

Extraction of Drug-Drug Interactions from Biomedical Texts

Carlo Alessi & Doria Saric

May 30, 2018

- 1 Software
- 2 Method
- 3 Learners
- 4 Results
- 5 Conclusions

Software

- Base language: Python
- XML parser: ElementTree
- Pandas
- Numpy
- Gensim: Word2Vec
- NLTK
 - a. Word tokenizer
 - b. Part-of-Speech tagger
 - c. English stemmer
- Scikit Learn
 - a. Data preprocessing
 - b. Pipelines
 - c. SVM
- Keras: ANN

Method

- Task9.1
- Task9.2

Learners

1. Feed-forward Neural Network
 - Kernel: radial basis function
 - $C = 1$
 - $\gamma = \text{auto}$
 - $\alpha = 0.001$
2. Support Vector Machine

Results

Conclusions