the definitions to ensure that we always get a ground atom, but raises problems in the unification case as the following example demonstrates.

Example 6. Suppose we want to know whether we can unify the list of terms $l_1 = [?x, ?y]$ with the list of constants $l_2 = [a, b]$, i.e. we look for a function $f : V \rightarrow C$ such that mapping every element of l_1 with f yields l_2 .

We start by trying to unify x and a . If we use groundings, we might use this function $g = x \mapsto a, y \mapsto a, z \mapsto a$.

Now we want to use this result and match another term y with the constant b . The variable y is already mapped to a different constant, but we cannot say whether this is due to a previous matching process or simply because we needed to define a value for every input.

One solution might be to use a special constant symbol that we map variables to that are really unmapped for now. Instead, we want to use substitutions that were already introduced in [5]. A substitution is a partial mapping from variables to constants. We implement this by mapping to an Option of constant.

```
def substitution ( $\tau$ : signature):=  $\tau$ .vars  $\rightarrow$  Option ( $\tau$ .constants)
```

This allows us to only specify what is necessary. We call the substitution that maps any variable to none the empty substitution.

If we apply a substitution to a term and this term is a constant, we do not change the term. If the term is a variable and the substitution does not map the variable to none we replace the variable with the result of the substitution.

```
def applySubstitutionTerm (s: substitution  $\tau$ ) (t: term  $\tau$ ): term  $\tau$ 
  :=
match t with
| term.constant c => term.constant c
| term.variableDL v =>
  if p: Option.isSome (s v)
  then term.constant (Option.get (s v) p)
  else term.variableDL v
```

Similar to groundings we can extend this to atoms (`applySubstitutionAtom`) and rules (`applySubstitutionRule`). As we no longer replace all variables with constants, this only results in atoms instead of ground atoms.

The main result we want to prove is the following.

```
theorem groundingSubstitutionEquivalence
  (r: groundRule  $\tau$ ) (r': rule  $\tau$ ):
  ( $\exists$  (g: grounding  $\tau$ ), ruleGrounding r' g = r)  $\leftrightarrow$ 
  ( $\exists$  (s: substitution  $\tau$ ), applySubstitutionRule s r' = r)
```

This allows us to replace the grounding check with a substitution check when trying to validate trees and by this, we can bypass the problems that were illustrated in the example above.

For the forward direction, we want to find an equivalent substitution for every grounding. We can do this by mapping every variable to the same value of the grounding and place this in the some constructor to obtain an element of the Option type.

```
def groundingToSubstitution (g: grounding  $\tau$ ): substitution  $\tau$ 
  := fun x => Option.some (g x)
```

We have to prove that these are equivalent. We defined in the previous section a coercion from ground atoms to atoms. We can also coerce constants to terms by the `term.constant` constructor which enables us to use an equality in the following lemma.

Lemma 8 (`groundingToSubstitutionEquivTerm`). *Let t be a term and g be a grounding. Then the grounding of t by g is equal to the application of `groundingToSubstitution g` to t .*

Proof. If t is a constant, then both elements will simply return t .

If t is a variable v , then the term grounding by g will return $g(v)$. By the definition of `groundingToSubstitution` all variables v' return a some type which contains $g(v')$. Therefore the application replaces v by $g(v)$ and both results are equal. \square

Using this result we can extend this equality to the atom level (`groundingToSubstitutionEquivAtom`), because in both cases only the terms change by the application of the functions of the previous lemma. Finally, we can extend this to the rule level (`groundingToSubstitutionEquivRule`).

For the back direction, we need an additional property illustrated by the following example.

Example 7. Consider the program $\mathcal{P} = \{P \leftarrow, Q \leftarrow P\}$ and the signature $C = \emptyset$, $V = \{?x, ?y, ?z\}$ and $P = \{p, q\}$

Any rule in \mathcal{P} is already a ground rule and there exists a substitution, the empty substitution that maps all variables to none, so that the rule is equal to itself as a ground rule.

There is however no grounding that can achieve this. We cannot define a grounding since we have no constant available but have variables that need to be mapped somewhere. Therefore the equivalence does not hold here.

For the equivalence to hold we always need at least one constant symbol. There exist at least two possibilities to achieve this in Lean. We could require that the type of constants is inhabited or that it is non-empty. We decided to use the first as the inhabited class offers directly a constant, the default value, whereas the class non-empty only proves that such an element exists and requires the axiom of choice to extract it.

```
def substitutionToGrounding [ex: Inhabited  $\tau$ .constants]
(s: substitution  $\tau$ ): grounding  $\tau$  :=
fun x =>
if p:Option.isSome (s x)
then Option.get (s x) p
else ex.default
```

Example 7 (continued). If we add the fresh symbol a to C , we can use the function $f : V \rightarrow C, v \mapsto a$. This will not cause any problems during the verification of a proof tree. While it increases the number of groundings it will not result in any match if used, because it does not occur in the proof tree.

This is similar to the Herbrand base where we also add a constant symbol if no constant is present.

For the back direction, we prove again that it is equivalent on terms and then the atom and rule case easily follow.

Lemma 9 (`substitutionToGroundingEquivTerm`). Let τ be a signature that contains at least one constant symbol. For any substitution s , term t and constant c such that applying s to t yields t , also grounding t by `substitutionToGrounding` yields c .

Proof. If t is a constant, then it must be equal to c and neither the grounding nor the substitution change the term.

If t is a variable v , then $s(v)$ must be c in order to be equal to c . Therefore $s(v)$ is defined and `substitutionToGrounding` replaces v by the same value, so that the equality holds as well. \square

In the end, we obtain the desired theorem with the additional requirement on the set of constants.

Theorem 2 (groundingSubstitutionEquivalence). *Let τ be a signature that contains at least one constant symbol. Then for any τ rule r' and ground rule r we have that there exists a grounding g such that grounding r' by g yields r iff there exists a substitution s such that applying s to r' yields r .*

When introducing substitutions, we had the goal to only add what is needed to a substitution and usually, we want the smallest possible substitution. In order to formalize this, we want to define a partial order on substitutions, that is denoted by \subseteq

Firstly, we define the substitution domain of a substitution as the set of variables for which the substitution is defined.

```
def substitution_domain (s: substitution  $\tau$ ): Set ( $\tau$ .vars) :=
  {v | Option.isSome (s v) = true}
```

A substitution s_1 is then a subset of a substitution s_2 , if both substitutions agree on the substitution domain of s_1 . Outside of this s_1 is never defined, whereas s_2 might be, so we view s_1 as smaller.

```
def substitution_subs (s1 s2: substitution  $\tau$ ): Prop :=
   $\forall$  (v:  $\tau$ .vars), v  $\in$  substitution_domain s1  $\rightarrow$  s1 v = s2 v
```

This relation is by definition reflexive (`substitution_subs_refl`). It is also anti-symmetric (`substitution_subs_antisymm`). Since $s_1 \subseteq s_2$ anywhere where s_1 is defined s_2 is defined as well and has the same value. Since $s_2 \subseteq s_1$ anywhere where s_2 is defined the same holds, so that they have the same substitution domain and on it the same values. By the principle of function extensionality, both functions are equal.

Lastly, it is also transitive (`substitution_subs_trans`), so it is an instance of a partial order.

4.2 Unification

We know that instead of finding a grounding, it suffices to find a substitution. Now we want to describe an algorithm that tells us whether the ground rule that is formed from a vertex of the proof tree and its children is in the ground program. For this, we take inspiration from the unification problem of first-order logic.

In the unification problem of first-order logic we are given a set of equations $s_1 = t_1 \dots s_n = t_n$ between first-order terms and are required to present the most general unifier. A unifier μ is a function from variables to term so that if we replace every variable v in s_i and t_i by $\mu(v)$ then all equations are fulfilled.

A unifier μ is called the most general unifier if for every unifier σ there exists a function τ with $\sigma = \tau \circ \mu$.

Our problem is similar and is in general an instance of the following problem.

Definition 2. Let A and B be types and $f : A \rightarrow \text{substitution} \rightarrow B$ be a function. We call the following problem an instance of the unification problem.

Input: A substitution s and elements a, b of A and B .

Question: Does there exist a substitution s' with $s \subseteq s'$ and $f(a, s') = b$. If such an s' exists we call it a solution. We call s' a minimal solution if we have $s' \subseteq s^*$ for any solution s^* .

An example would be to consider A as the type of terms and B as the type of constants. Then f is `applySubstitutionTerm`. Another instance would be with atoms and ground atoms. a is in general the normal object and b the ground object.

An algorithm to solve the first-order unification problem is the algorithm of Martelli and Montanari [20] and is depicted below.

Algorithm 1 Algorithm of Martelli and Montanari

```

while There exists some equation for which a transformation is possible do
  Pick this equation  $e$  and do one of the following steps if applicable
  1. If  $e$  is of the form  $t = t$ , then delete this equation from the set.
  2. If  $e$  is of the form  $f(t_1, \dots, t_n) = f(s_1, \dots, s_n)$ , then delete  $e$  and add  $n$  new
     equations of the form  $t_i = s_i$ 
  3. If  $e$  is of the form  $f(t_1, \dots, t_n) = g(s_1, \dots, s_m)$  with  $g \neq f$ , then stop and
     reject.
  4. If  $e$  is of the form  $f(t_1, \dots, t_n) = x$  for a variable  $x$  and delete  $e$  and add
     an equation with the swapped order to the set
  5. If  $e$  is of the form  $x = t$  for some variable  $x$ , then check if  $x$  occurs in  $t$ .
     If it does, then stop and reject. If not map all  $x$  to  $t$  in the set.
end while

```

This algorithm offers a good starting point for our algorithms, but we know that certain transformations can't occur in the limited syntactic form we operate in. Additionally, we want to output a substitution instead of just answering whether a substitution exists. It is sufficient to do it here, but will later be important. Instead of mapping all x to t as done there in step 5, we will add $x \mapsto t$ to a substitution that is presented as an input. If a variable occurs on the left side, we will check whether it is already in the domain of the substitution and if so check if its current value is consistent with the right side. Steps 2 and 3 deal with function symbols that are not allowed in our language apart from constants, so that we can replace them by checking whether two constants are equal in the term case. For atoms or rules, we use something similar and split the problem into multiple smaller problems for terms or atoms.

Finally, as the one side of the equation is always a ground object there will never be a variable on this side, so we do not have to swap the equation as in step 4.

We will start by matching a term to a constant with the following algorithm.

```
def matchTerm (t: term  $\tau$ )(c:  $\tau$ .constants) (s: substitution  $\tau$ ):
Option (substitution  $\tau$ ) :=
match t with
| term.constant c' =>
    if c = c'
    then Option.some s
    else Option.none
| term.variableDL v =>
    if p:Option.isSome (s v)
    then
        if Option.get (s v) p = c
        then
            Option.some s
        else
            Option.none
    else
        extend s v c
```

We are given a term t , a constant c and a current substitution s and want to return the minimal solution.

This is done by case distinction. If t is a constant, then we either return s if t is equal to c or return none as two different constants can not be unified by a substitution. If t is variable, we check if t is in the domain of s . If it is already defined we check if the value matches the required value. If it is not defined we extend s with the new mapping $v \mapsto c$. Formally extend is defined in the following way:

```
def extend (s: substitution  $\tau$ ) (v:  $\tau$ .vars) (c:  $\tau$ .constants) :
substitution  $\tau$ 
:= fun x => if x = v then Option.some c else s x
```

We now formally prove the correctness of this algorithm.

Lemma 10 ([matchTermFindsSolution](#)). *Let t be a term, c be a constant and s be a substitution. If `matchTerm t c s` returns a substitution s' , then s' is a solution.*

Proof. The proof is done via case distinction. Suppose firstly that t is a constant c' . Since `matchTerm` returned a substitution we must have that c and c' are the same constant and therefore s' is s . Applying a substitution to a constant does not change it, so $s't = s'(c') = c' = c$. Additionally since \subseteq is a partial order and $s' = s$, we have that $s \subseteq s'$

Now we assume that t is a variable v . Now we do another case distinction on whether sv is defined or not. If it is defined, v must already be mapped

to c and we return s as this is a solution as seen previously. If it would be mapped to something else, then `matchTerm` would return `none`, which would violate our assumptions. If it is not defined, we use `extend`. After that v is mapped to c , so that $s't$ will be equal to c . Now we finally have to show that $s \subseteq \text{extend } s \ v \ c$. We only change the value of v . Since sv was not defined earlier, for any variable in the domain of s , s and `extend $s \ v \ c$` agree, so that it is fulfilled(`s_subset_extend_s`). \square

We have proven so far that `matchTerm` returns a solution, but it might not be a minimal solution. As we can transform any grounding into a substitution a match function might return the grounding in example 6. As we are interested in matching rules and ground rules, we will have to match many terms so that the minimality is an important property for later proofs.

Lemma 11 (`matchTermFindsMinimalSolution`). *Let t be a term, c be a constant and s be a substitution. If `matchTerm $t \ c \ s$` returns a substitution s' , then s' is a minimal solution.*

Proof. We already know that s' is a solution and only have to show its minimality. Let s^* be an arbitrary solution.

We prove the minimality again via case distinction on t . If t is constant, then s' must be equal to s . Since s^* is a solution we have that $s \subseteq s^*$ and since $s = s'$, we gain the desired result.

Now we consider the case of t being a variable v and do a case distinction whether sv is defined. If it was already defined, then s' must again be equal to s , so that the claim is fulfilled by the argument above. If sv was not defined, we have to show that `extend $s \ v \ c$` is a subset of any such s^* . We assume for a contradiction that this is not the case. Then there must be a variable in the domain of `extend $s \ v \ c$` such that `extend $s \ v \ c$` and s^* differ. Suppose this variable is v . Then s^* would either not be defined for v or map v to some other constant c' . In both cases $s^*v \neq c$, so that s^* would not be a solution and we would have reached a contradiction. If it is some other variable v' , then the value of `extend $s \ v \ c$` is simply the value of s . Since s^* maps v to a different value compared to s , s would not be a subset of s^* and we have reached another contradiction. \square

So we know that if `matchTerm` returns a substitution then it is a minimal solution. This holds however already if we never return a solution. We have to prove that if no substitution exists, then there is no solution.

Lemma 12 (`matchTermNoneImpNoSolution`). *Let t be a term, c be a constant and s be a substitution. If `matchTerm $t \ c \ s$` returns `none`, then there exists no solution.*

Proof. This is again done via case distinction on the type of t . If t is a constant c' , then c' must be different from c , so that `matchTerm` returns `none`. Then no substitution can map t to c .

If t is a variable v , then sv must be defined and mapped to a different value compared to c . Then again no such s' can exist. If s would be a subset of s' ,

then s' would not unify t with c and if s' would unify t with c then s would not be a subset of s' . \square

After proving the correctness for terms we now want to move up to atoms. Unfortunately, we cannot use recursion directly on the term list of an atom because an atom requires a proof that the length of the list is equal to the arity of the relation symbol, which fails when we do recursion on the list. Therefore we first establish a new procedure that matches a list of terms with a list of constants, if possible.

```
def matchTermList (s: substitution  $\tau$ ) (l1: List (term  $\tau$ )) (l2:
  List ( $\tau$ .constants)): Option (substitution  $\tau$ ) :=
  match l1 with
  | List.nil =>
    match l2 with
    | List.nil =>
      some s
    | List.cons _ _ =>
      none
  | List.cons hd tl =>
    match l2 with
    | List.nil => none
    | List.cons hd' tl' =>
      let s' := matchTerm hd hd' s
      if p: Option.isSome s'
      then matchTermList (Option.get s' p) tl tl'
      else none
```

Here we are given as previously a substitution as an input with the two lists. We see that if both lists differ in length then the function always returns none. If both lists are the same length we start matching the front and recursively call matchTermList with the resulting substitution until no substitution is found or the entire list is processed.

Lemma 13 ([matchTermListFindsSolution](#)). *Let s be a substitution, l_1 a list of terms and l_2 a list of constants. If $\text{matchTermList } s \ l_1 \ l_2$ returns a substitution s' , then s' is a solution.*

Proof. We prove this by induction on l_1 for arbitrary s and l_2 . In the base case, l_1 is the empty list. Since matchTermList returns something, l_2 must also be the empty list. Then s' is equal to s . Applying this to an empty list returns an empty list and by reflexivity, we have that $s' \subseteq s$.

In the induction step we have that l_1 is of the form $hd :: tl$ and we can similarly assume that l_2 is of the form $hd' :: tl'$ since matchTermList returns a substitution. Since matchTermList returned a substitution, matchTerm $hd \ hd'$ also must return a substitution s^* .

We then use this as an input to gain s' from matchTermList $s^* \ tl \ tl'$. By the induction hypothesis, we know that $s^* \subseteq s'$ and applying s' to tl results in

it being equal to tl' . First, we have to show that $s \subseteq s'$. From the correctness proof of `matchTerm`, we know that $s \subseteq s^*$ and from the induction hypothesis we know that $s^* \subseteq s'$. Since \subseteq is transitive, the result follows.

Secondly, we prove that s' is a solution for $hd :: tl$ and $hd' :: tl'$. From the induction hypothesis, we know that mapping tl with s' yields tl' . We know that s^* was a solution for hd and hd' and since $s^* \subseteq s'$, s' agrees on the domain of s^* with s^* . Therefore it is still a solution for hd and hd' . If hd was a constant, then any substitution would do. If hd was a variable, then it is mapped to the same element as by s^* so that applying s' to hd yields hd' as well, so that applying s' to $hd :: tl$ yields $hd' :: tl'$. \square

After proving that it is a solution, we prove that the solution is minimal.

Lemma 14 (`matchTermListFindsMinimalSolution`). *Let s be a substitution, l_1 a list of terms and l_2 a list of constants. If `matchTermList` s l_1 l_2 returns a substitution s' , then s' is a minimal solution.*

Proof. We prove this again via induction on l_1 for arbitrary l_2 and s . If l_1 is empty then l_2 is empty as well and we return s . The claim is true by reflexivity.

In the induction step we have that l_1 has the form $hd :: tl$ and we can assume that l_2 has the form $hd' :: tl'$ because a substitution was returned. Let s' be the result of `matchTermList` l_1 l_2 s . We have to show that for any solution s^* for l_1 , l_2 and s , we have that $s' \subseteq s^*$. Let s^* be a solution for l_1 , l_2 and s . Then it also must be a solution for hd , hd' and s and due to lemma 11 we have that `matchTerm` hd hd' $s \subseteq s^*$.

We additionally know that s^* is a solution for tl , tl' and s . Since `matchTerm` hd hd' $s \subseteq s^*$ we also know that it is a solution for tl , tl' and `matchTerm` hd hd' s . The substitution s' is the result of `matchTermList` tl tl' (`matchTerm` hd hd' s). The induction hypothesis tells us that s' is the minimal solution for these input values. Therefore we know that $s' \subseteq s^*$ and the claim is proven. \square

Finally, the negative case that confirms that the return value of none does indeed state that there is no solution.

Lemma 15 (`matchTermListNoneImplNoSolution`). *Let s be a substitution, l_1 a list of terms and l_2 a list of constants. If `matchTermList` s l_1 l_2 returns none, then there exists no solution s' .*

Proof. We prove this again via induction on l_1 for arbitrary l_2 and s . If l_1 is the empty list then l_2 must be non-empty and then no solution exists.

In the induction step we have that l_1 has the form $hd :: tl$. If l_1 is empty, the claim is proven, so we assume that l_2 has the form $hd' :: tl'$. There are two cases where `matchTermList` can return none. The first case is when `matchTerm` hd hd' s returns none. If that is the case there is no s' with $s \subseteq s'$ and applying s to hd will make it equal to hd' . Therefore the two lists cannot be equal either and the proof is finished.

If `matchTerm hd hd' s` returns a substitution s^* , then `matchTermList s* tl` must return `none`. From the induction hypothesis, we know that there is no substitution s' with $s^* \subseteq s'$ and applying s' to tl bringing it equal to tl' . Since s^* is already the minimal solution to match hd with hd' there cannot exist a solution whose application will lead to hd and hd' being equal and tl and tl' being equal. \square

This can be used to create a `matchAtom` procedure. We are given an atom and a ground atom and check if the symbols are equal. If they are, we use `matchTermList` to find a unifier. If the symbols are different, we will never unify them and can already return `none`.

```
def matchAtom (s: substitution  $\tau$ ) (a: atom  $\tau$ ) (ga: groundAtom  $\tau$ ):
Option (substitution  $\tau$ ) :=
  if a.symbol = ga.symbol
  then
    matchTermList s a.atom_terms ga.atom_terms
  else none
```

The correctness proofs follow from the proofs of `matchTermList`.

Now we can create a `matchAtomList` function similar to `matchTermList` and use this to define a `matchRule` function. We first try to match the head of the rule with the head of the ground rule. If this returns a solution, we use this as the start for matching the bodies. If not, we will not find a solution and return `none`.

```
def matchRule (r: rule  $\tau$ ) (gr: groundRule  $\tau$ ):
Option (substitution  $\tau$ ):=
  let s := matchAtom emptySubstitution r.head gr.head
  if p: Option.isSome s
  then matchAtomList (Option.get s p) r.body gr.body
  else none
```

We now prove again that if this returns a substitution then this is a solution and if none is returned then there is no solution using our previous results. We want a statement with the quantifier to combine it with the theorem about the equivalence between groundings and substitutions. We can prove this statement by using the result of `matchRule` as a witness for the quantifier. For the back direction, we see that this statement implies that a solution exists, which we always find with `matchRule`.

Theorem 3 (`matchRuleIsSomeIffSolution`). *Let r be a rule and gr be a ground rule. Then there exists a substitution s such that applying s to r yields gr iff `matchRule` returns some substitution.*

We now have a method to replace this quantifier with a computable function and can finally devise the check for tree validation.

4.3 Tree validation

We have so far introduced substitutions and have shown a way to decide whether a substitution exists that maps a rule to a ground rule. Now we want to finally show a way to validate trees. As the first step, we want to find whether a ground rule gr we gain from the tree is in the ground program, i.e. that there exists a grounding g (or by theorem 2 a substitution s) and a rule r from the program P so that applying g to r yields gr . We want to use the `matchRule` function defined earlier for this. We can simply pass each rule of the program given as a list of rules into this function and stop when `matchRule` returns a some object. This works but requires in the worst case to match every rule in the program with a given ground rule. Suppose that the predicate symbol of the head of the given ground rule is T . We can safely discard any rule whose head is not an atom with the predicate symbol T . This can be generalized into the symbol sequence which is a list of predicate symbols that starts with the predicate symbol of the head and then the predicate symbols of the body atoms in the order the atoms occur in the body.

```
def symbolSequence (r: rule  $\tau$ ): List  $\tau$ .predicateSymbols :=
  r.head.symbol :: (List.map atom.symbol r.body)
```

The equality of symbol sequences is a necessary condition for the existence of substitution that matches a rule with a ground rule, as the application of any substitution does not change the symbols and therefore not the symbol sequence. (`symbolSequenceNotEq`).

Now we could iterate over the whole program and calculate the symbol sequence for every rule and only start the `matchRule` process for those whose symbol sequences are equal to the given ground rule. In the worst case, we still have to iterate over the whole program. It would be even better to preprocess the program into a look-up structure that returns a list of rules for a given symbol sequence.

We use a function of the type `List τ .predicateSymbols \rightarrow List (rule τ)` as the look-up structure and build it iteratively by adding the first rule of the given program to the right symbol sequence element.

```
def parseProgramToSymbolSequenceMap (P: List (rule  $\tau$ ))
  (m: List  $\tau$ .relationSymbols  $\rightarrow$  List (rule  $\tau$ )):
  List  $\tau$ .relationSymbols  $\rightarrow$  List (rule  $\tau$ ) :=
  match P with
  | [] => m
  | hd::tl =>
    let seq:= symbolSequence hd
    parseProgramToSymbolSequenceMap tl
    (fun x => if x = seq then hd::(m x) else m x)
```

If we start this with a function that maps any symbol sequence to the empty list and gain a function m' then for any list of predicate symbols l , $m'(l)$ returns exactly those rules in P whose symbol sequence matches l . This follows from the following lemma.

Lemma 16 (`parseProgramToSymbolSequenceMap_mem`). *Let P be a program and m be a function that maps a list of predicate symbols to the list of rules. Let m' be the result of `parseProgramToSymbolSequenceMap P m`. Then for any list of predicate symbols l and rule r , we have r is in $m'(l)$ iff r was in $m(l)$ or r is in P and the symbol sequence of r equals l .*

Proof. We prove this by induction on the structure of P for arbitrary m .

If P is empty, then $m' = m$ and no rule is a member of P so that the claim follows.

If P has the structure $hd :: tl$, then m' is the result of

```
parseProgramToSymbolSequenceMap tl (fun x => if x = (
  symbolSequence hd) then hd::(m x) else m x)
```

Applying the induction hypothesis we see that for any rule r and list l , r is in $m'(l)$ iff it is in tl and has the symbol sequence l or was in `(fun x => if x = (symbolSequence hd) then hd::(m x) else m x)`

The only element of P not yet considered was hd and is either obtained by its symbol sequence or was already there in m so that the proof follows. \square

Since at the start $m(l)$ has no member for any l , the desired property holds. Now we can use this to check whether a substitution exists. `List.any` checks whether any element in $m(symbolSequence gr.toRule)$ matches the ground rule. If that is the case, we simply return unit as a matching rule in P was found. If that is not the case, we return an error message. The `Except` type allows us to combine these two return types.

```
def checkRuleMatch (m: List  $\tau$ .predicateSymbols  $\rightarrow$  List (rule  $\tau$ ))
  (gr: groundRule  $\tau$ ): Except String Unit :=
  if List.any (m (symbolSequence gr.toRule))
    (fun x => Option.isSome (matchRule x gr)) = true
  then Except.ok ()
  else Except.error ("No match for " ++ ToString.toString gr)
```

Lemma 17 (`checkRuleMatchOkIffExistsRuleForGroundRule`). *Let P be a program, m be the result of `parseProgramToSymbolSequenceMap P (fun _ => [])` and gr a ground rule. Then `checkRuleMatch m gr` returns `ok` iff there exists a rule r in P and a grounding g such that grounding r with g yields gr .*

Proof. Let l be the symbol sequence of gr . If `checkRuleMatch` returns `ok`, then there exists a rule r in $m(l)$ there exists a grounding g such that grounding r by g yields gr . By lemma 16 this rule r is a member of P so the claim is proven.

If `checkRuleMatch` returns an error, then we have to show that for rule $r \in P$ and grounding g grounding r by g yields a different ground rule compared to gr . If r has the same symbol sequence as gr , then it occurred in $m(l)$ and since `checkRuleMatch` returned an error `matchRule r gr` must return `none` so that the property holds.

If r has a different symbol sequence to gr then this property holds as well due to `symbolSequenceNotEq` and theorem 2. \square

This function is almost enough to complete the rule case. We just need to also verify all subtrees. For this we introduce the `List.map_except_unit` function that applies a function of the type $A \rightarrow \text{Except } B \text{ Unit}$ to a list l until the first error occurs. If no error occurs, then we return `Unit` which is a type that has only one element `()` and is the placeholder because we have to return something.

```
def List.map_except_unit {A B: Type} (l: List A)
(f: A → Except B Unit): Except B Unit :=
match l with
| [] => Except.ok ()
| hd::tl =>
  match f hd with
  | Except.ok () => List.map_except_unit tl f
  | Except.error b => Except.error b
```

We can prove by induction that this function returns unit iff f returns unit on all elements of the list l (`List.map_except_unitIsUnitIffAll`).

Now we have the necessary tools to check whether a given proof tree is valid which is done by the `treeValidator`. For a tree with the root a and subtrees l , we do a case distinction whether l is empty. If a is additionally an element of the database, then we accept a because we are in the database case. If not we use `checkRuleMatch` to check for the rule case. If this returns ok, then we can already accept a because there are no subtrees.

If not we can only be in the rule case and first use `checkRuleMatch` and if this returns ok, then we check all subtrees using the `treeValidator` again called via `List.map_except_unit`.

```
def treeValidator (m: List  $\tau$ .predicateSymbols → List (rule  $\tau$ )) (
d: database  $\tau$ ) (t: proofTree  $\tau$ ) : Except String Unit :=
match t with
| tree.node a l =>
  if l.isEmpty
  then if d.contains a
    then Except.ok ()
    else
      match checkRuleMatch m {head:= a, body := List.map
root l} with
      | Except.ok _ => Except.ok ()
      | Except.error msg => Except.error msg
  else
    match checkRuleMatch m {head:= a, body := List.map root l}
    with
    | Except.ok _ => List.map_except_unit l.attach (fun ⟨x, _h⟩
=> treeValidator m d x)
    | Except.error msg => Except.error msg
```

Using the previous lemmas and induction over the height of the tree for the recursive calls, we prove the correctness of the `treeValidator`.

Theorem 4 (`treeValidatorOkIffIsValid`). *Let P be a program, m be the result of `parseProgramToSymbolSequenceMap P (fun _ => [])` and d be a database. For any proof tree t , `treeValidator m d t` returns `ok` iff t is valid for P and d .*

If we want to validate the whole result of a datalog program, we will often need many proof trees and a function that can validate them. Additionally, we need to preprocess the program to gain the map from symbol sequences to rules. We can reuse the `List.map_except_unit` function for this purpose

```
def validateTreeList (P: List (rule  $\tau$ )) (d: database  $\tau$ ) (l: List
  (proofTree  $\tau$ )) : Except String Unit :=
let m := parseProgramToSymbolSequenceMap P (fun _ => [])
List.map_except_unit l (fun t => treeValidator m d t)
```

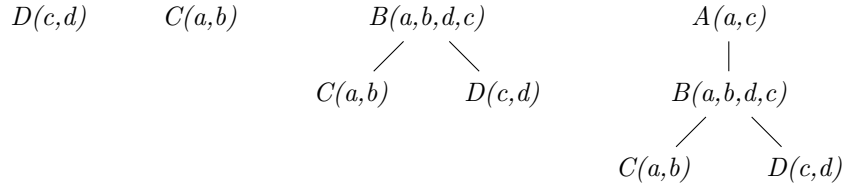
Using theorem 4 and `List.map_except_unitIsUnitIffAll`, we can show that this function returns `ok` iff all trees in l are valid. We know that a ground atom ga is in the proof-theoretic semantics if there exists a proof tree with the root ga so that we can conclude the set of roots of the trees in l is a subset of the semantics.

It turns out however that we can do a bit better and prove a stronger statement as the following example shows.

Example 8. *Consider the program P :*

$$\begin{aligned} A(?x, ?z) &\leftarrow B(?u, ?x, ?y, ?z). \\ B(?u, ?x, ?y, ?z) &\leftarrow C(?u, ?x), D(?z, ?y). \\ C(a, b). \\ D(c, d). \end{aligned}$$

The result of this program over an empty database is $\{A(a, c), B(a, b, d, c), C(a, b), D(c, d)\}$ and a proof tree is depicted below for each of them.



The proof tree for $A(b, d)$ already contains proofs for all the other elements of the program so it would be enough to check this single tree.

If we want to check a complete datalog result, we therefore do not have to include a proof tree for every element as they can occur in other proof trees already. We formalize this using the element member function that is true if this element is the root or a member in a subtree.


```

def elementMember (a: A) (t: tree A): Bool :=
  match t with
  | tree.node a' l => (a=a') ∨
    List.any l.attach (fun ⟨x, _h⟩ => elementMember a x)

```

A tree is valid if all subtrees are valid. In the rule case, we explicitly required this, whereas there are no subtrees in the database case. Any element member is the root of some subtree and therefore any element member of a valid proof tree is in the proof-theoretic semantics

(`allTreeElementsOfValidTreeInSemantics`).

This allows us to establish the alternative property of `validateTreeList`. We added the requirement of all proofs being valid to prove the back direction. As the validity of a proof tree is in general not preserved under permutation, it is not sufficient to just require every element member to be in the proof-theoretic semantics.

Lemma 18 (`validateTreeListUnitIffSubsetSemanticsAndAllElementsHaveValidTrees`).

Let P be a program, d be a database and l a list of proof trees. Then `validateTreeList` returns `ok` iff all element members of the trees in l are in the proof-theoretic semantics and all trees in l are valid.

Proof. For the forward direction, we already noted that if `validateTreeList` returns `ok`, then every tree in l must be valid. By the previous lemma therefore any element member of one of these trees is in the proof-theoretic semantics.

For the back-direction follows from the previous observation, that `validateTreeList` returns `ok` iff all trees in l are valid. \square

Now it is enough to pass just the proof tree for $A(a, c)$ from our example into the checker to validate the whole input and use in general fewer trees.

5 Validating graphs

In the previous section, we described an algorithm to verify proof trees. Proofs trees are a very natural way to encode certificates for datalog results but have some drawbacks.

Example 9. Consider the following program that computes the transitive closure in a different way.

$$\begin{aligned} T(?x, ?y) &\leftarrow E(?x, ?y). \\ T(?x, ?z) &\leftarrow E(?x, ?y), T(?y, ?z), T(?y, ?z). \end{aligned} \quad (4)$$

The database shall encode the elements of a chain $E(i, i + 1)$ for $i \in \{0, n\}$. In the proof tree representation we need the same proof tree twice for $T(?y, ?z)$ in the second rule and will need copies for it later, which results in large trees

Proof trees offer no reusability as they need to stay in the tree shape. Formally we can view the proof tree as a tree $T = (V_T; E_T)$ with a label function $label : V_T \rightarrow groundAtom$ because different vertices have the same label in a proof tree as in the example above.

A more compact representation would be a directed graph $G = (V_G, E_G)$ of T with $V_G = \{a \mid \exists v \in V_T, label(v) = a\}$ and $E_G = \{(a_1, a_2) \mid \exists v_1, v_2 \in V_T, label(v_i) = a_i \wedge (v_1, v_2) \in E_T\}$. Then a vertex can be considered equal to its label as no label occurs multiple times anymore.

Another advantage occurs when we want to validate multiple trees or even a complete result. Multiple trees will often share labels as well and combining them into a graph, i.e. the union of G_{T_i} for trees T_i will again include every label just once. In the previous section, we introduced the possibility of not having to check every proof tree as they occur in other proof trees as well. Selecting a minimal amount of proof trees to check a complete result is however still an instance of the NP-hard subset cover problem.

We require for this that if the successors of a ground atom are the same elements in every tree if the ground atom occurred in this tree. If not an atom would have more successors then previously in the tree and probably no matching ground rule. In practice, this requirement was always fulfilled.

Another requirement is acyclicity. We cannot use an atom to explain itself. Either we could have used it directly before or it cannot be explained by a proof tree at all.

5.1 Graph model

As a number of different results from mathematics are already formalized in mathlib, it is no surprise that graphs are already modelled in mathlib as depicted below:

```
structure SimpleGraph (V : Type u) where
  /-- The adjacency relation of a simple graph. -/
```

Show the exponential growth for clearer

Do a picture



Figure 7: An example of a pregraph

```

Adj : V → V → Prop
symm : Symmetric Adj
loopless : Irreflexive Adj

```

This graph representation is unfortunately not what we need in an algorithm. Firstly, the adjacency relation of this graph is always symmetric so that it does not encode the directed graphs we need. Secondly, the adjacency relation is a function to `Prop` and thus in general not computable but we want to use it in an algorithm. Finally, we need for the validation the successors of a vertex as they are supposed to represent the body of a ground rule. When we want to get them in this model we would have to ask the adjacency relation for every vertex if it would be computable.

Therefore we design our own implementation of a graph. We use an adjacency list approach so that we can quickly get all the successors of a vertex. This represented by a hash map whose keys are the vertices. Each vertex is mapped via the hash map towards its successors.

```

abbrev PreGraph (A: Type) [DecidableEq A] [Hashable A] :=
  Std.HashMap A (List A)

```

```

def vertices (g : PreGraph A) : List A := g.toList.map Prod.fst
def successors (g : PreGraph A) (a : A) : List A := g.findD a []

```

`Std.HashMap.toList` returns here a list of pairs of the type $(A, List A)$, i.e. a vertex and its successors. Using `Prod.fst` we obtain the first element of every pair in the list, which are all vertices. `Std.HashMap.findD a []` returns the value saved with a in the hash map which are the successors of a . If a vertex is not present (the value is not in the hash map), then `[]` is returned which is the fall back option provided to `Std.HashMap.findD`.

This approach contains all important elements in a graph, but there is a small technical problem with it.

Example 10. Consider the graph depicted in fig. 7. An element is colored, if it is in the list of vertices. The successors of a vertex are all elements that

We can view the type `Fin 5`, i.e. the natural numbers n with the property that $n < 5$. The vertices are the list $[1, 2, 3]$. The vertex 3 has the element 4 as a successor, but 4 is not a vertex of the graph. Such a behaviour is undesirable as searching for vertices that satisfy some criteria may lead outside of the graph.

We define a pregraph to be complete, if every successor of a vertex is a vertex itself and can then avoid the problem.

```

def complete (pg: PreGraph A) :=

```

```

 $\forall (a:A), pg.contains\ a \rightarrow \forall (a':A), a' \in (pg.successors\ a) \rightarrow pg$ 
 $.contains\ a'$ 

```

A graph is then a complete pre-graph and we can lift all previous operations to this new type.

```

abbrev Graph (A: Type) [DecidableEq A] [Hashable A] :=
  { pg : PreGraph A // pg.complete }

```

The completeness predicate was necessary, because our type is a superset than the vertices that occur in the graph. We do not expect every ground atom to be in the graph. An alternative would be to create a subtype of those ground atoms that occur in a list but this requires many types conversions when creating a graph from scratch as the list of vertices will change. Here we can add vertices to the graph and only have to prove that the completeness property still holds at the end.

We start by formalizing walks in a graph. Walks allow the repetition of vertices in itself in contrast to paths which suffices us.

A walk w in a graph G is a list of vertices that are all vertices of G and are connected via the successor relation of G , i.e. for every i with $0 < i < w.length$ we have that $w[i - 1] \in G.successors\ w[i]$. The starting vertex of the walk is consequently at the back.

```

def isWalk (l: List A) (G: Graph A): Prop :=
  ( $\forall (a:A), a \in l \rightarrow a \in G.vertices$ )  $\wedge$ 
  ( $\forall (i: \mathbb{N}), i > 0 \rightarrow \forall (g: i < l.length), l.get\ (Fin.mk\ i.\text{pred}$ 
    ( $\text{pred\_lt}\ i\ l.length\ g$ ))  $\in G.successors\ (l.get\ (Fin.mk\ i\ g))$ )

```

The `pred` function returns the predecessor of any natural number that is not zero and zero otherwise. Because we use the `List.get` function we not only have to specify a position but also a proof that this position is smaller than the length of the list to not get an error. These two elements are combined into a `Fin` type. In the future we usually omit these proofs in this work because of the length.

Due to the completeness predicate it would suffice to require that the starting vertex is in G but this variant is simpler to state. We also allow the empty walk `[]` as this will be beneficial in later applications.

Lists are beneficial in this use case in contrast to arrays because they can easily be extended at the front which we will use when explore a graph algorithmically. We can always extend a walk by a successor of the leading element.

Lemma 19 (`isWalk.extends_successors`). *Let w be a list of vertices and a and b be vertices. If $a :: w$ is a walk in G and b is a successor of a in G then $b :: a :: w$ is also a walk in G .*

A cycle is supposed to be a walk that starts and ends with the same node and an acyclic graph is a graph that has no cycles. As any list that only contains a single vertex of the graph would fulfill this property we require that a cycle has additionally at least length two.

```

def isCycle (l: List A) (G: Graph A): Prop :=
  if h: l.length < 2
  then False
  else

  isWalk l G ∧ l.get (Fin.mk 0 _) = l.get (Fin.mk l.length.pred _)
)

def isAcyclic (G: Graph A) := ∀ (l: List A), ¬ isCycle l G

```

5.2 Validation of a graph

After defining the graph model, we want to specify how the graph validation takes place. For these we defined the recursive predicate `isValid` and implemented it using `treeValidator`. As the graph we receive may not even be acyclic we cannot simply define a recursive predicate as this will then not terminate. Instead we define a local predicate for each vertex and its successor. Then we can check every vertex individually.

```

def locallyValid (P: program τ) (d: database τ) (v: groundAtom τ)
  (G: Graph (groundAtom τ)): Prop :=
  (∃(r: rule τ) (g:grounding τ), r ∈ P
    ∧ ruleGrounding r g = {head:= v, body:= (G.successors v) })
  ∨ ((G.successors v) = [] ∧ d.contains v)

```

Except for the recursive case this is very similar to `isValid`. Therefore we can reuse earlier concepts and build a checker for this. We expect the program again parsed into a look-up structure m , a database d , the vertex a and its successors passed as a list l .

```

def localValidityCheck (m: List τ.predicateSymbols → List (rule
  τ)) (d: database τ) (l: List (groundAtom τ)) (a: groundAtom τ
  ) : Except String Unit :=
  if l.isEmpty
  then
    if d.contains a
    then Except.ok ()
    else checkRuleMatch m (groundRule.mk a l)
  else
    checkRuleMatch m (groundRule.mk a l)

```

The correctness is proven similar as before assuming that m is the result of `parseProgramToSymbolSequenceMap` of the program P .

Lemma 20 (`localValidityCheckUnitIffLocallyValid`). *Let d be a database and P be a program. If m is the result of `parseProgramToSymbolSequenceMap` with a function that maps any element to the empty list and l is the list of successors of a ground atom a . Then the `localValidityCheck` returns ok iff a is locally valid with respect to P and d .*

In the next section we will show a method to show that a graph is acyclic and all elements are locally valid with respect to P and d . In the remainder of this section we assume that this is true and devise a method to show that then all vertices of the graph are in the proof-theoretic semantics of P and d .

Our goal is to create a proof-tree for every node. We devise a function that creates such a proof tree for a given node by recursively calling it on the successors.

```
def extractTree (a: A) (G: Graph A) (mem: a ∈ G.vertices) (
  acyclic: isAcyclic G): tree A :=
  tree.node a (List.map (fun ⟨x, _h⟩ => extractTree x G (G.
    complete a mem x _h) acyclic) (G.successors a).attach)
```

This function only terminates because the graph is acyclic. Therefore we require the fact as an explicit argument in the function above. We will argue with the vertices that are reachable from a vertex. These decrease the further we walk in the graph or else we would be able to reach a previous node and have a cycle which is not allowed. We start by formalizing the notion of reachability. We say that a vertex a **canReach** a vertex b if there exists a walk w from a to b . (The start of the walk is the last element in the list).

```
def canReach (a b: A) (G: Graph A):=
  ∃ (p: List A) (neq: p ≠ []),
    isWalk p G ∧ p.get (Fin.mk 0 _) = b
    ∧ p.get (Fin.mk p.length.pred _) = a
```

Any vertex a can reach it self via the walk $a(\text{canReach.refl})$.

We want to use the finite set of the vertices that are reachable from the current node so that we can argue that the cardinality decreases and by this the function terminates. This set will be a subset of the vertices of the graph G since any element reachable by a walk in G must be in the vertices of G . Ideally, we want to filter this finite set to only keep those vertices reachable from the current node. Unfortunately, `Finset.filter` requires a decidable predicate. This predicate is in principle decidable for example by Dijkstra's algorithm. Instead of implementing and validating it, we instead use classical logic to obtain that any predicate is decidable and reuse the previous filter version. This is now no longer computable but we only want to use it in the termination proof anyway.

```
noncomputable def Finset.mem_filter (p: A → Prop) (S: Finset A):=
  @Finset.filter A p (Classical.decPred p) S

lemma Finset.mem_filter_nc (a:A) (p: A → Prop) (S: Finset A): a ∈
  Finset.filter_nc p S ↔ p a ∧ a ∈ S
```

The global successors in a graph G of a vertex a are then all the vertices b that are reachable from a in G .

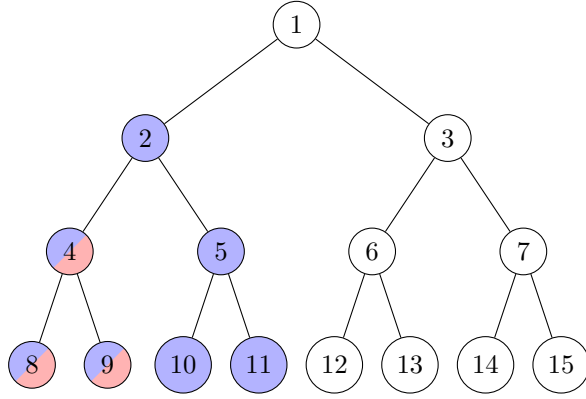


Figure 8: The global successors of 2 are colored blue, the global successors of 4 are colored red.

```
noncomputable def globalSuccessors (a:A) (G: Graph A): Finset A
  := Finset.filter_nc (fun b => canReach a b G) G.vertices.
  toFinset
```

We already see in fig. 8 that we have that $globalSuccessors(2) \subset globalSuccessors(4)$ and 4 is the successor of 2. We will generalize this. In any graph we have that for every vertex a and successor b of a we have that $globalSuccessors(b) \subseteq globalSuccessors(a)$, because any element c that b can reach can also be reached from a by first going to b and then following the walk from b to c

([globalSuccessorsSubsetWhenSuccessor](#)).

If the graph is additionally acyclic, we get that $globalSuccessors(b) \subset globalSuccessors(a)$, because a can reach a and b via the successor edge. If b could reach a , then this would lead to cycle which is not allowed.

([globalSuccessorsSSubsetWhenAcyclicAndSuccessor](#))

Since we call [extractTree](#) on all successors b of a and the global successors of b are a strict subset of the global successors of a the cardinality of the global successors of the vertex we call the function on always decreases.

```
def extractTree (a: A) (G: Graph A) (mem: a ∈ G.vertices) (
  acyclic: isAcyclic G): tree A :=
  tree.node a (List.map (fun ⟨x, _h⟩ => extractTree x G (G.
    complete a mem x _h) acyclic) (G.successors a).attach)
termination_by Finset.card (globalSuccessors a G)
decreasing_by
  simp_wf
  apply Finset.card_lt_card
  apply globalSuccessorsSSubsetWhenAcyclicAndSuccessor
  apply acyclic
  apply _h
  apply mem
```

We see from the definition that the root of the tree returned by `extractTree` is always the vertex a used in the arguments. (`rootOfExtractTree`)

It remains to be shown this leads to a valid proof tree.

Lemma 21 (`extractTreeStepValidProofTreeIffAllLocallyValidAndAcyclic`).

Let G an acyclic graph where all vertices are locally valid with respect to a program P and database d . Then `extractTree` results in a valid proof tree for any vertex a of G .

Proof. We prove this by strong induction on the cardinality of the global successors of a .

Since a is a vertex of G , a is locally valid. There are two cases for this. If a is locally valid because it has no successors and is in the database, then the map operation creates no subtrees so that the resulting proof tree is valid as well.

If a is locally valid because a and its successors form a ground rule from the ground program of P , then we create the subtrees with `extractTree`. By the previous observation the roots of the subtrees equal the successors of a , so that we have again a ground rule from the ground program. By the induction hypothesis all these trees are valid as well, so that the proof tree for a is valid. \square

Using this result the vertices of an acyclic graph whose vertices are all locally valid with respect to P and d are a subset of the proof-theoretic semantics of P and d .

5.3 Depth-first search

By the results of the previous result we know that in order to verify a graph we have to call the `localValidityCheck` on every vertex and its successors and determine whether the graph is acyclic. A typical solution to answer the second question is depth-first search, which we will implement and verify in this section. During depth-first search the algorithm has to visit every vertex. With the aim of reducing the iterations of the vertex list, we combine this with executing `localValidityCheck` on every vertex and its successors.

We generalize this by considering a function `f: A → List A → Except String Unit` for a `Graph A`. The `except` type is again chosen to return an error message to the user in the negative case.

We follow [12] when implementing depth-first search and use several helper functions that simplify the proofs. A first function `dfs` starts the process on every vertex and the actual exploration of the graph is done in `dfs_step`. So that we do not explore vertices multiple times, we store already explored vertices in a finite set. For performance reasons this is a hash set instead of the finite set from `mathlib`.

The function `isOkOrMessage` takes an arbitrary exception whose error type is a string and transform it into an `Except String Unit` object to have a similar output type as the `treeValidator`. This returns `ok` iff the original exception was also an `ok` element(`isOkOrMessageOkIffExceptionIsOk`).

We use `foldl_except_set` as a variant as the common *foldl* or *reduce* function. It stops and returns the first found exception. If no exception is found it execute the function on the first element with the input then and then calls itself recursively with the resulting set. We use this to use the set of already explored vertices in the further explorations using depth first-search.

```
def isOkOrMessage (e: Except String A): Except String Unit :=
  match e with
  | Except.error msg => Except.error msg
  | Except.ok _ => Except.ok ()

def foldl_except_set
(f: A → HashSet B → (Except String (HashSet B)))
(l: List A) (init: HashSet B): Except String (HashSet B) :=
  match l with
  | [] => Except.ok init
  | hd::tl =>
    match f hd init with
    | Except.error msg => Except.error msg
    | Except.ok S => foldl_except_set f tl S

def dfs (G: Graph A) (f: A → List A → Except String Unit) :
  Except String Unit :=
    isOkOrMessage (foldl_except_set (fun ⟨x,_h⟩ S => dfs_step x G
    f [] (isWalkSingleton G x _h) (List.not_mem_nil x) S) G.
    vertices.attach HashSet.empty )
```

The main work is done in `dfs_step` which we define next. This function takes a variety of arguments:

1. an element a that is the vertex we want to start exploring from,
2. a graph G ,
3. a function $f: A \rightarrow List A \rightarrow Except String Unit$ that shall be evaluated on every vertex and its successors,
4. a list *currWalk* of nodes that are the walk we used to arrive at a when exploring the graph,
5. a proof that $a :: currWalk$ are indeed a walk in G ,
6. a second proof that a is not in *currWalk* for the termination proof of this function and
7. a hash set *visited* of already explored vertices to allow earlier termination.

We start the algorithm by checking if a is already in *visited*. If that is the case, then we have no further exploration to do and return *visited*. If not we check if f raises an error on a and its successors and return it. If f returns

ok, we check for a cycle by intersecting a 's successors with the current walk. If any successor of a occurs in the current walk, we have found a cycle and return this as an error. If we find no cycle, then we recursively call `dfs_step` using `foldl_except_set` to update the set of visited vertices on every successor. If no error is found during this exploration, we add a using `addElementIfOk` to the return set and return this.

```
def addElementIfOk [Hashable A] (e: Except B (HashSet A)) (a:A):
  Except B (HashSet A) :=
  match e with
  | Except.ok S => Except.ok (S.insert a)
  | Except.error msg => Except.error msg

def dfs_step [Hashable A] (a: A) (G: Graph A)
(f: A → List A → Except String Unit) (currWalk: List A)
(walk: isWalk (a::currWalk) G) (not_mem: ¬ (a ∈ currWalk))
(visited: HashSet A) : Except String (HashSet A) :=
  if visited.contains a
  then Except.ok visited
  else
    match f a (G.successors a) with
    | Except.error msg => Except.error msg
    | Except.ok _ =>
      if succ_walk: (G.successors a) ∩ (a::currWalk) = []
      then
        addElementIfOk (foldl_except_set (fun ⟨x, _h⟩ S =>
          dfs_step x G f (a::currWalk)
          (isWalk_extends_successors walk x _h)
          (not_mem_of_empty_intersection succ_walk x _h) S)
          (G.successors a).attach visited) a
      else
        Except.error "Cycle detected"
```

This terminates because we only call `dfs_step` only on elements that do not occur on the current walk. Formally, we show that the cardinality of the finite set $(\text{List.toFinset } G.\text{vertices} \setminus \text{List.toFinset } \text{currWalk})$ decreases. In the recursive call we use $a :: \text{currWalk}$ as the current walk. Since any element of a walk is a vertices the cardinality of $\text{List.toFinset } \text{currWalk}$ is smaller then the cardinality of the set of vertices. In the recursive calls, the walk is instead $a :: \text{currWalk}$ and since a is not in currWalk the cardinality of $\text{List.toFinset } a::\text{currWalk}$ is larger than the cardinality of $\text{List.toFinset } \text{currWalk}$ and hence the overall cardinality decreases.

The desired theorem describing the behavior of depth-first search is the following:

Theorem 5 (*dfs_semantics*). *For any graph G and function $f: A \rightarrow \text{List } A \rightarrow \text{Except String Unit}$, $\text{dfs } G \ f$ returns ok iff the graph is acyclic and f*

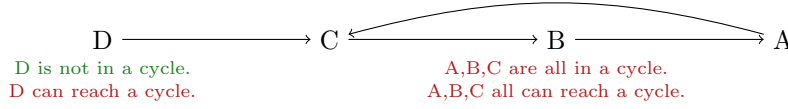


Figure 9: Propagating Acyclicity Check Results via Depth-First Search in example 11.

is evaluated to *ok* for every vertex a of G and its successors.

We defined acyclicity by stating that no list of elements of type A is a cycle in G but we do not check lists. Therefore we need another criteria for acyclicity that works on the vertex level. A first try is the membership in a cycle. A graph is obviously acyclic if no vertex is a member in a cycle. This is however not sufficient as the next example shows.

Example 11. Suppose that we start `dfs_step` on the vertex D on the graph depicted in fig. 9. We will start exploring D , then C , B and A . During the exploration of A we notice that its successor C already occurs in the current walk. Hence, we found a cycle. If we use membership in a cycle, then this information is not propagated back to D . `dfs_step` returned an error for D while D is not a member in a cycle. Instead we desire an iff relation between our criteria and the output of `dfs_step`. An alternative is the ability to reach a cycle. This is propagated back and a criteria for when `dfs_step` returns *ok*, which we will formalize now.

We reuse the `canReach` predicate for the new `reachesCycle` predicate. A vertex a reaches a cycle, if there exists a cycle c with a member b that a reaches.

```
def reachesCycle (a:A) (G: Graph A) :=
  ∃ (c: List A), isCycle c G ∧ ∃ (b: A), b ∈ c ∧ canReach a b G
```

We see from the definitions that a graph G is acyclic iff every vertex in G does not reach a cycle.

Lemma 22 (`acyclicIffAllNotReachCycle`). A graph G is acyclic iff all vertices of G do not reach a cycle.

Proof. If G is acyclic, then showing that all vertices of G are not reached from a cycle is equivalent to showing that any cycle in G cannot reach any element. Due to the acyclicity we know that no cycles exist in G so that the first direction is shown.

The back direction is proved via contradiction. Assuming that the graph is not acyclic, we know that there must exist a cycle c in G . Cycles in G are nonempty lists of vertices that are all in G . As any vertex can reach itself, there are vertices that are reached from a cycle in contrast to our assumption, so that we have reached the contradiction. \square

This property is propagated back from the successors.

@I originally created a worse looking diagram of the same example and Lukas created this better version for the paper. Can I use it here again ?

Lemma 23 ([NotreachesCycleIffSuccessorsNotReachCycle](#)). *A vertex a does not reach a cycle iff every successor of a does not reach a cycle.*

Proof. Both directions are proven via the contraposition. For the first direction, we have that there is a successor b of a that reaches a cycle. Then we can simply extend the walk by adding a at the back and then a can reach a cycle.

For the backdirection we assume that a reaches a cycle and try to show then one of its successors must also be reach a cycle. If a reaches a cycle c with an element b we consider two cases. If b would be a successor of a , we have shown our goal.

Now we assume that b is not a successor of a . Again we can consider two cases. If the walk is of length one then a and b must be equal and a is a member in a cycle itself. As long as a is not the first element in the cycle, we can simply pick the preceeding element in the cycle due to the connectness property of the walk. This does not work if a is the first element, but since it is a cycle a must also be the last element and we can pick the predecessor of the last element, which is a successor of a and in a cycle.

If a is reaches b with walk longer than length 1, then the walk also must have at least three elements since b is not a successor of a . Therefore the walk which must have the form $b \dots p a$ for a successor p of a . Then also p reaches a cycle. \square

As we also evaluate the function f during the depth-first search, we need a similar criteria that propagates the results of f back as we might encounter later a vertex where f is not evaluated to ok even if the starting vertex is evaluated to true similar to example 11.

We reuse the reachability predicate again to state that all vertices reachable from a vertice fulfill f .

Lemma 24 ([allTrueIfAllCanReachTrue](#)). *Consider a function $f: A \rightarrow List A \rightarrow Except String Unit$ and a graph G . Any vertex a in G is evaluated with its successors on f to ok iff any vertex b reachable in G from a vertex a in G is evaluated with its successors to ok.*

Proof. The forward direction holds because any vertex reachable in G must be a vertex of G itself. The backward direction holds because any vertex can reach themselves. \square

The propagation can be similarly stated but requires the vertex themselves to also be evaluated to ok. The proof follows from the recursive view on reachability.

Lemma 25 ([canReachLemma](#)). *Consider a function $f: A \rightarrow List A \rightarrow Except String Unit$ and a graph G . Any vertex reachable from a vertex a in G is evaluated with its successors on f to ok iff a is evaluated with its succesor on f to ok and any vertex reachable from a successor of a is evaluated to ok with its succesor on b .*

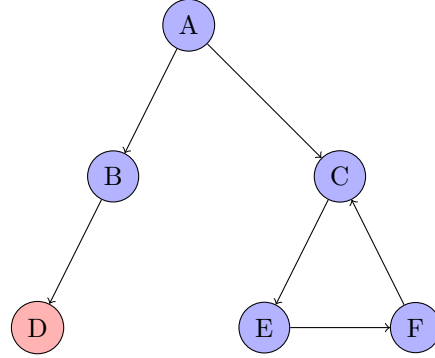


Figure 10: A graph for depth-first search. The result of f on a vertex and its successors is denoted by the color. Red denotes an error and blue denotes ok.

After establishing these criteria, we want to return to proving the correctness of the depth-first search algorithm. We start by proving the correctness of `dfs_step`. This function has a lot of arguments which we will call just as \vec{v} and if needed to an argument by its name.

Ideally, we would want to prove the following statement: Let a be the vertex in \vec{v} and G be the graph. Then `dfs_step` \vec{v} returns ok iff a does not reach a cycle in G and any vertex a reaches in G is evaluated with its successors to ok. This statement is not true because we can use the visited set to trick the algorithm.

Example 12. Suppose we start `dfs_step` on the vertex A in the graph depicted in fig. 10. Additionally, suppose that the set of visited vertices is $\{B, C, D, E, F\}$.

As A is not yet visited, we start the exploration process by evaluating f on A and its successors and as seen by the color this evaluates to ok. We have not seen any of the vertices before and therefore call `dfs_step` on B and C and it returns ok, since B and C are already in the set of visited vertices. Therefore the function in total returns ok for A .

This is however not what we expect it to do. A can reach the vertex D on which f evaluates to an error and it can reach the cycle $[C, E, F, C]$. These wrong evaluations occurred, because we assumed in the implementation that the visited set reports this property truthfully. Therefore we require this in every proof. If we would run this with an empty set of already visited vertices, then we gain the expected result.

Definition 3. Let a be a vertex in a graph G and $f: A \rightarrow \text{List } A \rightarrow \text{Except String Unit}$ be a function. We say that a has the `DfsStepSemantics` property if a does not reach a cycle in G and every vertex reachable from a in G evaluates to ok on f with its successors.

This statement allows us to state the theorem we want to prove for `dfs_step` more succinctly.

Theorem 6 (dfs_step_semantics). *Let \vec{v} be the input arguments with the vertex a and visited set V . If every element in V has the DfsStepSemantics property, then `dfs_step` \vec{v} returns ok iff a has the DfsStepSemantics property.*

We will prove this statement by induction on the amount of vertices not covered by the current walk, i.e. `(List.toFinset G.vertices \ List.toFinset currWalk)`, for arbitrary walks and sets of visited vertices. We note that the base case is always true, because if this is equal to zero, then every vertex of G is in the current walk. In \vec{v} we have however explicit proofs that a is a vertex in G and that a is not in the current path, so that we have a contradiction. The interesting case is therefore only the induction step. There we need a criteria for when `foldl_except_set` returns an error.

In general, this seems hard to do. Here we already see that the set of visited vertices plays no really important role in the result of the statement as long as all elements in it have the DfsStepSemantics property. As long as we preserve this property in creating the sets we can view `foldl_except_set` as individual calls when considering whether it returns an error or not.

Think of an example

Lemma 26 (foldl_except_set_is_ok). *Let l be a list of type A and S be a hash set of type B . Consider a property $p : B \rightarrow \text{Prop}$ and a function $f : A \rightarrow \text{HashSet } B \rightarrow (\text{Except String (HashSet B)})$, that fulfills the following two properties. Firstly, that f preserves p , i.e. if every element in S has the property p and f returns for the input a and S a set S' , then also every element in S' has the property p .*

Secondly, that f 's acceptance status does not change on sets whose elements fulfill the property p , i.e. for any hash sets S_1, S_2 such that all elements in each hash set fulfill the property p , we have that for any element a $f a S_1$ returns ok iff $f a S_2$ returns ok.

Then there exists a hashset S' that `foldl_except_set` $f l S$ returns S' iff $f a' S$ returns ok for any element a' in l .

Proof. We prove this by induction on the structure of l for arbitrary S .

If l is empty `foldl_except_set` always returns ok and there are no elements in l so that the claim holds.

If l has the shape $hd :: tl$, then `fhdinit` must return a set S . If not then `foldl_except_set` returns an error and there is an element a from $l = hd :: tl$ for which `fainit` is false so that both sides are false which finishes the prove.

The remaining result of `foldl_except_set` is therefore `foldl_except_set f tl S`. From the induction hypothesis we know that then any element a from tl we have that `faS` returns ok. We already have the required result for hd , it remains to get that always `f a init` returns ok. Since f preserves p and any element in `init` had the property p , also any element in S has the property p . Therefore we can use the second requirement for f to conclude that `f a init` returns true for any element a from $hd :: tl$. \square

The property in our case is the DfsStepSemantics property. The first requirement will follow from the induction hypothesis. Therefore it remains to show that `dfs_step` preserves the DfsStepSemantics property.

Lemma 27. *Let the*

5.4 HashSets for depth-first search

In the previous section, we used hash sets in the implementation of depth-first search. This offered a massive performance improve in practice in contrast to finite sets based on lists we used earlier. Unfortunately, while many lemmas for finite sets exist, there are currently almost no lemmas for hash sets. We try to bridge this gap partially by proving the correctness of insertion for hash sets.

```
lemma HashSet.contains_insert {S: HashSet A} (a:A ):  
∀ (a':A), (S.insert a).contains a' ↔ S.contains a' ∨ a = a'
```

6 Completeness

In the previous chapter we described a method why an atom is in the datalog semantics. This criteria is however not sufficient to recognize a solution. The empty list passes this test for any program while not being the semantics in most cases. Proof trees were a method to recognize why a ground atom is part of the semantics, but we are not aware of any simple way to describe why a ground atom is not in the semantics. Instead we want to show that the set of elements in the proof trees, S are complete in the sense that nothing else can be derived from it any more. This is the case when the T_P operator has a fixed-point for S or alternatively when S is a model. In this chapter we are going to create a certified model checker to show the completeness. If S passed the tree validation algorithm and is a model the following statements hold:

$$S \subseteq \text{proofTheoreticSemantics } P \text{ d} = \text{modelTheoreticSemantics } P \text{ d} \subseteq S$$

6.1 Partial ground rules

We defined the model property on the ground program $\text{ground}(P)$. In order to check if a set of ground atoms is a model for a program we therefore have to ground the program. We want to avoid simply grounding all the rules at once and instead do it in a more intelligent way because the amount of grounding is very large or even infinite. For this we introduce a new data structure, the partial ground rule. This bears some similarities to the rules we defined in chapter 3. It has a head that is an atom. The body is split into two lists. The first list contains the ground atoms and represents the atoms in the rule we already grounded, whereas the second list consists of the so far ungrounded atoms in the rule body. We want to move the ungrounded atoms one by one into the grounded list by applying substitutions, which map all variables of this atom to constants, so that we can transform this atom into a ground atom.

```
structure partialGroundRule ( $\tau$ : signature) where
  head: atom  $\tau$ 
  groundedBody: List (groundAtom  $\tau$ )
  ungroundedBody: List (atom  $\tau$ )
```

Example 13. A rule $r := q(X) : \neg r(a, b), t(X, c), s(c, d), u(d, X)$. may be viewed as the following partial ground rule $\text{pgr}_1 =$

```
{
  head := q(X),
  groundedBody := [],
  ungroundedBody := [r(a, b), t(X, c), s(c, d), u(d, X)]
}
```

This representation does not look any different to the rule itself as we do not use the grounded body at all. We can however move ground atoms from the

ungrounded body into the grounded body. The order of the atoms in the body does not matter semantically as we use a set definition when defining the criteria for a rule being true, so that we can simply move all ground atoms in the grounded body.

```
{
  head := q(X),
  groundedBody := [r(a, b), s(c, d)],
  ungroundedBody := [t(X, c), u(d, X)]
}
```

We can transform any rule into a partial ground rule by setting the head as the head, the body as the ungrounded body and setting the grounded body to be empty.

```
def partialGroundRuleFromRule (r: rule  $\tau$ ): partialGroundRule  $\tau$  :=
{
  head := r.head,
  groundedBody := [],
  ungroundedBody := r.body
}
```

Example 14. pgr_1 is exactly the result of *partialGroundRuleFromRule* r .

We choose this representation instead of the approach used for pgr_2 as this does not require iterating over the whole body to find ground atoms. As we will apply multiple substitutions in the grounding process, we will create anyways ground atoms in different place to the atom in the body we are currently trying to ground.

Any partial ground rule can also be transformed back into a rule by concatenating the grounded and ungrounded body and combining creating first a partial ground rule from a rule and then transforming the resulting partial ground rule yields the original rule. This does not hold if we swap the operations as we do not explicitly move atoms without variables from the start of the body into the grounded body.

```
def partialGroundRule.toRule (pgr: partialGroundRule  $\tau$ )
: rule  $\tau$  :=

{
  head := pgr.head,
  body := (List.map (groundAtom.toAtom) pgr.groundedBody)
  ++ pgr.ungroundedBody
}

lemma partialGroundRuleToRuleInverseToFromRule (r: rule  $\tau$ ):
  r = partialGroundRule.toRule (partialGroundRuleFromRule r)
```

Example 15. The application of `partialGroundRule.toRule` on `pgr1` yields `r` as predicted by the lemma.

If we swap both functions and first apply `partialGroundRule.toRule` to a partial ground rule and then convert the resulting rule back to a partial ground rule, we see that this is no longer equal. Applying `partialGroundRule.toRule` to `pgr2` results in the rule $q(X) : \neg r(a, b), s(c, d), t(X, c), u(d, X)$. Converting this back into a partial ground rule with `partialGroundRuleFromRule` we gain

```
{
  head := q(X),
  groundedBody := [],
  ungroundedBody := [r(a, b), s(c, d), t(X, c), u(d, X)]
}
```

which is different from `pgr2`

Using the transformation to rules we can lift definitions like safety or being true to partial ground rules.

```
def partialGroundRule.isSafe (pgr: partialGroundRule τ) :=
  pgr.toRule.isSafe

def partialGroundRule.ruleTrue (pgr: partialGroundRule τ)
(i: interpretation τ) :=
  ruleTrue pgr.toRule i
```

So far we only split the body into two parts and have the goal of applying substitutions to move everything into the grounded body. This is not too different to just applying groundings directly. This process allows us to potentially stop early. If the substitutions we applied so far resulted in a ground atom that is not part of the interpretation *i* we already know that the rule is true. No matter how the remaining variables are mapped, the body will never be a part of *i* and therefore the antecedent is false and the rule is therefore true. We call a partial ground rule *active in an interpretation i* where all ground atoms in the `groundedBody`

```
def active (pgr: partialGroundRule τ) (i: interpretation τ) :=
  ∀ (ga: groundAtom τ), ga ∈ pgr.groundedBody → ga ∈ i
```

6.2 Explore grounding

7 Evaluation

In the previous sections we proved the correctness the algorithms to check the soundness and completeness of datalog reasoning results. Now, we are interested in the practicability of these algorithms on actual data. We combined these algorithms into a command line tool that takes a file consisting of the problem and the certificates and tells us whether the result is correct according to the certificate.

7.1 Input format

The input format is JSON-based because Lean offers already direct support for JSON. Similarly, as we were able to derive decidable equality or inhabitedness, we can also derive functions that convert Lean objects to JSON objects or try to create a Lean object from a JSON object.

We can define mock terms similar to `term` but with the variables and constants as simple strings. Lean knows how to read and write strings into JSON hence we can derive the `Json` methods.

```
inductive mockTerm
| constant: String → mockTerm
| variable: String → mockTerm
deriving DecidableEq, Lean.FromJson, Lean.ToJson, Repr
```

Using this type we can similarly as to `atom` define `mockAtom`. In contrast to real atoms, we do not require a proof that the number of terms matches the arity of the predicate symbol as encoding such a proof is difficult and we have no information about the arity. The symbol is again just a string.

```
structure mockAtom where
  (symbol: String)
  (terms: List mockTerm)
deriving DecidableEq, Lean.FromJson, Lean.ToJson, Repr
```

Mock atoms form mock rules similarly as atoms form rules and a program is simply a list of mockRules. Lists are a basic feature of the `Json` decoder which allows us to get the from a json file.

Example 16. *The program*

$$P = \{ \begin{array}{l} T(?x, ?y) \leftarrow E(?x, ?y), Q(a). \\ \end{array} \} \quad (5)$$

is represented as following in json:

```
"program": [
{
```

```

    "head": {
      "symbol": "T",
      "terms": [
        {
          "variable": "?x"
        },
        {
          "variable": "?y"
        }
      ]
    },
    "body": [
      {
        "symbol": "E",
        "terms": [
          {
            "variable": "?x"
          },
          {
            "variable": "?y"
          }
        ]
      },
      {
        "symbol": "Q",
        "terms": [
          {
            "constant": "a"
          }
        ]
      }
    ]
  }
}

```

Afterwards we go twice through the program. In the first run, we collect all the predicate symbols and their arities ([parsingArityHelper](#)) and report an error if a predicate symbol is used in multiple atoms with different amounts of terms. Using such a list we can construct a signature ([parsingSignature](#)). We use for constants and variables simply the set of string as types and for predicate symbols the subset of strings that occurred as symbols. This choice allows us to directly inherit the requirements for the signature elements such as decidable equality or hashability.

In the second run, we then transform every mock object into the corresponding datalog object of the previously created signature.

The second part is the input file is either a list of trees or a graph. For

these we define again mock objects and transform them after we transformed the program.

```
inductive jsonTree (A: Type)
| node (label: A) (children: List (jsonTree A))
deriving Lean.FromJson, Lean.ToJson

-- graph validation
structure mockEdge where
  (vertex: mockAtom)
  (successors: List (mockAtom))
deriving DecidableEq, Lean.FromJson, Lean.ToJson

structure mockGraph where
  (edges: List mockEdge)
deriving Lean.FromJson, Lean.ToJson
```

Additionally, there are two command line options that can be set. Firstly, the option `-g` specifies that the input file as a graph instead of a list of trees which is the default option. Secondly, we tell the program with `-c` to also use the `modelChecker` to check for completeness.

7.2 Results

8 Conclusion

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