Customized automatic corpus annotations using AlvisNLP/ML

Robert Bossy, Mouhamadou Ba, Claire Nédellec

Mathématiques et Informatique Appliquées du Génome à l'Environnement – *Bibliome* Institut National de la Recherche Agronomique

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Who we are

and what we do

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Part of the National Institute of Agronomical Research

- Domains of interest: agriculture, food, biology, ecology, sustainable development.
- We are a reasearch team in a bioinformatics lab (MaIAGE).
- Our specialties: NLP applied to biology, domain-specific Knowledge Acquisition.

Our approach

- Our research and developments always start with and support applied services for end-users.
- We make developments as generic and reusable as possible.

Software

- AlvisAE: annotation editor.
- TyDI: terminology and ontology editor.
- AlvisIR: semantic search engine framework.
- AlvisNLP/ML: corpus processing engine.

Projects

- We joined the BioNLP Shared Task in 2011 (BB3 and SeeDev tasks).
- We are part of EU project OpenMinTeD:
 - Objective: offer a text-mining infrastructure for researchers.
 - Lessons could be drawn from this Hackathon.

AlvisNLP/ML

what it is and how it works

NLP workflow engine

Allows users to assemble processing modules into NLP pipelines.

Design goals

- Reproducibility of data processing.
- Easy adaptation to new documents/data/problems.
- Allow *sharing* pipelines.
- Scalability.

Library of modules

- I/O: read and write in a variety of formats.
- Linguistic: tokenization, POS-tagging, parsing, term extraction...
- Generic: lexicon projection, regexp...
- Machine-learning: Weka, Wapiti.

The Plan

The plan is the file where the user specifies their pipeline:

- The sequence of modules.
- Extenal resources to be used.
- Parameters for each module.

```
<alvisnlp-plan id="REN">
 <module id="reader" class="XMLReader2">
    <sourcePath filter="\.xml$">../corpus/Ouaero t3.2 gene dev+train-v1.1</>
    <xslTransform>../../bibliome/share/xslt/gene-train2alvisnlp.xslt</>
 </module>
 <import file="../resources/segmig.xml" id="segmig"/>
 <module id="tt" class="TreeTagger">
    <treeTaggerExecutable>/bibdev/install/tree-tagger-3.2/bin/tree-tagger</>
    <parFile>/bibdev/install/tree-tagger-3.2/lib/english.par</>
 </module>
 <module id="train" class="TrainingElementClassifier">
    <algorithm>weka.classifiers.bayes.NaiveBayes</>
    <classifierFile>classifier.model</>
    <relationDefinition>attributes.xml</>
    <examples>documents[@set=="train"].sections.layer:candidates</>
 </module>
</alvienla alons
```

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```

Command-line interface

The Most Useful Interface.

REST interface

- Exposes the documentation.
- Allows to expose selected plans and parameters.
- Users can run plans synchronously or asynchronously.

Our proposal

and the means to achieve it

Initial idea

- Make the REST service understand the PubAnnotation protocol.
- In order to turn any plan exposed by the REST service into a linked annotation tool.

Benefits

- More exposure of our developments.
- We bring a potential of a collection of NLP annotation tools at once.

Materials

Source code

- Java.
- ALv2.
- https://github.com/Bibliome/alvisnlp

Working deployment

- Ready-to-use plan:
 - bacterial taxon named-entity recognition.
- Running the application on my laptop, Jetty Maven plugin, maybe INRA's servers.

Workforce

- There are two of us.
- Main AlvisNLP/ML developer.