Discovering Alignments in Ontologies of Linked Data*

Rahul Parundekar, Craig A. Knoblock, and José Luis Ambite

University of Southern California Information Sciences Institute 4676 Admiralty Way Marina del Rey, CA 90292

Abstract

Recently, large amounts of data are being published using Semantic Web standards. Simultaneously, there has been a steady rise in links between objects from multiple sources. However, the ontologies behind these sources have remained largely disconnected, thereby challenging the interoperability goal of the Semantic Web. We address this problem by automatically finding alignments between concepts from multiple linked data sources. Instead of only considering the existing concepts in each ontology, we hypothesize new composite concepts, defined using conjunctions and disjunctions of (RDF) types and value restrictions, and generate alignments between them. In addition, our techniques provide a novel method for curating the linked data web by pointing to likely incorrect or missing assertions. Our approach provides a deeper understanding of the relationships between linked data sources and increases the interoperability among previously disconnected ontologies.

1 Introduction

The last few years have witnessed a paradigm shift from publishing isolated data to publishing data that is *linked* to related data from other sources using the structured model of the Semantic Web. By doing so, the publishers of the linked data are able to supplement their own knowledge base, by integrating data from different sources, and realize significant benefits across various domains. Most of the effort has been on identifying which objects from different sources are actually the same. For example, that object *geonames.org/5368361* is the same as *dbpedia:Los_Angeles*. Despite the increasing availability of linked data, the absence of links at the concept level has resulted in heterogenous schemas, challenging the interoperability goal of the Semantic Web. For example, of the 190 sources in the latest census of linked data¹ only 15 have mappings between their ontologies.

The problem of schema linking (aka schema matching in databases and ontology alignment in the Semantic Web) has received much attention [Bellahsene et al., 2011; Euzenat and Shvaiko, 2007; Bernstein et al., 2011; Gal, 2011]. In this paper we present a novel extensional approach to generate alignments between ontologies of linked data sources. Similar to previous work on instance-based matching [Duckham and Worboys, 2005; Doan et al., 2004; Isaac et al., 2007], we rely on linked instances to determine the alignments. Two concepts are equivalent if all (or most of) their respective instances are linked (by owl:sameAs or similar links). However, our search is not limited to the existing concepts in the ontology. We hypothesize new concepts by combining existing elements in the ontologies and seek alignments between these more general concepts. This ability to generalize allows us to find many more meaningful alignments in ontologies in which one-to-one concept equivalences might not exist. For example, the alignment of an impoverished ontology like GeoNames, which has only one class - geonames: Feature, with the well-developed DBpedia ontology is not particularly informative. To successfully link such ontologies, we first generate more expressive concepts, based on properties and values of the instances in the sources. For example, in GeoNames the values of the featureCode and featureClass properties can be used to find alignments with existing concepts in DBpedia, such as the alignment of the concept geonames:featureClass=P to dbpedia:PopulatedPlace.

Our approach finds alignments between concepts defined by conjunction and disjunctions of (RDF) type and value restrictions (cf. [Horrocks *et al.*, 2006]), which we call *restriction classes* henceforth. An *atomic restriction class*, $\{p = v\}$, is the set of objects having object (or data) property *p* (including rdf:type) with object (or literal) value *v*. These alignments are based on the linked instances between these composite concepts. This is an important feature of our approach; we model the *actual* contents and relationships between sources, as opposed to what ontologies disassociated from the data may lead us to believe based on class names or structure.

2 Sources with Heterogenous Ontologies

Linked data sources often conform to different, but related, ontologies that can be meaningfully linked [Cruz *et al.*, 2011; Jain *et al.*, 2011; Parundekar *et al.*, 2010; 2012]. Our algorithms are generic and can be used to align any two linked

^{*}The paper on which this extended abstract is based was the recipient of the best paper award in the research track at the 11th International Semantic Web Conference, 2012 [Parundekar *et al.*, 2012].

¹http://www4.wiwiss.fu-berlin.de/lodcloud/state/

sources. However, we will use two sources with geospatial data for better illustration of our approach. *GeoNames* (geonames.org), contains about 7.8 million geographical objects. It is described by a rudimentary ontology since its semantic web version was generated automatically by direct translation of a simple relational database model. All its instances belong to a single class, *Feature*, with the type of the geographical data (e.g. mountains, lakes, cities, etc.) encoded in the *feature-Class* and *featureCode* properties. *DBpedia* (dbpedia.org) is a knowledge base that covers multiple domains and includes approximately 526,000 geographical objects. It is described using a rich ontology with extensive concept hierarchies and numerous relations. At the time of our experiments, these two sources have over 86,000 pairs of instances linked using *owl:sameAs* assertions.

3 Finding Alignments Across Ontologies

We find three types of alignments between the ontologies of linked data sources. First, we extract equivalent and subset alignments between *atomic restriction classes*. These are the simplest alignments that we define. Though simple, they often yield interesting alignments. Moreover, we use them as seed hypotheses to find alignments that are more descriptive. Second, we find alignments between *conjunctive restriction classes* in the two sources. Finally, we find *concept coverings*, which are alignments where a concept from one source maps to a union of smaller concepts from the other source.

Before searching for alignments, we pre-process the sources to reduce the search space and avoid computation not leading to meaningful alignments. First, we only consider instances that are actually linked, thus removing unrelated instances and their properties. Second, we eliminate inverse (or quasi-inverse) functional properties, since a *restriction class* on such a property would only contain a single instance (or very few) and would not be a useful concept (e.g., the latitude and longitude properties generally point to one place).

3.1 Aligning Atomic Restriction Classes

Atomic restriction classes can be generated by combining properties and values in the sources and tested for alignments using the simple algorithm in Fig. 1. Fig. 2 illustrates the set comparison operations of our algorithm. We consider the two concepts equivalent if they significantly overlap each other. We use two metrics P and R to measure the degree of overlap between *restriction classes*. In a perfect equivalence alignment, the values for P and R would be both 1. However, to allow for missing links or errors in the sources, we use $P \ge \theta$ and $R \ge \theta$ ($\theta = 0.9$ in our experiments). For example, consider the alignment between restriction classes {geonames:countryCode=ES} and {dbpedia:country = dbpedia:Spain }. Based on the concept extensions, our algorithm finds $|Img(r_1)| = 3198$, $|r_2| = 4143$, $|Img(r_1) \cap r_2|$ = 3917, R = 0.9997 and P = 0.9454. Thus, the algorithm considers this alignment as equivalent in an extensional sense. This algorithm finds numerous equivalent and subset alignments between atomic restriction classes. For example, we find that each of {geonames:featureCode = S.SCH} and $\{geonames: featureCode = S.UNIV\}$ (i.e. Schools and Universities from *GeoNames*) are subsets of {*dbpedia:-EducationalInstitution*}.

```
function ATOMICALIGNMENTS(Source_1, Source_2)

for all properties p_1 in Source<sub>1</sub>, all distinct values v_1 \in p_1,

all p_2 in Source<sub>2</sub>, and all distinct v_2 \in p_2 do

r_1 \leftarrow \{p_1 = v_1\} // instances of Source<sub>1</sub> with p_1 = v_1

r_2 \leftarrow \{p_2 = v_2\}

Img(r_1) \leftarrow instances of Source<sub>2</sub> linked to those in r_1

P \leftarrow \frac{|Img(r_1)\cap r_2|}{|r_2|}, R \leftarrow \frac{|Img(r_1)\cap r_b|}{|r_1|}

alignment(r_1, r_2) \leftarrow [

if P \ge \theta and R \ge \theta then r_1 \equiv r_2

else if P \ge \theta then r_1 \subset r_2

else if R \ge \theta then r_2 \subset r_1

end if]

end for

end function
```





Figure 2: Comparing linked instances from two ontologies

3.2 Aligning Conjunctive Restriction Classes

The second type of alignments we detect are those between *conjunctive restriction classes*. For example, the *conjunctive restriction class* 'Schools in the US', {*geonames:countryCode=US* \cap *geonames:featureCode=S.SCH*}, is the intersection of the *atomic restriction classes* representing all schools in *GeoNames* and all things in the US.

We seed the search space with the alignments generated by ATOMICALIGNMENTS. Taking one hypothesis at a time, we can generate new hypothesis by conjoining *atomic restriction classes*. For example, we can extend the alignment [{geonames:featureCode=S.SCH}, {rdf:type=EducationalInstitution}] by conjoining {geonames:featureCode=S.SCH} with {geonames:countryCode=US}, and investigate the relationship between schools in the US and educational institutions.

The algorithm to find *conjunctive restriction classes* appears in Fig. 3 (cf. [Parundekar *et al.*, 2010]). To reduce the combinatorial search space, our algorithm prunes hypotheses that 1) do not have enough instances supporting either of the *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2) where the extension of the refined *restriction classes*; 2, where the extension of the refined *restriction classes*; 2, where *r'*; is a subclass of *r*; and *r*; *c*; *r*; since no immediate specialization is provided; and 4) would have been explored more than once (these are avoided by using a lexicographic ordering). Finally, the algorithm removes any

implied alignments that are generated because of the hierarchical nature of the algorithm. Specifically, it removes 1) relations that implied by the transitivity of the subset relations; 2) cyclic equivalences which may be generated due to the reduced threshold θ .

```
function CONJUNCTIVEALIGNMENTS(Source<sub>1</sub>,Source<sub>2</sub>)
    for all [r_1, r_2] \in \text{ATOMICALIGNMENTS}(Source_1, Source_2)
do EXPLOREHYPOTHESES(r_1, r_2, Source_1, Source_2)
    end for
    REMOVEIMPLIEDALIGMENTS
end function
function EXPLOREHYPOTHESIS(r_1, r_2, Source_a, Source_b)
    for all p_a in Source<sub>a</sub> occurring lexicographically after all the
properties in r_1 and distinct v_a associated with p_a do
        r_1' \leftarrow r_1 \cap \{p_a = v_a\}
        alignment \leftarrow FINDALIGNMENT(r'_1, r_2)
        if not SHOULDPRUNE(r'_1, r_2, alignment) then
            alignment(r'_1, r_2) \leftarrow alignment
            EXPLOREHYPOTHESES(r'_1, r_2)
        end if
    end for
    for all p_b in Source<sub>b</sub> occuring lexicographically after all the
properties in r_2 and distinct v_b associated with p_b do
        r_2' \leftarrow r_2 \cap \{p_b = v_b\}
        alignment \leftarrow FINDALIGNMENT(r_1, r'_2)
        if not SHOULDPRUNE(r_1, r'_2, alignment) then
            alignment(r_1, r'_2) \leftarrow alignment
            EXPLOREHYPOTHESES(r_1, r'_2)
        end if
    end for
```

end function

Figure 3: Aligning conjunctive restriction classes

3.3 Finding Concept Coverings

The CONJUNCTIVEALIGNMENTS algorithm may produce a very large number of subset relations. Analyzing the results of [Parundekar *et al.*, 2010], we noticed that these subset alignments follow common patterns. For example, we found that both Schools and Universities from *GeoNames* were subsets of Educational Institutions from *DBpedia*. However, the *union* of Schools, Colleges, and Universities from *GeoNames* was *equivalent* to *dbpedia:EducationalInstitution*, which is a more informative finding.

The algorithm for generating *concept coverings* appears in Fig. 4. We start with the subclass alignments found by ATOM-ICALIGNMENTS. Then we identify concepts from one ontology that are defined on the same property and are subsets of another concept in the other ontology. We test whether the union of the smaller concepts is equivalent to the larger concept based on the extensions of the concepts as before. Although we could explore more complex hypotheses, this approach is tractable and generates intuitive definitions.

Since all smaller classes are subsets of the larger *restriction* class, $P_U \ge \theta$ holds by construction. Thus, we just need to check that $R_U \ge \theta$ to determine whether the union *restriction* class is equivalent to the single concept. The smaller *restriction* classes that were omitted in ATOMICALIGNMENTS because of insufficient support size of their intersections (e.g.,

function CONCEPTCOVERINGS(Source₁,Source₂)

for all alignments $[U_L, r_2] \in ATOMICALIGN-MENTS(Source_1, Source_2)$, with larger concept $U_L = \{p_L = v_L\}$ from $Source_1$ and multiple classes $r_2 = \{p_S = v_i\}$ from $Source_2$ that can be partitioned on property p_S do for all smaller concepts $\{p_S = v_i\}$ do $U_S \leftarrow \{p_S = \{v_1, v_2, ...\}\}$ // union restriction class $U_A \leftarrow Img(U_L) \cap U_S, P_U \leftarrow \frac{|U_A|}{|U_S|}, R_U \leftarrow \frac{|U_A|}{|U_L|}$ if $R_U \ge \theta$ then $alignment(r_1, r_2) \leftarrow U_L \equiv U_S$

end if end for end for





 $\{geonames: featureCode = S.SCHC\}$) are included in constructing U_S for completeness.

Figure 5 illustrates the approach. ATOMICALIGNMENTS detects that {geonames:featureCode = S.SCH}, {geonames:featureCode = S.SCHC}, and {geonames:featureCode = S.UNIV} are subsets of {rdf:type = dbpedia:EducationalInstitution}. As can be seen in the Venn diagram in Figure 5, U_L is $Img({rdf:type = dbpedia:EducationalInstitution}), U_S$ is {geonames:featureCode = S.SCHC} \cup {geonames:featureCode = S.UNIV}, and U_A is the intersection of the two. Upon calculation we find that R_U for the alignment of *dbpedia:EducationalInstitution* to {geonames:featureCode = {S.SCHC}, S.UNIV} is 0.98 (greater than θ). We can thus confirm the hypothesis and consider U_L and U_S as equivalent.



Figure 5: Concept covering: Educational Institutions

4 Results

The results of the three alignment algorithms over *GeoNames* and *DBpedia* appear in Table 2. In all, we were able to detect about 580 (263 + 45 + 221 + 51) equivalent alignments including both atomic and complex *restriction classes*, along with 15,376 (4,946 + 5,494 + 4,400 + 536) subset relations.

Table 1 shows some of the representative alignments found by the three algorithms. We are able to detect alignments of *atomic restriction classes* with existing RDF concepts (i.e., defined with *rdf:type*) and with value restrictions. For example, alignment #1 shows that the feature class 'H' in *GeoNames* maps to a 'Body of Water' in *DBpedia*. Alignments #2 and #3 show equivalence and subset relations between value restrictions. Alignment #4 shows a conjunctive alignment for 'Populated Places in the US'. Finally, alignment #5

#	GeoNames concept	Rel.	DBpedia concept	P	R	$ I(r_1) \cap r_2 $
1	geonames:featureClass=geonames:H	=	rdf:type=dbpedia:BodyOfWater	0.91	0.99	1939
2	geonames:countryCode=ES	=	dbpedia:country=dbpedia:Spain	0.95	0.99	3917
3	geonames:featureCode=geonames:T.MT	C	rdf:type=dbpedia:Mountain	0.97	0.78	1721
4	geonames:featureClass=geonames:P &	=	rdf:type=dbpedia:PopulatedPlace &	0.97	0.96	26061
	geonames:countryCode=US		dbpedia:country=dbpedia:United_States			
5	geonames:featureCode =	=	rdf:type =	-	0.98	396
	{S.SCH, S.SCHC, S.UNIV}		dbpedia: Educational Institution			

Table 1: Representative alignments found in two sources

atomic restriction classes	Alignments	
Equivalent Alignments	263	
Subclasses with larger class from GeoNames	4,946	
Subclasses with larger class from DBpedia	5,494	
conjunctive restriction classes		
Equivalent Alignments	45	
Subclasses with larger class from GeoNames	4,400	
Subclasses with larger class from DBpedia	536	
concept coverings		
Coverings with larger class from GeoNames	221	
Coverings with larger class from DBpedia	51	

Table 2: Alignments found between GeoNames and DBpedia

shows the covering of 'Educational Institutions' in *DBpedia* with schools, colleges and universities in *GeoNames*. None of these alignments could be detected by previous algorithms that do not hypothesize concepts beyond the existing classes. Also, note that the alignments generated from *actual* data need not match the intentional similarity of the concepts. For example, Mountains from *GeoNames* are subset of Mountains in *DBpedia*, since *GeoNames* divides the concept by distinguishing Peaks, Hills, etc., from Mountains.

An interesting outcome of our approach is the detection of outliers, which suggest possible erroneous links or value assignments. For example, in alignment #2, *R* overlap (0.99) is not complete (1) since one outlier instance from *GeoNames* has 'IT' (Italy) as *countryCode*. However, this is likely an error since there is overwhelming support for 'ES' being the *countryCode* of Spain. Alignment #5 shows an interesting case where 8 instances could not be identified as Educational Institutions. They had either a missing *genomes:featureCode* (1) or a value for Library (1), Hospitals (1), Buildings (3), Establishments (1), and Museums (1). The detection of outliers provides a unique opportunity for identifying inconsistencies and automatically curate the web of linked data.

5 Related Work

Even though most previous work on linked data focuses on linking instances across different sources, several authors have considered aligning ontologies of linked data sources. Jain et al. [2010] describe the BLOOMS approach, which uses a central forest of concepts derived from topics in Wikipedia. It is, however, unable to find alignments because of the single *Feature* class in *GeoNames*. BLOOMS+ [Jain *et al.*, 2011] aligns linked data ontologies with an upperlevel ontology called Proton. Using contextual information, BLOOMS+ finds an greater number of alignments between GeoNames & Proton and DBpedia & Proton than its predecessor. Cruz et al. [2011] describe a dynamic ontology mapping approach called AgreementMaker that uses similarity measures along with a mediator ontology to find mappings using the labels of the classes. The advantage of our approach is that, by hypothesizing novel concepts (restriction classes), it can find a larger set of alignments than previous approaches, even from sources described using a rudimentary ontology, such as GeoNames. Völker et al. [2011] describe an extensional approach that uses statistical methods for finding alignments by generating OWL-2 axioms using an intermediate associativity table of instances and concepts and mining associativity rules from it. GLUE [Doan et al., 2004] is a instance-based matching algorithm, which predicts the concept in the other source that instances belong to by using machine learning. GLUE then hypothesizes alignments based on the probability distributions obtained from the classifications. In contrast, our approach is based on the existing links, and hence reflects the nature of the source alignments in practice. CSR [Spiliopoulos et al., 2008] aligns a concept from one ontology to a union of concepts from another ontology using the similarity of properties as features in predicting the subsumption relationships. It differs from our approach in that it uses a statistical machine learning approach for detection of subsets rather than the extensional approach. Atencia et al., [2012] provide a formalization of weighted ontology mappings that is applicable to extensional matchers like ours.

6 Conclusion

We described an approach to identifying alignments between atomic, conjunctive and disjunctive *restriction classes* in linked data sources. Our approach discovers alignments where concepts at different levels in the ontologies of two sources can be mapped even when there is no direct equivalence or only rudimentary ontologies exist. Our algorithm is also able to detect outliers that help identify erroneous links or inconsistencies in the linked instances. By using the *GeoNames* and *DBpedia* sources as an example, we showed that the results the algorithm generates can provide a deeper insight into the nature of the alignments of linked data.

In future work, we plan to explore more expressive concept descriptions and provide a curation system that not only signals outliers, but also proposes corrections automatically.

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