# MULTIPLE KERNEL LEARNING FOR ONTOLOGY INSTANCE MATCHING

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Abstract: This paper proposes to apply Multiple Kernel Learning and Indefinite Kernels (IK) to combine and tune Similarity Measures within the context of Ontology Instance Matching. We explain why MKL can be used in parameter selection and similarity measure combination; argue that IK theory is required in order to use MKL within this context; propose a configuration that makes use of both concepts; and present, using the IIMB bechmark, results of a prototype to show the feasibility of this idea in comparison with other matching tools.

## 1 INTRODUCTION

Ontology matching is the problem of determining correspondences between concepts, properties, and individuals of two or more different formal ontologies (Euzenat and Shvaiko, 2007). The aforementioned plays a key role in many different applications such as data integration, data warehousing, data transformation, open government, peer-to-peer data management, semantic web, and semantic query processing.

Currently, one of its main challenges is the selection and combination of similarity measures during the matching process (Shvaiko and Euzenat, 2008). Although it is broadly accepted that multiple similarity measures can help in finding better alignments and the general opinion supports the idea that there is not a similarity measure that is able to deal with all existing matching problems, we still require to find ways to orchestrate the available similarity measures in order to find the appropriate set of these for the matching task at hand.

Furthermore, even if it is possible to choose the measures that are likely to work within a specific context, the question on how to set the parameters of such functions remains open. Empirical results and literature tell us that similarity measures work but they require prior tuning steps.

To overcome these issues, most of the current proposals use probabilistic or machine learning techniques in order to find the correct combination of measures. This is a natural approach considering that the rules that define how the measures should be composed depend on the real application. Furthermore, even for a domain expert, such rules are not necessarily clear because similarity for humans is a relative and sometimes - contradictory concept (Laub et al., 2007). Within this context, a learning algorithm is a suitable option to find such rules.

In the present paper, our aim is to give new insights to this problem. We propose a matching solution based on the recent research in Multiple Kernel Learning (MKL) and Indefinite Kernels (IK). To our knowledge, there are not any current solutions that propose the use of the algorithms and techniques that are employed in this article. Our main concern is to explore other ways to find the weights that typically need to be determined when a process of aggregation of similarity measures is carried on; therefore, we assume the existence of an available library of similarity measures and aggregation functions from which both of these can be selected.

In a proof of concept prototype, the semisupervised learning paradigm is also integrated, as we believe it to be suitable for this problem. First, because of the volume of instances it is not feasible to compare all the possible instances to find the correct correspondences; for this reason, it is necessary to find rules that can be learned from a small subset of instances. Second, as previously stated, the rules that make two instances equivalent can be difficult to capture, thus, the use of a learning algorithm to find them is better. Third, a huge amount of unlabeled data can be easily obtained in many applications of this problem so it would be great if we could take advantage of such information.

This article is organized as follows: In the next

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section some of the related work is described. Section 3 discusses the suggested approach. Section 4 presents experimental results that validate our approach. Finally, Section 5 discusses the conclusions and future work.

### 2 BACKGROUND AND RELATED WORK

There is a lot of work related to ontology matching. Some of the available reviews are (Kalfoglou and Schorlemmer, 2005) and (Shvaiko and Shvaiko, 2005). However, these reviews focus on the schema level. This is probably because there are relatively few works that prioritize the instance level. In fact, to our knowledge, there is not a comprehensive review involving ontology instance matching systems.

Specifically, concerning the challenge of tuning and selecting similarity measures, the proposals typically attempt to find a linear combination  $\sum c_i S_i$ , where each  $S_i$  is a similarity measure which sometimes is called a matcher, an agent, an expert or a classification hyphotesis. What tends to change within the different works is the manner in which the coefficients are found.

One of the first solutions proposed was using values obtained through empirical evaluation. For example, this approach was used by (Castano et al., 2003) where they set the weights using the data of several real integration cases. Of course, some of the problems of this approach are that it can only be used in very static context and that the process of tuning the parameters can be very expensive or require the sampling of many scenarios in order to have a reliable estimation.

There are works that use different similarity measures as features of a sample so that they can employ toolbox Machine Learning Algorithms. For example (Wang et al., 2006) uses Support Vector Machine (SVM) as the classification model. The training is achieved by creating a set of matched instance pairs with positive labels and a set of non-matched instance pairs with negative labels. A binary classifier is trained by using different similarity measurements as features from the two pair sets. The classifier then acts as a pairing function taking a pair of instances (a,b) as input and generating decision values as output. Since from a Kernel Theory point of view this is equivalent to modifying the spectrum of the Gram matrix by replacing each of the eigenvalues with its square, our approach captures this kind of proposal.

(Ehrig et al., 2005) uses different machine learning techniques for classification (e.g. decision tree learner, neural networks, support vector machines) to assign an optimal internal weighting and threshold scheme for each of the different feature/similarity combinations of a given pair of ontologies. The machine learning methods like C4.5 capture relevance values for feature/similarity combinations. To obtain the training data, they employ an existing parametrization as input to the Parameterizable Alignment Method to create the initial alignments for the two ontologies. The user then validates the initial alignments and thus generates correct training data.

Some systems define a hierarchy of similarity measures that are combined through a preestablished process. This approach allows the systems to define different types of mapping in which the kind of features that are analyzed changes. A system of this kind is HMatch (Castano et al., 2005) that defines four matching models. The idea is that each model reflects different levels of complexity within the matching process. To combine the different similarities, it defines weights according to the characteristics of each feature. For example, each semantic relation has associated a weight  $W_{sr}$  which shows the strength of the connection expressed by the relation on the involved concepts.

(Marie and Gal, 2008) proposes creating an ensemble matcher by treating each similarity matrix M(S, S') as a weak classifier and finding a strong classifier using a modified version of Adaboost. They use a compound measure formed by Precision and Recall as the error function for each iteration. The principles behind kernel theory and boosting are different making it possible to complement this proposal with our ideas.

(Duchateau et al., 2008) introduces the notion of planning to the problem of similarity measure aggregation. Although this is a very interesting idea and we believe it can be used to extent most of the current approaches, the solution currently requires the user to manually create or modify a decision tree. This heavily depends on the user, who does not necessarily know exactly how the similarity measures should be parametrized and aggregated.

Some works propose different operators to combine different similarity measures. For example, (M. Nagy, 2010), based on Dempster Schafer Theory of Evidence (Diaconis, 1978), proposes using the Demspter Combination Rule  $m_{ij}(A) = m_i \oplus m_j =$  $\sum m_i(E_k) * m_j(E_k)$ , where  $m_i, m_j$  are similarity measures and  $E_k$  is the similarity value for a candidate correspondence. Another similar approach is found in (Ji et al., 2008) where they define what is called the Ordered Weighted Average (OWA) operator and use the linguistic quantifiers developed by Yager (Yager, 1988).

Finally, it is worth mentioning works that attempt to formalize the combination task. For example, (Stahl, 2005) investigates aspects of these approaches in order to support a more goal-directed selection as well as initiating the development of new techniques. The investigation is based on a formal generalization of the classic CBR cycle, which allows a more suitable analysis of the requirements, goals, assumptions, and restrictions relevant in learning similarity measures. To simplify the selection of accurate techniques within a particular application as well as for creating foundations for future investigations, the work proposes different categories for each of the following three dimensions of the task of combining similarity measures:

- Semantic of Similarity Measures: Determining the Most Useful Case, Ranking the Most Useful Cases, Approximating the Utility of the Most Useful Cases and Probabilistic Similarity Measures.
- Training Data: Relative Case Utility Feedback, Absolute Utility Feedback, Absolute Case Utility Feedback and Utility Feedback.
- Learning Techniques: Probabilistic Similarity Models, Local Similarity Measures and Feature Weights.

### 3 MKL FOR ONTOLOGY MATCHING

We want to find an appropriate combination of similarity measures for an instance matching task. Specifically, our interests lie in learning a linear combination of N similarity measures with nonnegative coefficients  $\beta_j$  that minimizes some error criteria e within a given dataset  $\Psi$ . The elements of such dataset are equivalent and non equivalent correspondences  $C \in (I_1, I_2)$ , where  $I_1$  and  $I_2$  are two homogeneous instance sets:

$$\min e(\sum \beta_j S_j(c), \Psi)$$
(1)  
$$\beta \ge 0$$
  
$$S_i \in (f_i, p_i, m_i), i \in 1..N$$

We see each similarity measure as a 3-tuple  $(f_i, p_i, m_i)$ :  $f_i$  is the actual similarity function,  $p_i$  a specific value set of parameters for the function and  $m_i$  a possible mapping between the properties of the instances. We note that according to this description the same similarity function can be part of two different similarity measures.

The advantage of incorporating the mapping between properties as an additional component of the similarity measure is that this allows us to conduct the instance matching process even though there is neither an unique property mapping nor certainty concerning the correct mapping at the end of the schema matching problem. In this case, it is sufficient to view the mappings as an additional variable during the selection of the similarity measures and allow the algorithm to select the mappings that provide better information to accomplish the task.

Our interest extends only to finding equivalence correspondences among homogeneous instance sets in which their elements belong to equivalence classes. For this reason, to find all the correspondences between all the instances of two ontologies, it would be necessary to carry on a schema matching process and then to transverse the instance tree of the two ontologies in post-order.

This condition leads us to suppose that the underlying similarity rules are globally shared by the individuals of the set. To see that, consider a set in which all its instances have a natural key but they can be members of the concept PERSON or the concept CAR. In this case, the natural key for each class will be obviously different as is the correct combination of similarity measures. While a person should be identified by its social security number, a car should be identified by its license plate number.

We argue that the problem in equation (1) can be solved using the MKL problem. In the following section we present the advantages of such algorithm and describe how it can be used in the instance matching context assuming all similarity measures are also a kernel. Then, we explain how it is possible to learn a kernel from a similarity measure so that the algorithm can be correctly employed.

### 3.1 MKL as Similarity Measure Aggregator

If we limit ourselves to kernel functions (Scholkopf and Smola, 2001) as similarity measures, define the set of candidate correspondences as the input space, label 1 for equivalent correspondences and -1 for non equivalent correspondences, the problem stated in equation (1) is equivalent to the MKL problem (Bach et al., 2004). Under this setting, (Bach et al., 2004) showed that this problem can be solved by the QCP problem of equation (2) whose basic idea is to train a classifier that minimizes the error in the dataset while also learns the optimal coefficients as part of the optimization problem.

$$\min_{\xi,\alpha} \xi - 2\mathbf{1}^T \alpha$$

$$subject 0 \le \alpha \le C, \alpha^T y = 0$$

$$\alpha^T D(y) S_j D(y) \alpha \le tr \frac{S_j}{c} \xi$$

$$\xi \in \mathfrak{R}, \alpha \in \mathfrak{R}^n$$
(2)

where D(y) is the diagonal matrix with diagonal y - the labels -,  $1 \in \mathbb{R}^n$ , the unit vector, and C a positive constant. The coefficients  $\beta j$  are recovered as Lagrange multipliers for the constraints  $\alpha^T D(y)S_jD(y)\alpha \leq tr\frac{S_j}{2}\xi$ .

There are several advantages of using MKL within the context of ontology matching:

First, MKL allows us to find a sparse and nonsparse combination of similarity measures by using various combinations of 1-norms and 2-norms. Primarily 1-norms algorithms form a sparse linear combination that can be useful in parameter selection where few kernels - the ones with the correct parametrization - encode most of the relevant information. On the other hand, 2-norm algorithms find a non-sparse combination that can be useful when features encode orthogonal characterizations of a problem (Marius Kloft and Sonnenburg, 2008); in other words, this may be used to combine complementary similarity measures such as Knowledge, String, and Structural based measures.

Second, there are very efficient methods to solve large scale MKL problems with a large number of kernels ((Rakotomamonjy et al., 2008), (Sonnenburg et al., 2006)). In fact, experimental results show that the available methods work for hundreds of thousands of examples or hundreds of kernels to be combined and that have been applied in demanding applications such as medical data fusion (Yu et al., ). This is extremely useful within the present context in which large and complex ontologies have started to be a concern.

Third, MKL directly addresses the problem of combining similarity measures by using such combination during the learning process. This is contrary to what happens when a neural network or any other classic machine learning algorithm uses similarity measures as features. Feature comparison and not instance comparison is being carried out under this condition, thus, the problem of instance matching is not being directly addressed.

Fourth, a sparse and linear combination of similarities such as the one producing MKL is simple and easy to interpret. If a given situation is observed, all a human has to do is analyze the larger and non zero terms to understand which similarities are important to classify a pair of instances, as different or equivalent.

Finally, it is worthwhile mentioning that MKL is an extension of the SVM algorithm that is capable of learning from small training sets of high-dimensional data with satisfactory precision (Wang et al., 2006).

#### **3.2 Indefinite Kernels**

Even though the configuration needed to use MKL as a solver for (1) is simple and set forth a typical scenario of binary classification, until now kernels have been considered as similarity measures. However, most of the current similarity measures for ontology matching are not explicitly presented as a Kernel. Furthermore, for most of the similarity functions the question on whether or not these are kernels has not even been raised.

On the one hand, kernels are very convenient functions from an optimization point of view. The PSD condition on the Gram matrix makes most of related optimization problems convex, and as a result, low cost computation algorithms for solving them such as interior-point methods (Boyd and Vandenberghe, 2004) - become available. The related problems would be nonlinear without this condition, and under the current technology, intractable.

Moreover, kernels have generalization advantages over regular similarity functions. The latter imply a significant deterioration in the learning guarantee. (Srebro, 2008) found that if an input distribution can be separated, in the sense of a kernel, with a margin  $\gamma$ and an error rate  $\varepsilon_0$ , then for any  $\varepsilon_1 > 0$ , this may also be separated by the kernel mapping viewed as a similarity measure, with similarity-based margin  $\varepsilon_0$  and error rate  $\varepsilon_0 + \varepsilon_1$ . Because  $\varepsilon_0$  and  $\varepsilon_1$  do not take negative values, a kernel-based margin is never smaller than a similarity-based margin.

On the other hand, kernels also come with compromise and trade-offs. Their mathematical expressions do not necessarily correspond to the intuition of a good kernel as a good similarity measure and the underlying margin in the implicit space is not usually apparent in natural representations of the data (Balcan and Blum, 2006). Therefore, it may be difficult for a domain expert to use the theory to design an appropriate kernel for the learning task at hand. Furthermore, the requirement of positive semi-definiteness may rule out most of the natural pairwise similarity functions for the given problem domain.

To use MKL in the context of ontology matching without losing the designability and interpretability of similarity functions, we suggest following the approaches that focus on finding a surrogate kernel matrix *K* derived from the original similarity matrix *S* (Wu et al., 2005).

To this regard, one of the first approaches was to consider all the negative eigenvalues as noise and apply the linear transformation (3) to the similarity matrix to replace all the eigenvalues with zero.

$$A_{clip} = U^T a_{clip} U \tag{3}$$

where  $a_{clip} = diag(I_{\lambda_1>0}, ..., .., .., I_{\lambda_N>0}).$ 

Another common approach consists in changing the signs of all negative eigenvalues - instead of making them zero - by using the following linear transformation:

$$A_{flip} = U^T a_{flip} U \tag{4}$$

where  $a_{flip} = diag(sign_{\lambda_1>0}, ..., ..., sign_{\lambda_N>0})$ .

What we propose is to use the optimization problem stated in (Chen et al., 2009) to alter the original similarity measure. This approach guarantees a consistent treatment of all the samples because the same linear transformation that is applied to the original similarity matrix i.e. the one that creates the original measure, can be applied to the new samples. Besides, by controlling a parameter  $\gamma$  the user can control how far to extend the search for the surrogate matrix.

The problem is presented in equation (5). Given a similarity matrix  $S_m$  calculated from a similarity measure S and whose eigendescomposition is  $U\Lambda U^T$ , this problem finds a linear transformation  $A = Udiag(a)U^T$  that modifies the original similarity matrix by solving a small QCP problem that can be handled by standard optimization packages:

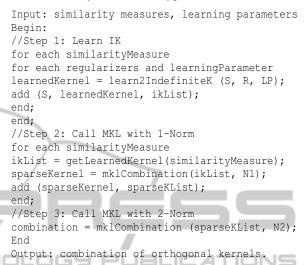
$$\min_{\substack{c,b,\xi,a}} \frac{1}{n} \mathbf{1}^{T} \xi + \eta c^{T} K_{a} c + \gamma h(a)$$
(5)  
ubject to  $diag(y)(K_{a} c + b\mathbf{1}) \ge 1 - \xi$   
 $\xi \ge 0, \Lambda a \ge 0$ 

where h(a) is a convex function that regularizes the search of the modified similarity matrix toward  $S_m$ ; for example, one can use  $h(a) = ||a - a_{clip}||$  to focus the search at the vicinity of the  $A_{clip}$  transformation. Since there may be different regularizers, we suggest employing several of these to find surrogate matrices and allow MKL to select the proper one. In other words, the regularization function and the parameter coefficient may be seen as other components of the similarity measure.

### **3.3 Putting the Ideas Together...**

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The following algorithm shows the suggested sequence to compose MKL and IK. There are three steps in the process. The first one calculates different transformations of similarity measures that use the same similarity function. The second one uses MKL with 1-norm to find a sparse combination of kernels for each similarity function. The last one calls 2-norm MKL to find a linear combination of similarity measures that analyzes different types of features.



#### 3.4 Labeling and Unbalanced Classes

Because MKL is a supervised learning algorithm, an oracle that labels a small set of candidate correspondences as positive is required. The negative labels can be constructed by crossing an instance of a positive correspondence with a random instance that is not within the set given by the oracle. Depending on the real application, the oracle can be a human or some other system that does not require labels to accomplish the alignment. In this case, what could be of value of our approach is the generalization capability of a supervised learning paradigm.

On the other hand, the unbalanced nature of our input space needs to be considered. An element of our space is a candidate correspondence  $(I_i, I_j)$ , where  $I_i, I_j$  are instances of the sets to be matched. Consequently, the cardinality of the input space is  $N \times M$ , where N and M correspond to the size of each of the sets to be mapped. Within this setting, there will be at most max(N, M) positive correspondences making the rest negative.

This is a typical scenario of unbalanced classes that can be treated with Cost-Sensitive or Sampling Techniques. For example, it is possible to choose the undersampling method which changes the training sets by sampling a smaller majority training set (Drummond and Holte, 2003). As the performance of every unbalance technique is highly dependent on the data set (McCarthy et al., 2005), we suggest selecting the technique by using cross validation.

### 4 EXPERIMENTS

A prototype in Java as a proof of concept was implemented. Its architecture is depicted by figure 1.

We used the MKL implementation of Shogun (Sonnenburg and Raetsch, 2010), and Mosek (Mos, 2010) to solve the QCP problem of equation (5). The employed libraries of similarity measures were (Chapman, 2009) for String measures and the Java WordNet Similarity library for Knowledge based measures. Jena OWL was the API to read the OWL ontologies and the alignment files. We also incorporated a TSVM classifier as a final matcher (module 8 in figure 1) whose training algorithm was implemented by (Joachims, 2002).

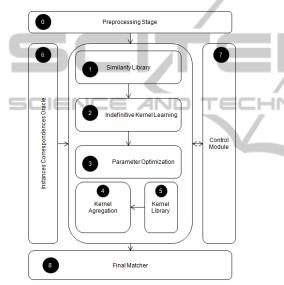


Figure 1: Architecture of the Prototype.

#### 4.1 Kernels and Similarity Measures

A composite kernel to compare two correspondences  $C_i, C_i$  was used:

$$K(C_i, C_j) = K_{internal}(C_i)K_{internal}(C_j).$$
 (6)

where  $K_{internal}$  refers to a kernel that measures the similarities between the instances  $I_1, I_2$  of each candidate correspondence. Considering that the product of two numbers is greater when they are close to each other, this kernel takes greater values when the two correspondences share a similar estimation of similarity.

Two type of measures were employed as internal kernels  $K_{internal}$ :

The first one was the function  $K_{internal}(I_1, I_2) = P_{m_{i,j}}(I_1, I_2)$  where  $m_{i,j}$  is a specific alignment between two properties lists of each ontology given by the

mapping  $m_i$  and P is a local similarity function that compares how similar the values of the two properties of the two instances are. The idea behind this kernel is to follow a natural key approach where the identity of the instances is captured within the value of a few properties.

The second internal kernel was an adapted version of the tree-like function described in (Xue et al., 2009) that is not stated as a Kernel. This measure aims to find structural and semantic similarity. Its basic principle is to find how far apart two instance trees are by computing the operations needed to transform one tree into another. Since both functions require to measure similarity between the values of the properties, the following list of local similarity functions was used:

- String based: BlockDistance, ChapmanLength-Deviation, CosineSimilarity, DiceSimilarity, EuclideanDistance, JaccardSimilarity, JaroWinkler, Levenshtein, MatchingCoefficient, MongeElkan, SmithWatermanGotoh.
- Knowledge based: Lin, Resnik, Path, WuAnd-Palmer.

Clearly, the fact that these similarity measures were employed at the local level makes most of our kernels indefinite.

#### 4.2 Test Set and Results

We used the IIMB bechmark (Ferrara et al., 2008) as basis for our preliminary experiments. The IIMB is an evaluation dataset for the OAEI conference track which consists of several transformations to a reference ontology. This ontology contains 5 named classes, 4 object properties, 13 datatype properties and 302 individuals. We make clear that we have not participated in the official campaign.

There are a total of 37 matching tasks in the benchmark. Each one introduces a class of modifications over the original value/s of a specific property within the source ontology. For example, there are typographical error simulations, changes in the aggregation level, and instantiation on different subclasses of the same individual. Only the first 19 matching tasks were tested because similarity measures designed to capture logic heterogeneity were not employed.

The standard set of parameters for Ontology Matching was used as evaluation measures:

- Precision: the number of correct retrieved mappings / the number of retrieved mappings.
- Recall. the number of correct retrieved mappings / the number of expected mappings.

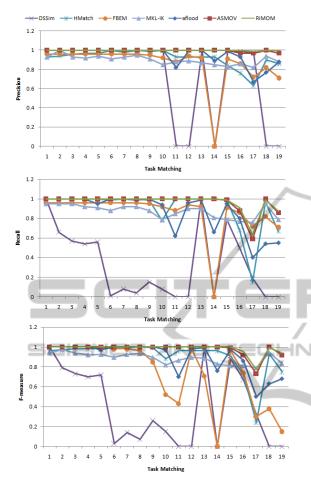


Figure 2: Protoype Behavior vs Other Systems.

• F-measure. 2 x (precision x recall) / (precision + recall).

The following figures reveal the results of the prototype (MKL-IK) and the systems that participated in the OAEI 2009 (Euzenat et al., 2009).

Our prototype was comparable to the other tools in the selected matching tasks. Besides, although most of the time it was overcome by another system, it showed a consistent behavior across every data set. Both facts leads us to believe that the suggested approach should be further explored.

#### **CONCLUSIONS AND FUTURE** 5 WORK

In the present paper, we proposed to use Multiple Kernel Learning (MKL) to combine similarity measures within the context of Ontology Instance Matching. We described the advantages of MKL and explained how it can be used to address the problem. The need to find surrogate similarity matrices to be able to use such algorithm within this context, has been explained and a possible approach to accomplish the task submitted. This approach consists in computing a linear transformation that searches for the surrogate matrix within the vicinity of the original similarity matrix.

We also suggested an algorithm that makes use of both concepts and pointed how the unbalanced class issues of the suggested configuration can be faced. In addition, we implemented a proof of concept prototype and partially tested it using the IIMB benchmark. The results suggest that our approach is feasible and should be explored to extent current matching solutions or to create new ones.

Our current research follows different directions. We are particularly studying the internal behavior of the algorithm and conducting a profound assessment of the prototype through the use of other bechmarks and test dataset. Furthermore, we are analyzing possible performance issues that may appear with this approach.

PUBLICATIONS

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