

A Constituency Parsing Tree based Method for Relation Extraction from Abstracts of Scholarly Publications

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Abstract

We present a simple, rule-based method for extracting entity networks from the abstracts of scientific literature. By taking advantage of selected syntactic features of constituent parsing trees, our method automatically extracts and constructs graphs in which nodes represent text-based entities (in this case, noun phrases) and their relationships (in this case, verb phrases or preposition phrases). We use two benchmark datasets for evaluation and compare with previously presented results for these data. Our evaluation results show that the proposed method leads to accuracy rates that are comparable to or exceed the results achieved with state-of-the-art, learning-based methods in several cases.

1 Introduction

As a public and formal record of original contributions to knowledge, scientific literature is a critical resource that promotes the progress of science and technology in society. To help researchers to comprehend the large and growing amount of scientific literature, automated methods can be used to extract and organize information from corpora of publications, e.g., in terms of scientific key concepts and their relationships. Leveraging prior work that has achieved high accuracy for entity recognition (Lample et al., 2016; Habibi et al., 2017), in this paper, we focus on identifying extracting relationships between entities.

Overall, prior studies consider the task of relation extraction from two perspectives: 1) identifying if a relationship exists between a given pair of identified entities (Gábor et al., 2018), which is also the goal with this paper, and 2) further labeling or classifying the identified relationships (Luan et al., 2018a; Mintz et al., 2009).

Prior work has used different methods for relation extraction. Rule-based algorithms primarily rely on lexical patterns, such as word co-occurrences (Jenssen et al., 2001) and dependency templates (Fundel et al., 2006; Kilicoglu and Bergler, 2009; Romano et al., 2006). Supervised learning-based methods mainly use either feature engineering (Kambhatla, 2004; Chan and Roth, 2011) or kernel functions (Culotta and Sorensen, 2004; Zelenko et al., 2003; Bunescu and Mooney, 2005). Using supervised learning to detect the existence (and type) of relationships between concepts in scholarly publications may further require domain expertise for annotation. Accounting for the fact that humans tend to use their background knowledge to identify relations, Chan and Roth (2010) showed that using external knowledge, such as Wikipedia, improves the accuracy of relation extraction. Recently, deep learning models have also been used for supervised learning (Luan et al., 2018a, 2019). For example, Luan et al. (2019) developed a dynamic span graph framework based on sentence-level BiLSTM for multi-task information extraction. Since data annotation by humans is expensive, semi-supervised learning methods, such as the snowball system (Agichtein and Gravano, 2000), as well as unsupervised learning, such as clustering methods (Daelemans and Van den Bosch, 2005; Davidov and Rappoport, 2008), have also been explored in prior studies. Though remarkable contributions have been made, one remaining limitation with prior machine-learning based work is that these approaches may involve time and related costs for parameter optimization and labeling. Moreover, data-driven training methods may be limited by the domain specificity of learned models. Finally, algorithms trained on deep learning models lack interpretability.

To address the above-mentioned limitations, we

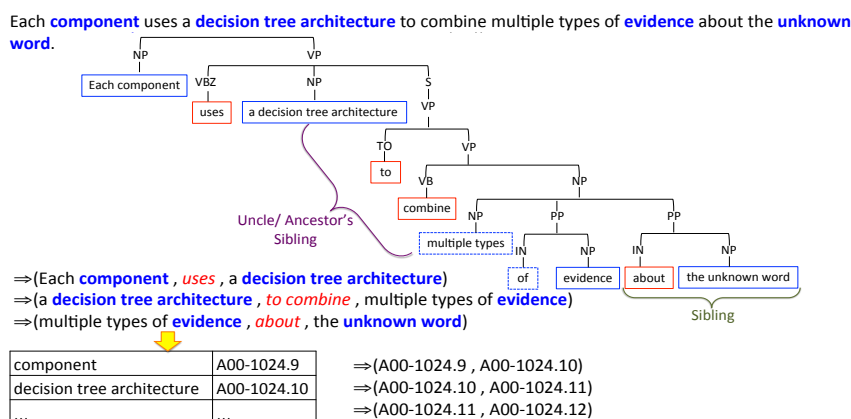


Figure 1: An illustrative example of building a constituency-based concept network (CTN) for a sentence.

propose a rule-based concept network construction method. In the resulting networks, nodes represent annotated entities per document, and edges are constructed based on constituency parsing. Our approach is motivated by the observation that scientific literature typically use formal language with clear syntactic structure and comparatively fixed word order (e.g., term phrases). Inspired by prior work that defines a relationship as either an interaction or an association between two entities (Jurafsky, 2000), we take advantage of the structured information provided by constituency parse trees built for each sentence. More specifically, We capture two types of tuples: 1) (noun phrase, verb-based connection, noun phrase), and 2) (noun phrase, preposition-based connection, noun phrase). To avoid over-fitting on the given domain, our rules are generated on the basis of a context-free grammar.

We evaluate our method against two benchmark datasets that have been previously labeled for entities and relations. Our experimental results show that the proposed constituency-based concept networks achieve comparable accuracy to or can even outperform state-of-the-art, learning-based methods for identifying entity networks. We find that relationships of the type *used-for* and *part-of* are better captured by our approach than other types of relationships. Finally, we describe differences between domain-level concept networks.

2 Method

The construction of constituency-based concept network (CTN) has three stages: 1) data pre-processing, 2) the identification of nodes (we used entities given in annotated data), and of edges based on a constituency parsing tree, and 3) map-

ping entities from the constituency parsing tree to labeled entities. Figure 1 provides an illustrative example of the process of building a CTN for a sentence.

Data Pre-processing In this stage, we focus on two aspects. First, we segment each document into a set of sentences that are the input to constituency parsing. Since scientific texts may have some long sentences with complex sentence structures, we further segment sentences by using regular expressions. Second, we identify the annotated entities per sentence by extracting the labeled entity id and corresponding entity phrase, as well as the entities’ index in the sentence.

Node and Edge Identification To generate a parsing tree for each sentence, we use the AllenNLP constituency parsing (Joshi et al., 2018) As shown in Figure 1, we extract noun phrases at the lowest layer (i.e., children are noun-based end-nodes) as candidates of CTN nodes. We made this decision to capture unique and specific (to the sub-field of science) noun phrases from scientific literature. For linking nodes, we capture keywords that occur between two adjacent node candidates in the parse tree. According to the parse tree structure, these keywords usually occur in two types of positions: 1) the sibling of a CTN node candidate, and 2) the uncle or the ancestor’s sibling of a potential CTN node. Based on node candidates and connecting keywords, we generate an edge if two node candidates are linked by a connecting keyword. One specific issue in this process is the of-phrase: according to our empirical observation, we believe that of-phrases (e.g., “computational model of discourse”) often represent a single concept. Based on this assumption, we merge

	SCIERC	SemEval18
#entities	8089	7482
#relations	4716	1595
#relations/doc	9.4	3.2
cross-sentence relations	yes	no

Table 1: Data statistics.

edge candidates connected by the keyword “of” as a single CTN node, and remove the original edges.

Entity Mapping We remove nodes that had been identified by the constituency parsing tree process described above, but are not labeled as entities in the ground truth data. As shown in Figure 1, the final output of CTN is a set of nodes with ids, and edges.

3 Experiments

3.1 Experimental Setup

Data We perform experiments on two publicly available datasets where humans annotated scientific entities and relations from abstracts of scientific publication. Table 1 provides a brief summary of both datasets. *SemEval18 Dataset* has 500 abstracts prepared by Gábor et al. (2018) for the shared task 7 (subtask 2) of SemEval 2018. All abstracts in this dataset are from published papers in the field of computational linguistics. The annotated relations are divided into six types of semantic relationships between scientific concepts. The *SCIERC Dataset* was provided by Luan et al. (2018a). This dataset has 500 scientific abstracts from 12 AI conference/workshop proceedings that cover five research areas: 1) artificial intelligence (AI), 2) natural language processing (NLP), 3) speech, 4) machine learning (ML), and 5) computer vision (CV).

Baselines For the SemEval data, we compared the results from our network construction method (CTN) with the official baseline, which was generated by using a memory-based k-nearest neighbor (k-nn) search (Gábor et al., 2018). The top three reported submissions in the SemEval leaderboard were: UWNLP (Luan et al., 2018b), ETH-DS3Labl (Rotsztein et al., 2018), and SIRIUS-LTG-UiO (Nooralahzadeh et al., 2018). For SCIERC, we compared our method with two state-of-the-art (SOTA) systems: SciIE (Luan et al., 2018a) and DyGIE (Luan et al., 2019). The original outputs from SciIE and DyGIE also identified

	Dev			Test		
	P	R	F1	P	R	F1
SciIE	62.0	47.7	53.9	66.4	46.7	54.9
DyGIE	55.0	48.6	51.6	63.3	52.2	57.2
CTN (ours)	73.4	47.3	57.5	75.4	46.5	57.5

Table 2: Comparison with previous methods for relation extraction on SCIERC dataset.

and categorized the type and direction of relations, and the boundary of entities. Since we do not identify these elements, we also do not consider them for comparison.

3.2 Results

Extraction Performance Table 2 and Table 3 compare our concept network CTN with baselines for SCIERC and SemEval18, respectively. Compared to SOTA on SCIERC, our approach outperformed the best previously reported results on both the development and testing data. Specifically, CTN achieved a noticeable improvement in precision (we achieved $\sim 74\%$) compared to prior methods, which benefits our F1 value. Further comparing each method’s performance on the development data versus testing data, we observe that our rule-based CTN produces more stable or consistent results than the considered, prior, learning-based methods.

For SemEval18, we find that CTN outperformed the baseline by $\sim 15\%$ (our F1 score was 41.6%), and the third best reported prior result, but could not come close to the top two prior results. From the presented results, we conclude that our rule-based concept network approach can serve as a strong baseline method for identifying related entity pairs in scientific texts.

Ablation Study In order to explore the isolated contribution of each rule considered for adding edges in the CTN, we conduct an ablation analysis

	P	R	F1
UWNLP	-	-	50.0
ETH-DS3Lab	-	-	48.8
SIRIUS-LTG-UiO	-	-	37.4
SemEval Baseline	-	-	26.8
CTN (ours)	33.6	54.8	41.6

Table 3: Comparison with previous methods for relation extraction on SemEval18 dataset.

We find that the most central (in terms of degree) nodes from the CV corpus mainly represent general scientific concepts, such as “method”, “algorithm” and “approach”, while in the ML corpus, key nodes represent domain-specific terms such as “robust PCA” and “side information”. Comparatively, the size of the CTN from the CV domain is larger than that from the ML domain, which might be due to different numbers of abstracts in each domain.

4 Conclusions

In this paper, we have proposed and evaluated a rule-based network construction method that leverages constituency parsing to extract relations between entities in scientific texts. Our method does not require machine learning or domain knowledge. Experiments on two benchmark datasets show that the proposed CTN achieve comparable performance with state-of-the-art learning-based methods in multiple cases. Even though our method could not outperform the two best performing systems built for one of the considered datasets, our results suggest that the demonstrated approach can work as a baseline method for relation extraction. In addition, we find that entities with a relationship of *used-for* and *part-of* are more likely to be connected in our network. Based on the corpus-level CTN, we further saw that key nodes in the networks based on the CV corpus are mainly general scientific terms, while for the abstracts from the ML domain, key nodes represent domain-specific terms. To improve the construction of CTN in the future, we plan to consider cross-sentence link formation and link label detection. Finally, the rules used to build CTN can further support the development of learning-based algorithms for relation extraction.

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