

Documentation for

# JULIE Lab UIMA Wrapper for OpenNLP Chunker

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## 1 Objective

JULIE LAB UIMA WRAPPER FOR OPENNLP CHUNKER is part of the Julie Lab NLP tool suite<sup>1</sup> which contains several NLP components (all UIMA compliant) from sentence splitting to named entity recognition and normalization as well as a comprehensive UIMA type system. The OPENNLP CHUNKER<sup>2</sup> provides chunks to tokens in IOB format. For more detailed information on the functioning of the OPENNLP CHUNKER check <http://www.opennlp.org>.

UIMA WRAPPER FOR OPENNLP CHUNKER is currently available in version 2.1

## 2 Requirements and Dependencies

JULIE LAB UIMA WRAPPER FOR OPENNLP CHUNKER is written in Java 1.6 using Apache UIMA version 2.1.0-incubation<sup>3</sup>. It was not tested with other UIMA versions.

The input and output of an AE takes place by annotation objects. The classes corresponding to these objects are part of a *Julie Lab UIMA Type Systems*<sup>4</sup>.

The wrapper comes as a UIMA pear file. Run the Pear-Installer (e.g., `./runPearInstaller.sh` for Linux) from your UIMA-bin directory. After installation, you will find a subfolder desc

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<sup>1</sup><http://www.julielab.de/>

<sup>2</sup><http://www.opennlp.org>

<sup>3</sup><http://incubator.apache.org/uima/>

<sup>4</sup>The *Julie Lab UIMA type systems* can be obtained from <http://www.julielab.de/>

in your installation folder. This directory contains a descriptor `ChunkAnnotator.xml`. You may now e.g. run UIMA's Collection Proccessing Engine Configurator (`cpeGUI.sh`) and add the wrapper as a component into your NLP pipeline.

This pear package also contains models for chunking. The models are trained on two bio-medical corpora respectively: GENIA ([OTK02]) and PennBioIE<sup>5</sup>. An accuracy of 93.6% is yielded on the GENIA using 10-fold cross-validation [BWPH06]. You will find the models in the directory `resources`.

### 3 Using the AE – Descriptor Configuration

In UIMA, each component is configured by a descriptor in XML. In the following we describe how the descriptor required by this AE can be created with *Component Descriptor Editor*, an Eclipse plugin which is part of the UIMA SDK.

A descriptor contains information on different aspects. The following subsection refers to each sub aspect of the descriptor which is, in the Component Descriptor Editor, a separate *tabbed page*. For an indepth description of the respective configuration aspects or tabs, please refer to the *UIMA SDK User's Guide*<sup>6</sup>, especially chapter 12 on "Component Descriptor Editor User's Guide".

To define your descriptor go through each tabbed pages mentioned here, make your respective entries (especially in page *Parameter Settings* you will be able to configure OPENNLP CHUNKER to your needs) and save the descriptor as `DescriptorName.xml`.

**Overview** This tab provides general informtion about the component. For the OpenNLP Chunker you need to provide the information as specified in Table 1.

**Aggregate** Not needed here, as this AE is a primitive.

**Parameters** See Table 2 for a specification of the configuration parameters of this AE. Do not check "Use Parameter Groups" in this tab.

**Parameter Settings** The specific parameter settings are filled in here. For each of the parameters defined in 3, add the respective values here (has to be done at least for each parameter that is defined as mandatory). See Table 3 for the respective parameter settings of this AE.

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<sup>5</sup><http://bioie.ldc.upenn.edu/>

<sup>6</sup><http://incubator.apache.org/uima/>

Subsection	Key	Value
Implementation Details	Implementation Language	JAVA
	Engine Type	Primitive
Runtime Information	updates the CAS	yes
	multiple deployment allowed	yes
	outputs new CASes	no
	Name of the Java class file	de.julielab.jules.ae.opennlp.ChunkAnnotator
Overall Identification Information	Name	JULES-OPENNLP-CHUNK-AE
	Version	2.1
	Vendor	julielab
	Description	see above

Table 1: Overview/General Settings for AE.

Parameter Name	Parameter Type	Mandatory	Multivalued	Description
modelFile	String	yes	no	Path to the OPENNLP CHUNKER model
posTagSetPref	String	no	no	POS (Part-of-Speech) annotations to get from CAS (see <i>julie-morpho-syntax-types.xml</i> )
mappings	String (multi-valued)	yes	yes	Mappings between chunks provided by the OpenNLP Chunker and CAS types (see <i>julie-morpho-syntax-types.xml</i> )

Table 2: Parameters of this AE.

Parameter Name	Parameter Syntax	Example
modelFile	model.bin.gz	resources/ChunkGenia.bin.gz
posTagSetPref	CAS sub-type of <i>POSTag</i>	de.julielab.jules.types.GeniaPOSTag
mappings	OpenNLP Name;CAS Name	NP;de.julielab.jules.types.NP

Table 3: Parameter settings of this AE.

**Type System** On this page, go to *Imported Type* and add the *julie-morpho-syntax-types.xml* type system. (Use “Import by Location”).

**Capabilities** See Table 4.

Type	Input	Output
de.julielab.jules.types.Sentence	✓	
de.julielab.jules.types.Token	✓	
de.julielab.jules.types.POSTag	✓	
de.julielab.jules.types.Chunk		✓

Table 4: Capabilities of this AE.

**Index** Nothing needs to be done here.

**Resources** Nothing needs to be done here.

## 4 Copyright and License

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The license is approved by the Open Source Initiative, and is available from their website at <http://www.opensource.org>.

## References

- [BWPH06] Ekaterina Buyko, Joachim Wermter, Michael Poprat, and Udo Hahn. Automatically adapting an NLP core engine to the biology domain. In Hagit Shatkay, Lynette Hirschman, Alfonso Valencia, and Christian Blaschke, editors, *Proceedings of the Joint BioLINK-Bio-Ontologies Meeting. A Joint Meeting of the ISMB Special Interest Group on Bio-Ontologies and the BioLINK Special Interest Group on Text Data Mining in Association with ISMB*, pages 65–68. Fortaleza, Brazil, August 5, 2006, 2006.
- [OTK02] Tomoko Ohta, Yuka Tateisi, and Jin-Dong Kim. The GENIA corpus: An annotated research abstract corpus in molecular biology domain. In M. Marcus, editor, *HLT 2002 – Human Language Technology Conference. Proceedings of the 2nd International Conference on Human Language Technology Research*, pages 82–86.

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