

**Term Project**  
**CS 710 - Artificial Intelligence**

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**Coreference Resolution in Biomedical Text:  
Improving Performance with Ontologies and Domain Specific  
NER**

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# 1 Introduction

For my final project, I built a biomedical coreference resolution system using python open source libraries. Coreference Resolution is hard and important problem as there are many applications that could benefit or have benefited from coreference resolution. Some of those applications are: question answering, text summarization, information extraction, and machine translation.

## 2 Recent Work

### 2.1 Coreference

Coreference resolution is a classical natural language processing problem where the task is to identify which mentions in a text refer to the same real-world entity.

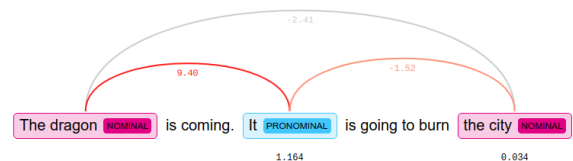


Figure 1: Example of resolving referring expression *it* to antecedent dragon

To better isolate specific sub-tasks, it is helpful to think of the reference the generic algorithm for anaphora and coreference resolution proposed by Ng et al., 2002. Algorithms and techniques described later in the paper will focus on certain sub-tasks to improve the performance of cluster detection.

1. **Identification of referring expressions** Identify all noun phrases in the text.
2. **Characterization of referring expressions** Composed of two sub-tasks: first, define a representation of a discourse entity. This representation determines the knowledge regarding a discourse entity that the algorithm needs in order to perform the remaining steps. Second, compute the information specified in the representation for each entity.
3. **Anaphoricity determination** Determine whether a discourse entity is anaphoric or not. Non-anaphoric entities, do not possess an antecedent and that algorithm will not search for these entities.
4. **Generation of antecedent candidates** Once a NP is determined to be anaphoric, the algorithm identifies scope of the NP and gathers those NPs that falling within its scope to generate a list of candidate antecedents for it.
5. **Filtering** Remove unreasonable candidate antecedents for each anaphoric noun phrase. Can be based off of a set of rules or hard constraints. This step reduces the amount of processing that needs to be performed by the algorithm.
6. **Scoring/ Ranking** Score that indicates the likelihood that two NPs co-specify.
7. **Searching/ Clustering** Select an antecedent for a given anaphor from the list of candidate antecedents returned by the previous step. If this list is empty then no antecedent will be selected for this anaphor.

## 2.2 Neural Network Systems

Since this task requires knowledge at all levels of language processing, including lexical, syntactic, semantic, and world knowledge, success in this problem is indicative of advancement of language technologies. Appropriately, due to advancements in computational power and reinforcement learning, coreference resolution has experienced a revival of sorts with many performant systems proposed in recent times. More specifically, Neural Network based coreference systems employing entity-level information and reinforcement learning have produced new scores outperforming the previous state-of-the-art. An example of such a system is the Deep Reinforcement Learning model proposed by Clark and Manning, 2016.

Specifically, this system is a deep neural network that learns distributed representations of coreference clusters. To gain better intuition of this system, it is better to think of this network comprising three sub-networks: mention-pair encoder, cluster-pair encoder, cluster-ranking model.

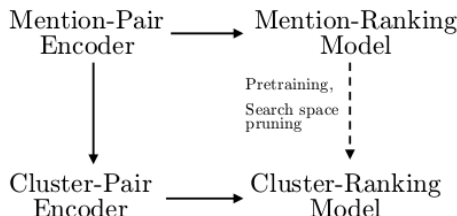


Figure 2: System Architecture defined in Clark and Mannings 2016 paper.

## 2.3 Open-Source Implementation

Though the python source code from *deepcoref* is publically available (Clark and Manning, 2016), it is reliant on StanfordCoreNLP components implemented in Java and is not optimized for speed<sup>1</sup>. For this project, I used the *neuralcoref* library - an open-source cython library built by *Hugging Face*. *Neuralcoref* is greatly inspired by Clark/Mannings coreference system. It is SpaCy pipeline extension so it uses SpaCys document annotations (which I will discuss in further detail in just a moment). Flexibility regarding SpaCy models and other pipeline components (EntityLinker, EntityRuler, etc.). *Neuralcoref* ships pretrained on the CoNLL 2012 dataset and tuned vectors (word embeddings) trained on the OntoNotes 5.0 dataset.

## 3 Dataset and Baseline Evaluation

To evaluate the performance of this system, I used the annotated coreference resolution dataset from the 2011 BioNLP shared task. The goal of this shared task is to find anaphoric expressions to proteins. The annotated data includes 2,300 pubmed abstracts with corresponding protein and gene annotations as well as coreference cluster annotations. Given that the *neuralcoref* library evaluates performance based off a file in the tabular CoNLL format, I wrote an evaluation script that closely follows the match guidelines outlined in the shared task<sup>2</sup>. The system was evaluated

<sup>1</sup>The *deepcoref* system takes roughly 7 days to train on the CoNLL dataset on a GTX Titan GPU.

<sup>2</sup>Writing these custom evaluation scripts proved to be quite helpful when investigating why and where the system was performing poorly.

using F1, precision, and recall scores. These metrics are calculated by comparing predicted cluster pairs vs. gold annotated cluster pairs. A true positive is counted when a predicted cluster comprises mentions of which both are either within a minimum span of a gold mention<sup>3</sup> or that a mention subsumes a gold mention of a gold cluster.

	Prec.	Rec.	F1
SpaCy Web Md	15.15%	8.37%	10.72%

## 4 Improvements

So given a baseline performance score and influenced by recent research, I chose to focus on introducing domain-specific named entity recognition (Choi et al., 2014), dependency parsing, word vectors as well as Knowledge Bases or Ontologies to improve performance. Libraries I used to introduce these were: BioWordVec 200 dimension word vectors trained using Fasttext algorithm on large pubmed corpus And for Domain-specific NER and dependency parsing, SciSpaCy which provides full SpaCy pipelines trained on biomedical text. Regarding word embeddings: NeuralCoref uses static and tuned 50 dimensional word vectors that are used when calculating both mention embeddings as well as pair embeddings. Given that BioWordVec uses 200 dimensional vectors, I trained a new set of 50 dimensional word vectors on a corpus comprising 1 million Pubmed abstracts and a sampling of random walk sequences from a MeSh graph (Node2Vec).

## 5 Results

Given the high number of False Positives, I chose to improve precision as much as possible to avoid a needle in the haystack problem when using this system in an nlp task.

The most performant combination was the use of the SciSpacy model, and pruning using a named entity recognition module trained on the JNLPBA corpus.

We see that in the third row of the table, very poor performance using word vectors trained on biomedical text. This can likely be attributed to the fact that the system was trained using the 50 dimensional google news vectors, so to see any performance gains using domain specific vectors we would need to retrain the system using these vectors.

	Prec.	Rec.	F1
JNLPBA NER	34.14%	11.17%	16.83%
SciSpacy MD	29.86%	14.29%	19.33%
SciSpacy MD BWV	2.02%	7.27%	3.16%

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<sup>3</sup>Some mentions in the annotated text have a minimum span that *must* be included in the mention span. These minimum spans are generally genes or proteins.

## References

- [1] Miji Choi, Karin Verspoor, and Justin Zobel. Analysis of coreference relations in the biomedical literature. In *Proceedings of the Australasian Language Technology Association Workshop 2014*, pages 134–138, Melbourne, Australia, November 2014.
- [2] Kevin Clark and Christopher D. Manning. Improving coreference resolution by learning entity-level distributed representations. In *Proceedings of the 54th Annual Meeting of the Association for Computational Linguistics (Volume 1: Long Papers)*, pages 643–653, Berlin, Germany, August 2016. Association for Computational Linguistics.
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