

Dep_search: Efficient Search Tool for Large Dependency Parsebanks

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Abstract

We present an updated and improved version of our syntactic analysis query toolkit, \textit{dep\_search}, geared towards morphologically rich languages and large parsebanks. The query language supports complex searches on dependency graphs, including for example boolean logic and nested queries. Improvements we present here include better data indexing, especially better database backend and document metadata support, API access and improved web user interface. All contributions are available under open licences.

1 Introduction

Huge text collections crawled from the Internet have become a popular resource for natural language processing and linguistic studies. Recently, massive corpora with automatically analyzed full syntactic structure became available for 45 languages (Ginter et al., 2017). To efficiently find and access specific types of sentences or syntactic structures from corpora with billions of tokens, powerful search systems are needed for both tree and tree based queries, and combinations thereof.

The SETS dependency tree search tool (Luotolahti et al., 2015) is developed for efficient and scalable tree search in dependency treebanks and parsebanks, supporting complex searches on words, lemmas, detailed morphological analyses and dependency graphs. It is implemented using efficient data indexing and turning search expressions into compiled code.

In this paper we present the \textit{Turku dep\_search} tool, an improved version of SETS. In addition to changing the name, our main contributions are: 1) major speed-up and reduction of stored index size by changing backend from SQLite to Solr\textsuperscript{1} and LMDB\textsuperscript{2}; 2) API access and improved web user interface; 3) supporting arbitrary sentence/document metadata enabling for example sub-corpora and multilingual searches; 4) Universal Dependencies\textsuperscript{3} treebanks and automatically analyzed UD parsebank data for 45 languages publicly available through \textit{dep\_search} API and online interface.

2 Query Language

The query language is in many ways inspired by languages used in existing tree query software such as TGrep (Rohde, 2004). The query language defines criteria for tokens and their relations in dependency graphs to be queried. It allows the user to specify graph structures in the syntactic trees, tokens by their properties, linear order of the tokens, and combinations thereof. Boolean logic is supported and queries can be combined and nested. The query itself is parsed into a tree structure, reflecting the nature of syntactic trees, and it is transformed into executable code with Cython\textsuperscript{4}. The query parser is able to differentiate between token word forms and tags present in the queried corpora.

The query language is best presented by examples. Arguably the most simple query is querying for all tokens. This is achieved by the query \_ underscore representing any token. To restrict the token by a word form or a tag the query is the word form itself, for example, to query the word cat the expression is cat. The lemma cat can be queried with the expression L=cat. Similarly to find tags, a query to find nouns is NOUN. Token queries can easily be combined using boolean operators. For example to search for a token with the lemma cat or dog, that is not in genitive case, one can query: (L=cat | L=dog) & !Case=Gen.

\textsuperscript{1}http://lucene.apache.org(solr/
\textsuperscript{2}http://www.lmdb.tech/doc/
\textsuperscript{3}http://universaldependencies.org/
\textsuperscript{4}http://cython.org
We can add dependency restrictions to our query to search for syntactic structures in the dependency graphs. A simple query like this, looks like cat > _ which finds tokens with word form cat with a dependent token. In a similar vein, we can search for a token with a typed dependent, for example to find the token cat with an amod dependent, query is: cat > amod _. The query for cat with two separate amod dependents query is cat > amod _ > amod _ and to look for chains of amod dependencies from the token cat, one needs to use parenthesis in their query: cat > amod (_ amod > _).

Boolean operators can be used with dependents similarly as they can be used with the tokens. To negate a dependency relation, two options are offered: The query cat ! amod _ looks for token cat without an amod dependent, where as the query cat > amod _ searches for the token cat with a dependent not of the type of amod. Since all parts of the query support boolean logic, subtrees can be negated, for example querying cat > amod ! pretty would find token cat if it has an amod dependent which is anything but the word pretty. Besides using negation, one can use OR operator to query dependency relations. For example, the query cat > amod | amod _ finds cat with amod or nmod dependents.

The third class of query operators has to do with the sentence; linear order of the tokens and set operations on subqueries. To query for tokens next to each other in the sentence, query syntax is: first . second, query to search for tokens in a window is cat < lin:2:3 NOUN, which finds cat - tokens with a noun within two to three tokens from it. The operation can be limited to a particular direction by adding @R/@L - operator. The previous query limited to only tokens to the right is: cat < lin:2:3@R NOUN. The @R/@L -operator can also be applied to all dependency types. The universal quantifier operator (->) allows searching for sentences in which all tokens of certain type have a property. For example query: (. < nsubj _ ) - > (Person=3 < nsubj _ ), finds sentences in which all subjects are in third person. The plus operator allows us to find sentences with multiple properties, eg. to find sentences with both tokens cat and dog, where dog is a subject, query is: (dog < nsubj _) + cat. In addition to these, the size of the result can be set by adding {len_set=limit} after the query. For example: Clitic=Han {len_set=2}.

### 3 Design

The search is executed in two main steps: 1) fetching candidate sentences from the indexed data, so that in a returned candidate sentence, all restrictions must be individually met, and 2) evaluating these candidates to check whether the configuration of the sentence fully matches the query. As the full configuration evaluation (part 2) stays mostly untouched compared to earlier version of the search tool, it is only briefly discussed here, and more detailed information can be found from Luotolahiti et al. (2015).

For fast retrieval of candidate sentences, we use the Solr search engine to return a list of sentence ids, where a sentence has to match all query restrictions individually (for example, sentence has a specific word, morphological tag, and/or relation type), but no relations of these individual restrictions are evaluated at this point. For fast retrieval of candidate sentences, an index is build individually for all possible attributes (e.g. words, lemmas, morphological features and dependency relations). The actual sentence data is stored separately in a fast memory-mapped database, LMDB, where for each sentence the data is already stored in the binary form used when the full sentence configuration is evaluated. The sentences can be fetched from the LMDB using sentence ids given by the search engine.

Together with the different attribute indices, the search engine index can be used to store any necessary metadata related to sentences and documents and further restrict the search also on metadata level. Metadata can be used for example to restrict the search to a specific language, sub corpora or time span, naturally depending on the metadata available for a corpus in use. This way we can keep all data from different corpora and languages in one database, giving us also the possibility to search similar structures across languages.
Especially in the case of Universal Dependencies, cross-linguistically consistent treebank annotation, this gives a great opportunity to study similar structures across languages without compromising on speed or ability to limit the set of interesting languages.

After fetching the candidate sentences, their full configuration is evaluated against the actual search expression. The search expression is turned into a sequence of set operations, which can be compiled into native binary code. This compilation is done only once in the beginning of the search, taking typically less than a second. All sentences passing this step are then returned to the user.

4 Benchmarks

Generally, by changing the database backend we are able to gain in terms of speed and disk space. We are also able to remove two major bottleneck queries.

In the previous version of the search tool, one major bottleneck was queries involving rare lexical restrictions. When the SQLite backend was used, the data index needed to be divided into multiple small databases in order to keep retrieving fast. This however resulted slower queries when rare lexical items were queried because it was needed to iterate through many of these small databases with very few hits in each in order to find enough hits for the user. The new Solr backend is able to hold the whole data in one index, giving us a major speed-up when rare lexical items are searched.

Even bigger speed-up in the new version of dep_search is noticed when the query involves OR statements. Solr handles alternative restrictions much faster than SQLite is able to do, removing the most computationally heavy part of these queries.

In addition to faster queries, dep_search needs now much less disk space for the data index. In the index of 20M trees the disk usage is ~45G, which is about half of the size compared to what the old version was using.

5 Web User Interface and API

In addition to the search tool, we also provide a graphical web interface, which can be set to talk to dep_search API, and render results in browser using Python Flask\(^5\) library, Ajax and BRAT annotation tool (Stenetorp et al., 2012). Dep_search API is accessed through http, and receives the query and the database name, and returns the matching trees as a response.

We maintain a public server for online searches at http://bionlp-www.utu.fi/dep_search/, where we have indexed all 70 Universal Dependencies v2.0 treebanks (Nivre et al., 2017), as well as automatically analyzed parsebank data for all the 45 languages present in UD parsebank collection. For each of these 45 languages, 1M sentences are made available in the dep_search web interface. Additionally, dep_search is used through its API to provide automatic content validation in the Universal Dependencies project,\(^6\) automatically reindexed upon any update of a UD treebank development repository. In the past 6 weeks, the server processed over 11,000 requests — mostly related to the automated UD validation.

6 Conclusions

In this paper we presented the Turku dep_search dependency search tool with expressive query language developed to support queries involving rich morphological annotation and complex graph structures. The search tool is made scalable to parsebanks with billions of words using efficient data indexing to retrieve candidate sentences and generating algorithmic implementation of the actual search expression compiled into native binary code. Dep_search tool supports indexing varying sentence and document metadata making it possible e.g. to focus the search to a specific time span or a set of languages. Source code for the search backend and the web user interface is publicly available at https://github.com/fginter/dep_search and https://github.com/fginter/dep_search_serve, respectively.

Additionally, we provide a public, online search interface, where we host all Universal Dependencies version 2.0 treebanks, together with UD parsebank data for 45 different languages.

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\(^5\)http://flask.pocoo.org/

\(^6\)http://universaldependencies.org/svalidation.html
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