

# **Application note**

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# Improving spaCy dependency annotation and PoS tagging web service using independent NER services

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Dependency parsing is often used as a component in many text analysis pipelines. However, performance, especially in specialized domains, suffers from the presence of complex terminology. Our hypothesis is that including named entity annotations can improve the speed and quality of dependency parses. As part of BLAH5, we built a web service delivering improved dependency parses by taking into account named entity annotations obtained by third party services. Our evaluation shows improved results and better speed.

Keywords: dependency parsing, named entity recognition, natural language processing Availability: The code for our web service is publicly available (https://github.com/OntoGene/ blah5).

### Introduction

Dependency parsing might be used as a component for tackling many text mining problems. It is often used as a feature to train machine learning algorithms or used for rule-based approaches for relation extraction [1,2]; and can be used to train better word embeddings [3]. However, as continued research and community efforts in the form of the CoNLL shared tasks, for example, show, dependency parsing in general is still not a problem solved completely [4]. Making a fast, reliable dependency parser readily available for the wider research community for further processing to build upon will help spur efforts in event and relation extraction.

As previous research has shown, providing named entity information to the dependency parser can improve the accuracy of the parses [5,6]. The reasoning is that dedicated named entity recognition (NER) tools perform much better in their specific domain, and by extracting named entities with higher accuracy will facilitate appropriate parsing tools that are not trained on biomedical data.

spaCy is a open-source natural language processing (NLP) library written in Python that performs tokenization, Part-of-Speech (PoS) tagging and dependency parsing. It is the fast-est NLP parser available, and offers state-of-the-art accuracy [2,7].

Services such as PubDictionaries and OGER perform dictionary-based entity look up [8]. Other state-of-the-art taggers are the Jensen tagger [9] and TaggerOne [10].

In this work, we present our contribution to BLAH5. We proposed to build a public web service that can be used by researchers and other services, such as PubAnnotation, to obtain improved dependency annotations, based upon integration with a NER service (Fig. 1). We report on its efficacy here.

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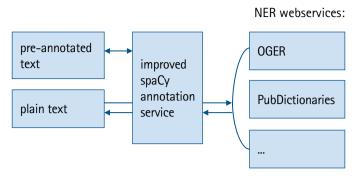


Fig. 1. Architecture of our web service.

# **Related Work**

The most recent extensive evaluation of existing dependency parsers has been performed by Choi et al. [11]. They evaluate 10 different off-the-shelf parsers for accuracy and speed; reporting labeled attachment scores (LAS) of 85% to 90%. While spaCy does not perform the most accurate in their evaluation, it performs fastest maintaining comparable accuracy. Similar scores are reported in the 2015 SemEval task [12].

Pletscher-Frankild and Jensen [9] note that in text mining literature, tools are often evaluated for accuracy only, omitting practical considerations such as speed and robustness.

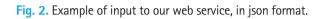
Since the publication of the previously mentioned evaluation of dependency parsers [11], new machine learning-based parsers have been published, surpassing existing approaches according to the respective authors' own evaluation [13,14]. Amongs them, spaCy was updated to employ a neural network to improve performance [7]. As Yang et al. [15] show, however, domain adaptation remains an issue.

# Methods

Our web service is implemented using Flask (http://flask.pocoo. org/), a Python library that facilitates the creating of web interfaces, and the aforementioned spaCy (https://spacy.io). spaCy's structure is modular, and processing text happens by calling the respective modules. For our approach, we added a further processing step between "tokenization" and "dependency parsing," during which we will recompute token segmentation taking into account named entities provided. Then we pass the adjusted token offsets to the dependency parsing module.

#### Input format

Input is provided either as plain text or as a JSON string. The latter



may either contain just a text field, or additionally annotations in PubAnnotation's JSON format (http://www.pubannotation.org/docs/annotation-format/) (Fig. 2).

The input is then internally converted into a spaCy object. From there, spaCy is used to perform tokenization, PoS tagging and dependency parsing, taking into account existing annotations if available.

This is done by tokenizing the text provided, and iterating through the given annotations to form a new sequence of tokens. This new sequence keeps the NER annotations as individual tokens and does not break them down further. In case of conflicting and overlapping annotations, we use the leftmost longest match.

Using this new token sequence, a new spaCy representation object is created on which dependency parsing is performed.

If no annotations are provided manually, our web service will automatically call another service to automatically obtain named entity annotations. While this could be any service, it is currently set to using OGER.

### **Output format**

The resulting object is then converted back into JSON, and token offsets are computed. The output format is PubAnnotation JSON, and can directly be uploaded to PubAnnotation. This also allows to make use of PubAnnotation's annotation visualizer, *TextAE*, as seen in Fig. 3.

#### Usage

The code for our web service is publicly available (https://github. com/OntoGene/blah5), and installation is straight-forward. For demonstration purposes, we are running an instance of the web service.

Improved dependency annotations can be obtained using curl or wget in the fashion shown in Fig. 4.

#### [R0] ROOT

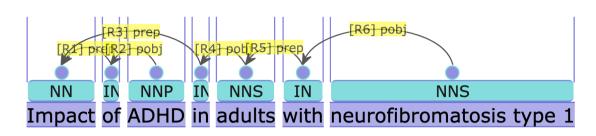


Fig. 3. Visualizing the web service's output in TextAE.

curl -F file='@data/sample.json' https://pub. cl.uzh.ch/projects/ontogene/blah5/ > out.json

**Fig. 4.** Example of complete query to the dependency annotation server: sample.json is the input file, out.json the output file, both in json format.

#### Adaptation

The application can be run locally, provided the necessary Python libraries are available. To add other independent web services to provide NER information, refer to the documentation of the code.

### Results

We evaluated our service in two regards: speed and accuracy.

#### Speed

In order to evaluate speed of our web service, we created a set of 10,000 randomly selected PubMed abstracts, (1) which we sent to the web service directly to obtain "default" dependency parses, and (2) which we first tagged for named entities using OGER, and then sent to the web service to obtain "improved" parses. Table 1 lists the time it took our tool to complete the dependency parsing step for both cases.

We attribute this decrease in processing time to the diminished

**Table 1.** Time required to process a batch of 10,000 PubMed abstracts, in the default case (without prior named entity recognition) and with added named entity recognition

	Default	Improved
Total time (s)	1018	931
Per abstract (s)	0.10	0.09

complexity of computing the parses.

In a practical setting, however, our web service needs to recompute token offsets in order to correctly integrate NERs in the token stream.

While spaCy in the "default" case does this in a internally optimized fashion, our service currently is not optimized to the same degree, which in practice results in overall lower processing times. The integration of NERs still produces an advantage, as the generated parses avoid some errors that otherwise would have been generated (see below). In practice, when using the web service over the network, total processing time will be around 0.3 s per abstract.

#### Accuracy

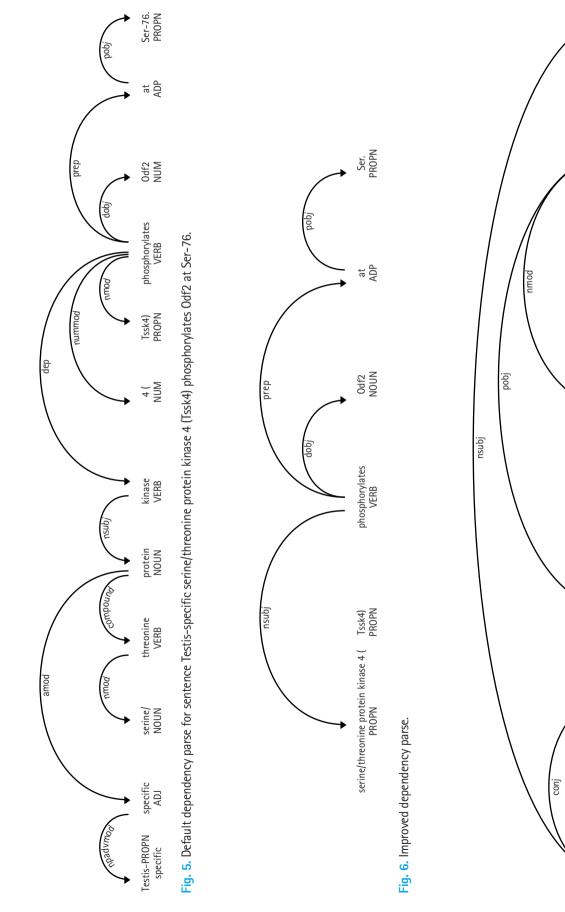
We expect our service to perform well in situations where multiword named entities cause problems for the parser. Below we demonstrate two examples in which our service performs well in such situations.

As Figs. 5 and 6 show, complex named entities are problematic for the dependency parser and tagger, causing it to tag "kinase" as a verb; when in fact it is part of the compound entity "Testis-specific serine/ threonine protein kinase 4 (Tssk4)." This results in an incorrect parse tree.

The second example of Figs. 7 and 8 shows the case where the parse is not technically wrong, but convoluted and useless to downstream tasks.

### Discussion

The examples shown above point to the efficacy of our web service. However, a more thorough evaluation, while useful, is difficult. This is mainly due to the fact that manual creation of parsing annotations is prohibitively expensive, and the fact that existing annotated corpora are annotated in various incompatible formats. For example, depending on the annotation schema, conjunctions can be represented



detected. VERB

SBCADD) NOUN

deficiency ( NOUN

dehydrogenase NOUN

CoA CCONJ

acyl-NOUN

chain NOUN

branched ADJ

short/ ADJ

of ADP

analysis NOUN

mutation NOUN

and CCONJ

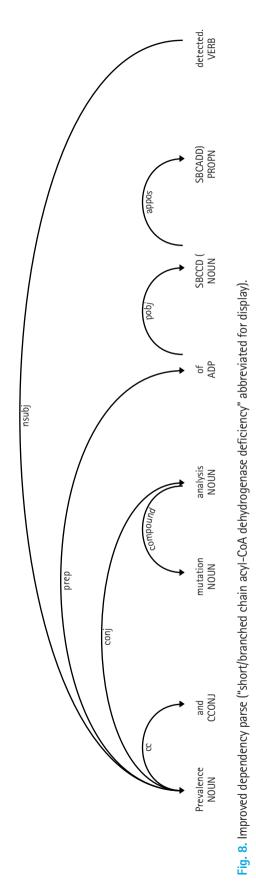
Prevalence NOUN

nmod

Fig. 7. Default dependency parse for the sentence "Prevalence and mutation analysis of short/branched chain acyl-CoA dehydrogenase deficiency (SBCADD) detected."



4/6



in various ways, or relationships may be either undirected or have opposite directionality. Even though there are efforts to homogenize annotation schemes, such as Universal Dependencies [16], they are not yet widely adopted. This makes automatic comparison of results very difficult. However, since dependency parsing is not usually a goal in itself, but a component used for downstream tasks, an extrinsic evaluation could be carried out on such a downstream task such as relation extraction.

### Conclusion

We have presented an improved service for dependency parses, and show that it can deliver better parses for phrases containing complex tokens.

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# **Authors' Contribution**

Conceptualization: NC, FR. Data curation: NC, FR. Formal analysis: NC, FR. Funding acquisition: NC, FR. Methodology: NC, FR. Writing – original draft: NC, FR. Writing – review & editing: NC, FR.

# **Conflicts of Interest**

No potential conflict of interest relevant to this article was reported.

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# References

- Nguyen TH, Grishman R. Relation extraction: perspective from convolutional neural networks. In: 2015 Conference of the North American Chapter of the Association for Computational Linguistics - Human Language Technologies (NAACL HLT 2015), 2015 May 31-Jun 5, Denver, CO, USA. Stroudsburg: Association for Computational Linguistics, 2015. pp. 39-48.
- 2. Colic N. Dependency parsing for relation extraction in biomedical literature [thesis]. Zurich: University of Zurich, 2016.
- 3. Levy O, Goldberg Y. Dependency-based word embeddings. In:

Proceedings of the 52nd Annual Meeting of the Association for Computational Linguistics (Short Papers) (Toutanova K, Wu H, eds.), 2014 Jun 23-25, Baltimore, MD, USA. Stroudsburg: Association for Computational Linguistics, 2014. pp. 302-308.

- 4. Zeman D, Hajic J, Popel M, Potthast M, Straka M, Ginter F, et al. CoNLL 2018 shared task: multilingual parsing from raw text to universal dependencies. In: Proceedings of the CoNLL 2018 Shared Task: Multilingual Parsing from Raw Text to Universal Dependencies (Zeman D, Hajic J, Popel M, Straka M, Nivre J, Ginter F, et al., eds.), 2018 Oct 31-Nov 1, Brussels, Belgium. Stroudsburg: Association for Computational Linguistics, 2018. pp. 1-21.
- 5. Rinaldi F, Dowdall J, Hess M, Kaljurand K, Koitand M, Kahusk N, et al. Terminology as knowledge in answer extraction. In: Proceedings of the 6th International Conference on Terminology and Knowledge Engineering (TKE 2002) (Melby A, ed.), 2002 Aug 28-30, Nancy, France. Le Chesnay: INRIA, 2002. pp. 107-113.
- 6. Lample G, Ballesteros M, Subramanian S, Kawakami K, Dyer C. Neural architectures for named entity recognition. Ithaca: arXiv, Cornell University, 2016. Accessed 2019 Apr 2. Available from: https://arxiv.org/abs/1603.01360.
- Honnibal M, Johnson M. An improved non-monotonic transition system for dependency parsing. In: Proceedings of the 2015 Conference on Empirical Methods in Natural Language Processing, 2015 Sep 17-21, Lisbon, Portugal. Stroudsburg: Association for Computational Linguistics, 2015. pp. 1373-1378.
- **8.** Furrer L, Jancso A, Colic N, Rinaldi F. OGER++: hybrid multitype entity recognition. J Cheminform 2019;11:7.
- **9.** Pletscher-Frankild S, Jensen LJ. Design, implementation, and operation of a rapid, robust named entity recognition web service. J Cheminform 2019;11:19.
- 10. Leaman R, Lu Z. TaggerOne: joint named entity recognition

and normalization with semi-Markov Models. Bioinformatics 2016;32:2839-2846.

- 11. Choi JD, Tetreault J, Stent A. It depends: dependency parser comparison using a web-based evaluation tool. In: Proceedings of the 53rd Annual Meeting of the Association for Computational Linguistics and the 7th International Joint Conference on Natural Language Processing (Zong C, Strube M, eds.), 2015 Jul 26-31, Beijing, China. Stroudsburg: Association for Computational Linguistics, 2015. pp. 387-396.
- Oepen S, Kuhlmann M, Miyao Y, Zeman D, Cinkova S, Flickinger D, et al. SemEval 2015 task 18: broad-coverage semantic dependency parsing. In: Proceedings of the 9th International Workshop on Semantic Evaluation (SemEval 2015) (Nakov P, Zesch T, Cer D, Jurgens D, eds.), 2015 Jun, Denver, CO, USA. Stroudsburg: Association for Computational Linguistics, 2015. pp. 915-926.
- Dozat T, Manning CD. Deep biaffine attention for neural dependency parsing. Ithaca: arXiv, Cornell University, 2016. Accessed 2019 Apr 2. Available from: https://arxiv.org/abs/1611.01734.
- 14. Kiperwasser E, Goldberg Y. Simple and accurate dependency parsing using bidirectional LSTM feature representations. Trans Assoc Comput Linguist 2016;4:313-327.
- 15. Yang H, Zhuang T, Zong C. Domain adaptation for syntactic and semantic dependency parsing using deep belief networks. Trans Assoc Comput Linguist 2015;3:271-282.
- 16. Nivre J, de Marneffe MC, Ginter F, Goldberg Y, Hajic J, Manning CD, et al. Universal dependencies v1: a multilingual treebank collection. In: Proceedings of the Tenth International Conference on Language Resources and Evaluation (LREC 2016) (Calzolari N, Choukri K, Declerck T, Goggi S, Grobelnik M, Maegaard B, et al., eds.), 2016 May 23-28, Portoroz, Slovenia. Paris: European Language Resources Association, 2016. pp. 1659-1666.