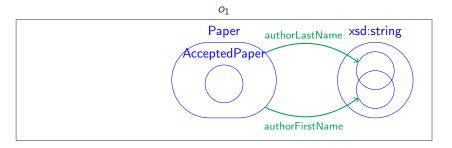
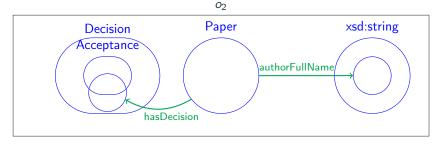
Task-Oriented Complex Ontology Alignment: Two Alignment Evaluation Sets

Elodie Thiéblin Ollivier Haemmerlé Nathalie Hernandez Cássia Trojahn 15th Extended Semantic Web Conference

IRIT & Université de Toulouse 2, Toulouse, France firstname.lastname@irit.fr

Context & Definitions





Retrieving all the accepted papers

```
o_1 o_1:myPaper1: o_1:AcceptedPaper
```

 o_2 (o_2 :myPaper2, o_2 :myAcceptance): o_2 :hasDecision

 o_2 :myAcceptance : o_2 :Acceptance

Retrieving all the accepted papers

- o_1 $o_1:myPaper1: o_1:AcceptedPaper$
- o₂ (o₂:myPaper2,o₂:myAcceptance) : o₂:hasDecision o₂:myAcceptance : o₂:Acceptance
 - Ontology merging
 - Query rewriting
 - Data translation

Retrieving all the accepted papers

- $o_1 o_1:myPaper1: o_1:AcceptedPaper$
- o₂ (o₂:myPaper2,o₂:myAcceptance) : o₂:hasDecision o₂:myAcceptance : o₂:Acceptance
 - Ontology merging
 - Query rewriting
 Ontology Alignments!
 - Data translation

Ontology Alignment [Euzenat and Shvaiko, 2013]

$$A_{o_1 \rightarrow o_2} = \{c_0, c_1, ..., c_n\}$$

Source ontology

Ontology Alignment [Euzenat and Shvaiko, 2013]

$$A_{o_1 \rightarrow o_2} = \{c_0, c_1, ..., c_n\}$$

Target ontology

Ontology Alignment [Euzenat and Shvaiko, 2013]

$$A_{o_1 \rightarrow o_2} = \{c_0, c_1, ..., c_n\}$$

Set of correspondences

Correspondence

$$c_i = \langle e_{o_1}, e_{o_2}, r \rangle$$

Correspondence

$$c_i = \langle e_{o_1}, e_{o_2}, r \rangle$$

Relation
$$(\equiv, \sqsubseteq, \supseteq)$$

Correspondence

$$c_i = \langle e_{o_1}, e_{o_2}, r \rangle$$

Members: Atomic entities or constructions

Simple vs complex

Simple correspondence

- e_{o_1} atomic entity
- e_o, atomic entity
- $o_1:Paper \equiv o_2:Paper$

Simple vs complex

Simple correspondence

- e_{o1} atomic entity
- e_{o_2} atomic entity
- $o_1:Paper \equiv o_2:Paper$

Complex correspondence

- at least one of e_{o_1} , e_{o_2} contains a constructor or a transformation
- constructor o₁:AcceptedPaper ≡ ∃o₂:hasDecision.o₂:Acceptance
- transformation function
 o2:authorFullName =
 o1:authorFirstName + " " +
 o1:authorLastName

Correspondence patterns

Correspondences can be categorised and decomposed into correspondence patterns [Scharffe, 2009].

Class By Attribute Type (CAT): $A \equiv \exists b. C$

 o_1 :AcceptedPaper $\equiv \exists o_2$:hasDecision. o_2 :Acceptance is a CAT

Artefact for analysing the correspondences

Can guide the matching process

Tasks

Ontology merging: Create a new ontology from o_1 and o_2 using $A_{o_1 \rightarrow o_2}$ and $A_{o_2 \rightarrow o_1}$

Query rewriting: Transformation of a query for o_1 into a query for o_2 using $A_{o_1 \rightarrow o_2}$

Tasks

Ontology merging: Create a new ontology from o_1 and o_2 using $A_{o_1 \rightarrow o_2}$ and $A_{o_2 \rightarrow o_1}$

Query rewriting: Transformation of a query for o_1 into a query for o_2 using $A_{o_1 \rightarrow o_2}$

Assumption: an alignment may be different for query rewriting or ontology merging

 o_2 :authorFullName = o_1 :authorFirstName + " " + o_1 :authorLastName

State of the art

Complex ontology matchers

Many complex matching approaches on different kinds of schemata ¹ Here, focus on ontology matching

	Pattern based	No pattern	
Ontology based	[Ritze et al., 2009,	[Jiang et al., 2016]	
	Ritze et al., 2010]		
Instance based	[Parundekar et al., 2010,	[Qin et al., 2007, Nunes et al., 2011]	
	Parundekar et al., 2012,		
	Walshe et al., 2016]		

¹http:

^{//}semantic-web-journal.net/content/survey-complex-ontology-matching

Evaluation of complex matchers so far

Approach	Dataset	Metrics	
[Ritze et al., 2009]		Precision	
[Ritze et al., 2010]	OAEI Conference ²	Precision	
[Jiang et al., 2016]		Precision, Recall	
[Parundekar et al., 2010]	DBpedia, Geonames,	None	
[Parundekar et al., 2012]	Geospecies, LinkedGeoData,	Precision, Recall	
[Farundekar et al., 2012]	GeneID, MGI	(subset)	
[Walshe et al., 2016]	DP-adia Vara	Precision, Recall	
[vvaisile et al., 2010]	DBpedia, Yago	(subset)	
[Nunes et al., 2011]	None	None	
[Qin et al., 2007]	LIMD, CMII	Precision, Recall	
[Qili et al., 2007]	UMD, CMU	(subset)	

²http://oaei.ontologymatching.org/

Where we are so far

- Complex ontology matching: a developing field
- No reference dataset
- No adapted evaluation metric

Methodology and dataset

Global methodology

Assumptions/choices

- Equivalence prefered over subsumption
- Simple correspondences prefered over complex ones

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Assumptions/choices

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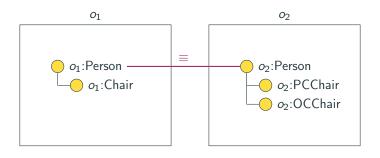
Global methodology

- 1. Simple equivalence correspondences o_1 o_2
- 2. Creation of the complex correspondences given a task
- 3. Write the correspondences in a reusable format (e.g., EDOAL)

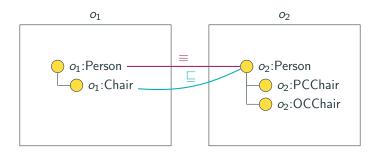
Methodology for an ontology merging alignment

Top-down approach for ontology merging

- 1. Create a new ontology importing o_1 and o_2
- 2. Insert the simple equivalence correspondences as axioms
- 3. Find for each entity of o_1 (then o_2) an equivalence or the closest super-entity not inferred
 - Top-down approach : allows inferrence

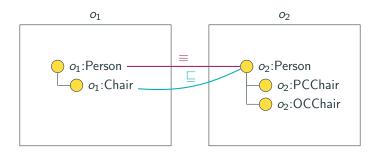


Simple equivalence : $o_1:Person \equiv o_2:Person$



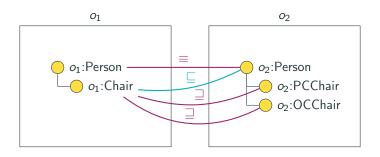
Simple equivalence : $o_1:Person \equiv o_2:Person$

Inferred axiom: $o_1:Chair \sqsubseteq o_2:Person$



Simple equivalence : $o_1:Person \equiv o_2:Person$ Inferred axiom: $o_1:Chair \sqsubseteq o_2:Person$

ullet $o_1
ightarrow o_2$: No equivalence for o_1 :Chair + axiom inferred



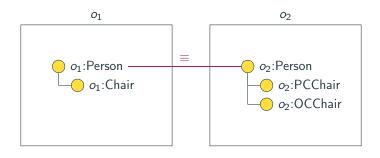
Simple equivalence : $o_1:Person \equiv o_2:Person$ Inferred axiom: $o_1:Chair \sqsubseteq o_2:Person$

- $o_1 \rightarrow o_2$: No equivalence for o_1 : Chair + axiom inferred
- ullet $o_2
 ightarrow o_1$: New correspondences added
 - o₂:PCChair ⊆ o₁:Chair
 - o₂:OCChair ⊑ o₁:Chair

Methodology for a query rewriting alignment

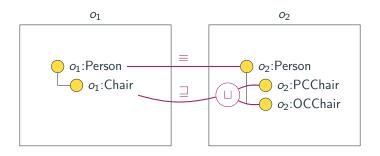
- 1. Find for each entity of o_1 (then o_2) an equivalence or the closest sub-entity or construction
 - The expressiveness of the alignment may prevent a reasoning task from being decidable
 - Precision favoured over recall

Example of query rewriting methodology



Simple equivalence : $o_1:Person \equiv o_2:Person$

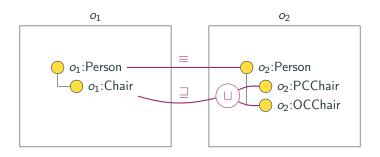
Example of query rewriting methodology



Simple equivalence : $o_1:Person \equiv o_2:Person$

• $o_1 \rightarrow o_2$: o_1 :Chair $\supseteq o_2$:PCChair $\sqcup o_2$:OCChair

Example of query rewriting methodology



Simple equivalence : $o_1:Person \equiv o_2:Person$

- $o_1 \rightarrow o_2$: o_1 :Chair $\supseteq o_2$:PCChair $\sqcup o_2$:OCChair
- ullet $o_2
 ightarrow o_1$: No new correspondences
 - o₂:PCChair has no subsumed entity/construction
 - o2:OCChair has no subsumed entity/construction

Alignment sets & Evaluation

Alignment sets

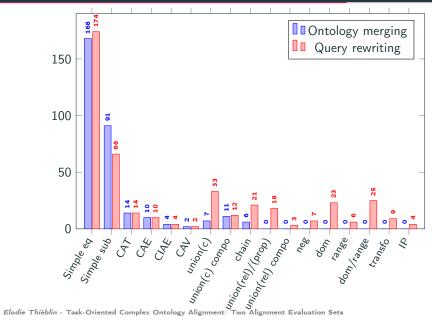
5 ontologies of the OntoFarm Conference dataset $2 \times (10 \text{ pairs } (20 \text{ oriented pairs}))$ cmt, conference, confOf, edas, ekaw

	Simple	Complex	TOTAL
Ontology merging	259	54	313
Query rewriting	240	191	431

Format: EDOAL, OWL (merging), FOL Available under CC-BY license

https://doi.org/10.6084/m9.figshare.4986368.v7

Analysis of the dataset by pattern



Manual evaluation of existing approaches on the dataset

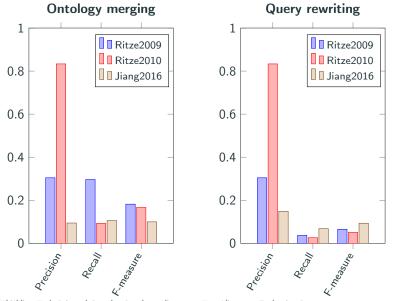
Manual evaluation of the complex correspondences (precision, recall)

Only consider the \mathcal{SROIQ} correspondences for ontology merging

Consider all correspondences for query rewriting

Relation (\equiv , \sqsubseteq , \supseteq) not taken into account

Manual evaluation of existing approaches on the dataset



Discussion & Conclusion

Discussion

Manual creation of the alignments

Bias on the interpretation of the ontologies Consensus dataset currently in work

Limitations (and future works)

Ontologies are not populated

No task-oriented evaluation

Metrics for automatic evaluation: semantic precision and recall ?

Conclusion

- Need of complex alignments
- A complex alignment evaluation dataset
- Transformation into a benchmark with instances and evaluation metrics

https://doi.org/10.6084/m9.figshare.4986368.v7

Thank you! Questions?

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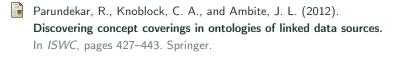


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