A Context-based Measure for Discovering Approximate Semantic Matching between Schema Elements

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- Finding semantic correspondences between 2 schemas still a challenging issue
- Semi automatic matchers available based on several approaches (combination of terminological measures, structural rules, ...)

Motivations

Terminological measures are not sufficient, for example:

- $\bullet\,$ mouse (computer device) and mouse (animal) $\Rightarrow\,$ polysemia
- $\bullet\,$ university and faculty $\Rightarrow\,$ totally dissimilar labels

Structural measures have some drawbacks:

- propagating the benefit of irrelevant discovered matches to the neighbour nodes increases the discovering of more irrelevant matches
- not efficient with small schemas

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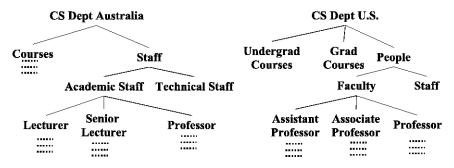


Figure: Two schemas from the university domain.

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Our approach: Approxivect

Based on the work of [1], Approxivect evaluates the similarity between two terms from different schema trees. It has the following properties:

- it is based on the combination of terminological measures (Levenhstein and n-grams) and structural measures (cosine measure applied to contexts)
- it is both automatic and not language-dependent
- it does not rely on dictionaries or ontologies
- it provides an acceptable matching quality





Figure: XML schemas relative to university.

- 3grams(Courses, GradCourses) = 0.2
- Lev(Courses, GradCourses) = 0.42
- \Rightarrow StringMatching(Courses, GradCourses) = 0.31



Figure: In the second schema, *Courses* replaces *GradCourses* due to StringMatching value.

- StringMatching(Faculty, University) = 0.002
- Context(Faculty) = Faculty, Courses, Professor
- Context(University) = University, Courses, Professor

\Rightarrow CosineMeasure(Context(Faculty), Context(University)) = 0.37

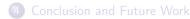
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Context of node n_c

- represents the most important neighbour nodes n_i for n_c
- each neighbour n_i is assigned a weight depending on the relationship n_c

$$\omega(n_c, n_i) = 1 + rac{K}{\Delta d + |level(n_c) - level(n_a)| + |level(n_i) - level(n_a)|}$$

String Matching is the average between

- Levenhstein distance
- 3-grams

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Discovering semantic similarities:

- String Matching between 2 node labels
- if above a given threshold, replacement of one of the label by the other.

Cosine Measure using context:

• due to replacements, the contexts of two nodes can be very similar

Similarity between two nodes

It is the best value between String Matching and Cosine Measure.

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- NB_LEVELS restricts the context by limiting the number of levels
- MIN_WEIGHT restricts the context by keeping only nodes with a weight above this threshold
- REPLACE_THRESHOLD if the StringMatching between two node labels is above this replacement threshold, then one label is replaced by the other
- K represents the importance given to the context

Flexibility

These parameters allow more flexibility. Tuning them is required in some specific scenarii.

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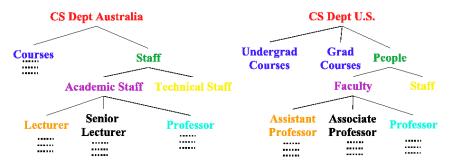


Figure: Mappings discovered by an expert between the schemas.

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Element from schema 1	Element from schema 2	Similarity value	Relevance
Professor	Professor	Professor 1.0	
CS Dept Australia	People 0.46		
Courses	Grad Courses	rad Courses 0.41	
CS Dept Australia	CS Dept U.S. 0.36		+
Courses	Undergrad Courses	Courses 0.28	
Academic Staff	Faculty 0.25		+
Staff	People	0.23	+
Technical Staff	Staff	0.21	+
Senior Lecturer	Associate Professor	ciate Professor 0.16	

Table: Approxivect similarity ranking between the two schemas

Element from schema 1	Element from schema 2 Similarity value		Relevance
Professor	Professor 0.5354546		+
Technical Staff	Staff 0.5300107		+
CS Dept Australia	CS Dept U.S.	0.52305263	+
Courses	Grad Courses	rses 0.5041725 +	
Courses	Undergrad Courses	0.5041725	+

Table: COMA++ discovered mappings between the two schemas

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	Precision	Recall	F-measure
COMA++	1	0.56	0.72
Approxivect	0.62	0.89	0.73

Table: Results of COMA++ and Approxivect on the XML schemas

Note that Approxivect parameters are set to default. An optimal configuration enables to obtain a 0.82 F-measure.

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COMA++ [2]

- combination of many terminological measures and a user-defined synonym table
- a matrix is built for each couple of elements and for each measure
- a strategy is applied to select the mappings
- mappings are modified and/or validated by the user

Similarity Flooding [3]

- a simple string matching algorithm to provide initial matchings
- structural rules and propagation to refine the matchings
- mappings are modified and/or validated by the user

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An automatic schema matching approach

- based on the combination of terminological and structural measures
- with an acceptable quality of matching
- flexible thanks to the parameters

However

- tuning is not automatic, but some tools could handle this step (eTuner)
- more heterogeneity in the experiments

Ongoing work

performance aspect

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