Event-Object Reasoning with Curated Knowledge Bases: Deriving Missing Information

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Abstract. The broader goal of our research is to formulate answers to why and how questions with respect to knowledge bases, such as AURA. One issue we face when reasoning with many available knowledge bases is that at times needed information is missing. Examples of this include partially missing information about next sub-event, first sub-event, last sub-event, result of an event, input to an event, destination of an event, and raw material involved in an event. In many cases one can recover part of the missing knowledge through reasoning. In this paper we give a formal definition about how such missing information can be recovered and then give an ASP implementation of it. We then discuss the implication of this with respect to answering why and how questions.

1 Introduction

Our work in this paper is part of two related long terms goals: answering “How”, “Why” and “What-if” questions and reasoning with the growing body of available knowledge bases [1], some of which are crowd-sourced. Although answering “How” and “Why” questions are important, so far little research has been done on them. Our starting point to address them has been to formulate answers to such questions with respect to abstract knowledge structures obtained from knowledge bases. In particular, in the recent past we considered Event Description Graphs (EDGs) [1] and Event-Object Description Graphs (EODGs) [2] to formulate answers to some “How” and “Why” questions with respect to the Biology knowledge base AURA [3].

Going from the abstract structures to reasoning with real knowledge bases (KBs) we noticed that the KBs often have missing pieces of information, such as properties of an instance (of a class) or relations between two instances. For example, AURA does not encode that Eukaryotic translation is the next event of Synthesis of RNA in eukaryote; this may be because the two subevents of “Protein synthesis” were encoded independently. The missing pieces make the KB and the Description Graphs constructed from it fragmented and as a result answers obtained with respect to them are not intuitive. Moreover, the KBs like AURA often have two or more names that refer to the same entity. To get intuitive answers they need to be resolved and merged into a single entity.

1 See for example, http://linkeddata.org/
Such finding of non-identical duplicates in the KB and merging them into one is referred in the literature as entity resolution [4,5].

In this paper, we start with introducing knowledge description graphs (KDGs) as structures that can be (without much reasoning) obtained from frame based KBs such as AURA. We discuss underspecified knowledge description graphs (UDGs) and formulate notions of reasoning with respect to these graphs to obtain certain missing information. We then present our approach of entity resolution and use it in recovering additional missing information. We give an Answer Set Programming (ASP) encoding of our formulation. We conclude with a discussion on the use of the above in answering “why” and “how” questions.

2 Background: Frame-based Knowledge Bases; ASP

The KB we used in this work is based on AURA [3] and was described in details in [6]. AURA is a frame-based KB manually curated by biology experts; it contains a large amount of frames describing biological entities events (or processes). One important aspect of our KB is the class hierarchy. For example, its basic class is Thing, which has two children classes: Entity and Event. Entity is the ancestor of all classes of biological entities; Event, of biological events. For instance, Spatial entity, Eukaryote, Nucleus and mRNA are descendants of Entity, while “Eukaryotic translation”, “Eukaryotic transcription” are descendant of Event.

Our KB is a set of facts of the form “has(A, slot_name, B)” where A and B are either classes or instances (of classes), slot_name is the name of the relation between A and B such as instance_of, raw_material or results. The statement “eukaryotic translation is based on mRNA” is represented in our KB as follows.

\[
\begin{align*}
&\text{has(euca\_transl4191, instance\_of, event)}. \\
&\text{has(euca\_transl4191, instance\_of, eukaryotic\_translation)}. \\
&\text{has(euca\_transl4191, base, mrna4642)}. \\
&\text{has(mrna4642, instance\_of, mrna)}. \\
\end{align*}
\]

This snippet reads as “eukaryotic_translation4191 is an instance of class event and an instance of class eukaryotic_translation. eukaryotic_translation4191 is based on mrna4642, which is an instance of mrna”.

For the declarative implementation of our formulations, we use ASP [7]. That allows us to use our earlier work [6] on using ASP to reason with frame-based knowledge bases. ASP’s strong theoretical foundation [8] and its default negation and recursion are useful in our encoding and in proving results about them.

3 Knowledge Description Graphs

An Underspecified Knowledge Description Graph (UDG) is a structure to represent the facts about instances and classes of events, entities and relationships between them. An UDG is constructed from knowledge bases such as AURA. Formal definition of the UDGs is given in the following.

**Definition 1.** An UDG is a directed graph with one type of node and five types of directed edges: compositional edges, ordering edges, class edges, locational edges and participant edges. Each node represents an instance (of a class) or a class in our KBs.

\(^2\) Our examples are either directly from AURA, or are slightly modified from it.
Table 1. Types of edges in an UDG. An edge of relation $Y$ from a node $X$ to $Z$ represents $X[Y] = Z$, meaning the slot $Y$ of the entity $X$ has value $Z$.

We used the slot names in KM [9] and AURA as a guide to categorize four types of edges (Table 1).

A **Knowledge Description Graph (KDG)** (a slight generalization of EODGs in [2]) is constructed from an UDG. A node in a KDG represents either an instance of a biological entity, an instance of a biological process, or a class of biological entity/event. The KDG structure allows us to answer “How” and “Why” questions. More formally, a KDG is defined as follows.

**Definition 2.** A KDG is a directed graph with: (i) three types of nodes: event nodes, entity nodes, and class nodes; and (ii) five types of directed edges: compositional edges, class edges, ordering edges, locational edges and participant edges. A KDG has the property that there are no directed cycles within any combination of compositional, locational and participant edges.

Figure 1 shows the types of edges in a KDG and the corresponding sources and destinations of the edges. Edges in a KDG are from the edges of the UDG, with additional type constraints of the source and destination nodes. For example, ordering edges must be from events to events; compositional edges are from events to events or from entity to entity, depending on their specific relations.

Since UDGs and KDGs can be huge, we usually work on their smaller subgraphs that are rooted at an entity or an event. They are defined as follows.

**Definition 3.** Let $Z$ be a node in a KDG $G$. The Knowledge Description Graph (KDG) rooted at $Z$ is the subgraph of $G$ composed of: (1) The set $N$ of all the nodes of $G$ that are accessible from $Z$ through compositional edges, class edges,
locational edges or participant edges; and (2) All the edges of $G$ connecting two nodes in $N$. We denote the KDG rooted at $Z$ as $KDG(Z)$ or the KDG of $Z$.

The UDG rooted at $Z$, denoted as $UDG(Z)$, is defined similarly. Figure 2 shows an example of a KDG rooted at Eukaryote where every other node can be reached from Eukaryote through edges with solid lines (compositional edges, class edges, locational edges or participant edges).

**Fig. 2.** A KDG rooted at the entity Eukaryote. Event, entity, and class nodes are respectively depicted by rectangles, circles and hexagons. Compositional edges are represented by solid-line arrows; ordering edges by dashed-line arrows; participant edges by lines with a black-square pointed to the entity node; class edges by diamond head arrows and locational edges by lines with a circle pointed to the event node.

4 Reasoning about Missing Info. in UDGs and KDGs

In this section, we discuss about missing information in the UDGs and the KDGs and how we can recover some of it through reasoning.

4.1 Event, Next Event, First Sub-event and Last Sub-event

One can directly obtain event names by looking at facts of the form “has(E, instance_of, event)” in the KB; and concluding E in it as an event. However, for some events such facts may be missing. In that case, we may be able to get the fact from the UDG’s edges and the edge constraints of the KDGs (Figure 1).

More formally:

**Definition 4.** Let $E$ be a node in the $UDG(Z)$. $E$ is an event if there is (i) a participant edge or an ordering edge from $E$; (ii) a locational edge or an ordering edge to $E$; (iii) a compositional edge (of subevent or first subevent relation) from/to $E$; or (iv) a path of class edges from $E$ to the class event.

Based on Definition 4, we can get that photosynthesis is an event because it has compositional edges (of subevent relation) to light reaction and calvin cycle.

Next-event, first subevent and last subevent are amongst the most important properties in describing events. However, they are not always directly available in our KB. Fortunately, in many cases, we can recover them from other properties.

**Definition 5.** Let $E$ and $E'$ be two events in the $KDG(Z)$. Event $E'$ is the next event of $E$ if $E$ enables, causes, prevents or inhibits $E'$.
In other words, \( E' \) is the next event of \( E \) if there is an ordering edge from \( E \) to \( E' \).

**Definition 6.** Let \( S \) be the set of subevents of an event \( X \) in the KDG(\( Z \)). Event \( E \) in \( S \) is the first subevent of \( X \) if there exists no other event \( E' \) in \( S \) such that \( E \) is the next event of \( E' \). Similarly, event \( E \) in \( S \) is the last subevent of \( X \) if there exists no other event \( E' \) in \( S \) such that \( E' \) is the next event of \( E \).

Here we assume that \( S \) was properly encoded in that there is only one chain of subevents in \( S \). In our KB, light reaction and Calvin cycle are two subevents of photosynthesis and light reaction enables Calvin cycle. But their orders are not defined. However, using Definition 5 and 6, we can identify that: Calvin cycle is the next event of light reaction; light reaction is the first subevent of photosynthesis; and Calvin cycle is the last subevent.

### 4.2 Input/Output of Events

**Two types of events:** In our KB there are two types of events: transport events and operational events. In a transport event, there is only a change in the locations; the input location and output location are different from each other while the input entity and output entity are the same. All other events are operational events. In an operational event, there is usually no change in its location. We differentiate two types of events by their ancestor classes; transport events are descendants of the classes move_through, move_into and move_out_of.

**Input, Output, Input Location, Output Location:** To reason about the KDG, we need the input and output of each event as well as the input location and the output location, which are not always available. In the following, we show how to use various event’s relations - such as raw-material, destination, location and others - to create four new relations (IO relations): input, output, input-location and output-location. After that, we propose rules to complete the KDG’s IO relations.

We created the IO relations of an event based on specific relations as shown in Table 2. The meaning of relation “base” from AURA depends on the context. For transport events, it is for input-location; for operational events, it is for input.

| Event type                  | IO relation type | Relation(s)       |
|-----------------------------|------------------|-------------------|
| Transport event             | input            | object            |
| Transport event             | output           | object            |
| Transport event             | input-location   | base, origin      |
| Transport event             | output-location  | destination       |
| Operational event           | input            | object, base, raw-material |
| Operational event           | output           | result            |
| Operational event           | input-location   | site              |
| Operational event           | output-location  | destination       |

**Table 2.** The IO properties of events and their corresponding relations.

**Completing Missing Information of Input, Output, Input Location, Output Location:** We can obtain missing IO properties of an event from its subevent(s). For instance, an input of the first subevent of \( E \) is also an input of \( E \).
**Definition 7.** Let $FSE$ and $LSE$ respectively be the first subevent and last subevent of event $E$ in $KDG(Z)$.

Let $InputRelation$ be the input relation, input-location relation or one of their corresponding relations Table 2. If $InputRelation$ is a relation from $FSE$ to $X$ then $InputRelation$ is also a relation from $E$ to $X$.

Let $OutputRelation$ be the output relation and output-location relation or one of their corresponding relations. If $OutputRelation$ is a relation from $LSE$ to $X$ then $OutputRelation$ is also a relation from $E$ to $X$.

In our KB, photosynthesis has two subevents: light reaction and calvin cycle, the next event of light reaction. Sunlight is the raw-material of the light reaction, sugar is the result of calvin cycle. Using Definition 7 we have that sunlight is the raw-material of photosynthesis and sugar is its result. Moreover, we also have: sunlight is the input of light reaction as well as photosynthesis; sugar is the output of both calvin cycle and photosynthesis.

Similarly, the output location of an operational event is often not defined in the KB but we can use input location as the default value for output location.

**Definition 8.** Let $E$ be an event in $KDG(Z)$, $E$’s input location is also the output location if $E$’s output location has not been specified.

Figure 3 shows the IO properties of events in Fig 2. The properties in bold are the ones that were recovered using Definitions 7 and 8.

**Fig. 3.** The IO properties of events in Fig 2. The five blocks contain IO properties of events: Synthesis of RNA in eukaryote, Eukaryotic translation, Move_out, and Eukaryotic transcription. The middle rectangle of each block contains the event name. The top rectangles are for input and output locations; the bottom rectangles are for input and output. The properties in bold were recovered using Definitions 7 and 8.

### 4.3 Main Class of an Instance

In our KB, one instance can belong to many classes. For example, dna_strand19497 - the input of Eukaryotic transcription - is an instance of dna_strand, dna_sequence, nucleic_acid and polymer.

However, to reason about the equality between in-

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*For the sake of simplicity, in the previous figures and descriptions, we usually referenced the entities and events by their “main” class(es) and not by the instances’ names although our KB and our implementation works on instances’ names.*
stances, we need the “main” class(es), the most specific class(es) of that instance. Our formal definition of “main” class is given below.

**Definition 9.** Let $E$ be an instance in $KG(Z)$. Class $B$ is a main class of instance $E$ if (1) it is a class of $E$ and (2) it is not the case that there is a Class $A$ which is a class of $E$ and (a) Class $B$ is ancestor of Class $A$ or (b) Class $B$ is a general class but Class $A$ is not; where general classes in our KB are thing, event, entity, spatial_entity, tangible_entity, and chemical_entity.

The main classes of $dna\_strand19497$, according to the Definition 9, are $dna\_strand$ and $dna\_sequence$; the other classes of $dna\_strand19497$ are ancestors of those two.

## 5 Entity Resolution

In the KBs such as AURA, the curation was done in many sessions and probably by many people. (Same is true with respect to many other KBs; especially the ones that are developed using crowd-sourcing.) The results are, in many cases, (i) two different instance names were used when they are probably the same instance; and (ii) parts of some biological process were encoded as independent events. For example: the input of *Eukaryotic translation* (Figure 2) is $mrna4642$ whereas the output of *Move_out* is $mrna22911$; *Synthesis of RNA in eukaryote* and *Eukaryotic translation* should be subevents of “Synthesis of protein in eukaryote” but they are encoded as two separate events.

In this section, we propose methods to solve the first problem. These methods are then used to solve the second problem in the next section. In order to compare two instances in a KB, we define a match relation. Generally speaking, instance $A$ can match with instance $B$ if $A$ can be safely used in a context where a term of $B$ is expected. We defined matching relation with many confidence levels for greater flexibility in future works.

**Definition 10.** Let $A$ and $B$ be two instances in $KG(Z)$. Let Class $A$ and Class $B$ be main classes of $A$ and $B$ respectively.

1. $A$ matches with $B$ with high confidence if one of the following is true
   (a) $A = B$ ($A$ and $B$ are the same instance)
   (b) $A$ is cloned from $B$ (Shortcut in AURA to specify that $A$ has all the properties of $B$)
   (c) Class $A$ is an ancestor of Class $B$.
2. $A$ matches with $B$ with medium confidence if $A$ and $B$ are both cloned from an instance $C$.
3. $A$ matches with $B$ with low confidence if Class $A = $ Class $B$ ($A$ and $B$ are instances of the same main class).
4. $A$ matches with $B$ with confidence $\min(\text{Conf}_1, \text{Conf}_2)$ if
   (a) $A$ matches with $C$ with confidence $\text{Conf}_1$ and
   (b) $C$ matches with $B$ with confidence $\text{Conf}_2$
5. Otherwise, $A$ does not match with $B$. 
Using Def.10, we can match mrna4642 - the input of Eukaryotic translation - with mrna22911 - the output of Move_out, because both have mrna as the main class.

While Def.10 can match all the input and output in our aforementioned example, it is not sufficient for matching location. For example, we can not match an instance of cytoplasm to an instance of cytosol. However when we say Event A occurs in cytosol, we can understand that Event A occurs in cytoplasm. To overcome this shortcoming, we define the relation Spatially match as follows.

**Definition 11.** Instance A in KDG(Z) is a location instance if the class ClassA of A is a descendant of the class spatial entity.

**Definition 12.** Let A and B be two location instances in KDG(Z). Let ClassA and ClassB be main classes of A and B respectively.

1. Location A spatially matches with location B with confidence $Conf$ if instance A matches with instance B with confidence $Conf$.
2. Location A spatially matches with location B with high confidence if one of the following is true:
   (a) $B$ is inside $A$ (the relation inside is encoded in our KB by slot name is_inside).
   (b) $B$ is part of $A$ (the relation “part of” is encoded in our KB by slot name part_of).
3. Location A spatially matches with location B with confidence $\min(Conf_1, Conf_2)$ if
   (a) A spatially matches with location C with confidence $Conf_1$, and
   (b) C spatially matches with B with confidence $Conf_2$.

Suppose that in our KB we have: cytosol234 and cytosol987 all are instances of cytosol; cytoplasm322 is an instance of cytoplasm and cytosol987 is inside cytoplasm322. We can then conclude: cytosol234 and cytosol987 match with each other with low confidence, according to Def.10.3; cytoplasm322 spatially matches with cytosol987 with high confidence (Def.12.2.a); cytosol987 spatially matches with cytoplasm324 with low confidence (Def.12.1); and cytoplasm322 spatially matches with cytosol234 with low confidence (Def.12.3).

6 Finding the Possible Next Events

In this section, we demonstrate the usefulness of matching instances (Definition 10 and 12) in finding the possible next event(s) of a given event. While in simple cases (Section 4.1), we can find the next event $E'$ of an event $E$ by using Definition 3 there are still cases where there exists no ordering edges from $E$ to $E'$. For examples, Alteration of mrna ends and RNA splicing are two subevents of RNA processing but no other relation between them was defined. However, they all occur in nucleus16421 and Alteration of mrna ends’s output, pre_mrna7690, matches with RNA splicing’s input, rna8697. This information hints us that RNA splicing is Alteration of mrna ends’s next event.

Following this intuition, our approach for finding the next event is that $E'$ is the next event of $E$ if the output of $E$ matches the input of $E'$ and output
location of $E$ matches the input location of $E'$. In the example in Figure 2, this assumption holds in all three events: Eukaryotic transcription, RNA processing and Move_out; all of which are already defined in our KB as consequent events. This assumption also suggests that Eukaryotic translation can be the next event of either Synthesis of RNA in eukaryote or Move_out. Armed with Definition 10 and 12, we define the following join relation.

**Definition 13.** Let $A$ and $B$ be two events in $KDG(Z)$. Event $A$ joins to event $B$ if all of the following conditions are true:

1. The output of $A$ matches with the input of $B$ or vice versa.
2. The output location of $A$ spatially matches with the input location of $B$ or vice versa.

Applying this definition, we have: Alteration of mrna ends joins to RNA splicing, Eukaryotic transcription joins to RNA processing; RNA processing joins to Move_out; and both Synthesis of RNA in eukaryote and Move_out are joined from Eukaryotic translation. Since we want Eukaryotic translation to be a possible next event of Synthesis of RNA in eukaryote instead of its subevent Move_out, we define the possible next event as follows.

**Definition 14.** Let $A$ and $B$ be two events in $KDG(Z)$ where $A$ joins to $B$. $B$ is a possible next event of $A$ if none of the following conditions is true:

1. $A$ joins to Ancestor$B$ where Ancestor$B$ is the ancestor event of $B$ (in other words, there is a non-empty path of subevent relation from Ancestor$B$ to $B$).
2. Ancestor event Ancestor$A$ of $A$ joins to $B$.
3. $A$ is an ancestor event of $B$.
4. $B$ is an ancestor event of $A$.
5. $A$ and $B$ have the same ancestor event.

In our example, condition 14.2 gives us that Eukaryotic translation is not the possible next event of Move_out while 14 concludes that Eukaryotic translation is the possible next event of Synthesis of RNA in eukaryote. We assume that an event and its subevents are put in our KB as a whole, so the next event relations between them are well defined. Thus conditions 14.3-5 take those relations out of consideration.

When we have a path of possible next events, we can create an event $SE$, which is the super event of all events in the path, and add suitable next event or subevent relations. This new event would link the events that were mistakenly encoded as independent events (that we mentioned earlier).

### 7 ASP Encodings

In this section we give ASP encodings of our formulations in the previous sections.

**Encoding the Entities and Events:** Rules t1-t2 in the following state that an instance $X$ is an event or an entity if and only if it is the instance of event
class or entity class respectively. Rules t3-t4 identify E as an event if there is an ordering edge to E (Definition 4(ii)). Rule t5 encodes Definition 4(iv). “has(X, ancestorclass, Y)” denotes the transitive closure of “has(M, superclass, N)” and is encoded the standard way (rules t6-t7). The rest of Definition 4 are encoded similarly in rules t8-t21.

t1: event(X) :- has(X, instance_of, event).
t2: entity(X) :- has(X, instance_of, entity).
t3: ordering_edge(next_event; enables; causes; prevents; inhibits).
t4: has(E, instance_of, event) :- has(X, S, E), ordering_edge(S).
t5: has(E, instance_of, event) :- has(X, instance_of, ClassY), has(ClassY, ancestorclass, event).

Finding Next Events, First Subevents and Last Subevents: Rules e1-e2 find the next events (Definition 5) and rules e3-e6 find the first subevents and the last subevents (Definition 6).

e1: predicates(ordering_edge, enables; causes; prevents; inhibits).
e2: has(E1, next_event, E2) :- has(S1, Predicate, E2), predicates(ordering_edge, Predicate).
e3: not_fse(Z, E) :- has(Z, subevent, E), has(Z, subevent, E2), E2 != E, has(E2, next_event, E).
e4: not_lse(Z, E) :- has(Z, subevent, E), has(Z, subevent, E2), E2 != E, has(E, next_event, E2).
e5: has(Z, first_subevent, E) :- has(Z, subevent, E), not not_fse(Z, E).
e6: has(Z, last_subevent, E) :- has(Z, subevent, E), not not_lse(Z, E).

Encoding Transport Events and Operational Events: t_event(E) or o_event(E) is used to indicate a transport event or an operational event, respectively.

ev1: predicates(t_event, move_through; move_into; move_out_of).
ev2: t_event(E) :- has(E, instance_of, Transport_class), predicates(t_event, Transport_class), event(E).
ev3: o_event(E) :- event(E), not t_event(E).

Encoding the Inputs and Outputs of Operational Events: We denote the input/output/input location/output location of an event by input, output, input_loc and output_loc respectively. Rules i1-i5 get the IOs of operational events. IOs of transport events are encoded similarly (rules i6-i10).

i1: input(E, A) :- has(E, object, A), o_event(E).
i2: input(E, A) :- has(E, base, A), o_event(E).
i3: input(E, A) :- has(E, raw_material, A), o_event(E).
i4: output(E, A) :- has(E, result, A), o_event(E).
i5: input_loc(E, A) :- has(E, site, A), o_event(E).

Getting the Missing Inputs and Outputs: Rule i11 gets the input of an event from its first subevent (Definition 7). Rule i12 gets the object property of a transport event from its first subevent. Other rules to get the input location, output and output location as well as other properties, such as raw-material, result, are encoded in a similar way (rules i13-i24). Rule i25 gets the default output location of an event (Definition 8).

i11: has(E, input, A) :- has(SE, input, A), has(E, first_subevent, SE).
i12: has(E, object, A) :- has(SE, object, A), has(E, first_subevent, SE), transport_event(SE).
i25: has(E, output_location, A) :- not has(E, output_location, A2), has(E, input_location, A), entity(A2), event(E), A2 != A.
Encoding the Main Class(es) of an Instance: \textit{ClassA} is a main class of instance \textit{A} if \textit{ClassA} is one of \textit{A}'s classes and we do not have \textit{not\_main\_class}(\textit{A}, \textit{ClassA}) (which mean \textit{ClassA} is not the main class of \textit{A}).

\begin{itemize}
  \item \texttt{m1: general\_class(thing; event; entity; spatial\_entity; tangible\_entity; chemical\_entity).}
  \item \texttt{m2: not\_main\_class(A, ClassB) :- has(A, instance\_of, ClassA), has(A, instance\_of, ClassB).}
  \item \texttt{m3: not\_main\_class(A, ClassB) :- has(A, instance\_of, ClassA), has(A, instance\_of, ClassB), general\_class(ClassB), not general\_class(ClassA).}
  \item \texttt{m4: main\_class(A, ClassA) :- has\_class(A, ClassA), not not\_main\_class(A, ClassA).}
\end{itemize}

Encoding Instance Matching: We use predicate \texttt{match\_with}(\textit{A}, \textit{B}, \textit{Confidence}) to represent \textit{match\_with} relation (Definition 10) from instance \textit{A} to \textit{B}; \textit{Confidence} can be either \textit{low}, \textit{medium} or \textit{high}. Rule \texttt{ma1} encodes the sub-case 10.1.a of Definition 10. The last rule is for Definition 10.4, matching \textit{A} to \textit{B} transitively through \textit{C}. \texttt{lowest\_confidence(Con1, Con2, Con)} means \textit{Con} is the lowest confidence in \textit{Con1} and \textit{Con2} (Rules \texttt{l1-lc7}). Rules for other cases of Definition 10 are skipped (rules \texttt{ma2-ma5}); locational instance matching is encoded in a similar way (rules \texttt{sma1-sma4}).

\begin{itemize}
  \item \texttt{ma1: \texttt{match\_with}(A,B,\texttt{high}) :- \texttt{main\_class}(\textit{A}, \textit{ClassA}), \texttt{main\_class}(\textit{B}, \textit{ClassB}), \textit{A==B}.}
  \item \texttt{ma6: \texttt{match\_with}(A,B,\texttt{Conf}) :- \texttt{match\_with}(A,C,\texttt{Conf1}), \texttt{match\_with}(C,B,\texttt{Conf2}), \textit{A!=B, A!=C, B!=C, lowest\_confidence(Conf1, Conf2, Conf)}.}
\end{itemize}

Encoding Possible-next-event Relation: In this section, we show how Definition 14 is encoded. We use \texttt{has(A, tc\_subevent, B)} to represent transitive closure of \textit{subevent} relation between \textit{A} and \textit{B} (encoded by \texttt{has(A, subevent, B)}), which is defined in the standard way (rules \texttt{tcsub1-tcsub3}). We also use \texttt{\_join(A, B)} to encode that \textit{A} joins to \textit{B} according to Definition 13 (rules \texttt{j1-j3}). The two rules below is corresponding to the sub-case 14.1. Other cases are skipped (rules \texttt{n2-n5}).

\begin{itemize}
  \item \texttt{n1: _notNextEvent(A, B) :- \_join(A, SuperB), \_join(A, B), has(SuperB, tc\_subevent, B).}
  \item \texttt{n6: possible\_next\_event(A,B) :- \_join(A, B), not _notNextEvent(A, B).}
\end{itemize}

Correctness of the ASP Rules:

Definition 15. The ASP program $\Pi_Z$ is the answer set program consisting of the facts of the form \texttt{has(X,S,V)}" that are generated from all the nodes and edges of \textit{KDG} in the following way:

1. For each node \textit{N}, generate \texttt{has(N,instance\_of,event)} if \textit{N} is event node, \texttt{has(N,instance\_of,entity)} if \textit{N} is entity nodes.
2. For each edge of relation \textit{R} (Table 7) from \textit{E1} to \textit{E2}, generate \texttt{has(E1, R, E2)}".

Definition 16. The ASP program $\Pi$ is the answer set program consisting of the following rules: t1 to t14 for events and entities, e1 to e6 for next events, first subevents and last events, ev1 to ev3 for two types of events, i1 to 25 for inputs, outputs of events, m1 to m4 for main class(es), lc1 to lc7 for the lowest confidence, ma1 to ma6 for match relation, sma1 to sma4 for spatially match relation, tcsu1 to tcsu2 for transitive closure of subevents, j1 to j3 for joined events, and n1 to n6 for possible next events.
Proposition 1: \( E \) is the last subevent of \( X \) in \( KDG(Z) \) iff 
\[ \Pi_Z \cup \Pi \models \text{has}(Z, \text{last} \_\text{subevent}, E) \]

Proposition 2: \( A \) is the main class of \( E \) in \( KDG(Z) \) iff 
\[ \Pi_Z \cup \Pi \models \text{main class}(E, A) \]

Proposition 3: Let \( A \) and \( B \) be two instances in \( KDG(Z) \). \( A \) matches with \( B \) with the confidence level \( \text{Conf} \) iff 
\[ \Pi_Z \cup \Pi \models \text{match with}(A, B, \text{Conf}) \]

Proposition 4: Let \( A \) and \( B \) be two events in \( KDG(Z) \). \( A \) is a possible next event of \( B \) iff 
\[ \Pi_Z \cup \Pi \models \text{possible next event}(A, B) \]

8 Discussion: Answering “How” and “Why” Questions

In Section 4, we showed how to recover missing information using properties of 
\( KDG \)’s structure. Completing this information not only allows us to improve the \( KB \) that was used to construct the \( KDG \), but also make it possible to reason 
about large curated \( KB \) using \( KDG \). In Section 5 and 6, we also solved an 
important step in bringing the \( KB \)’s usage out of small examples: we proposed the 
methods to compare instances and demonstrated their power in finding possible 
next events.

Those efforts have enabled us to answer deep reasoning questions, such as 
“How” and “Why” questions. We give examples of a few of them in the following.
Details about answering them are explained in another work of ours [2].

1. The answer of “How does \( X \) occur?” is simply a structure that basically 
contains \( KDG(X) \) and all the nodes connected to/from \( X \) through ordering 
edges.
2. The answer of “How does \( X \) produce \( Y \)?” is similar to “How does \( X \) occur?” 
but \( X \) must produce \( Y \).
3. The answer of “How are \( X \) and \( Y \) related?” is a simplified structure of 
\( KDG(Z) \) that contains: two paths of component edges to \( X \) and \( Y \) from 
their lowest common ancestor and all paths of ordering edges linking two 
nodes in those two paths.
4. Similarly, the answer of “Why \( X \) is important to \( Y \)?” is the answer of “How 
are \( X \) and \( Y \) related?” plus the path on “important” links which explains 
why \( X \) is important to \( Y \). An “important” link from \( A \) to \( B \) is defined in 
\( KDG \) to indicates that \( A \) is important to \( B \).
5. Other questions that \( KDG \) can answer includes “How does \( X \) participate in 
process \( Y \)?”, “How does \( X \) do \( Y \)?”, “Why does \( X \) produce \( Y \)?” and others.

9 Conclusion

In this paper we have shown how to derive certain missing information from large 
knowledge bases. Often such knowledge bases are created by multiple people; 
sometimes even through crowd-sourcing. This often leads to some information 
being not explicitly stated, even though the knowledge base contains clues to 
derive that information. In our larger quest to formulate answers to “why” and 
“how” questions, we focused on the frame based knowledge base AURA, noticed 
several such omissions, and using those as examples, developed several general 
formulations regarding missing knowledge about events. We also gave an ASP
implementation of our formulations and used them in answering “why” and “how” questions. We briefly discussed some of those question types and how their answer can be obtained from Knowledge Description Graphs (KDGs). Thus, by being able to obtain missing information and enriching the original KDGs one can obtain more accurate and intuitive answers to the various “why” and “how” questions.

One of our formulations was about entity resolution where we resolve multiple entities that may have different names but may refer to the same entity. Our method is different from other methods in the literature \[4,5\]. Since each entity resolution method heavily relies on the properties of the database it is working on, and no other system we know of is about AURA or similar event centered knowledge bases we were unable to directly compare our method with the others.

Our approach to use rules (albeit ASP rules) to derive missing information is analogous to use of rules in data cleaning and in improving data quality \[10,12\]. However those works do not focus on issues that we discussed in this paper.

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