



Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models

Yifan Peng¹, Anthony Rios^{1,2}, Ramakanth Kavuluru^{2,3}, Zhiyong Lu¹

¹National Center for Biotechnology Information, National Library of Medicine, National Institutes of Health

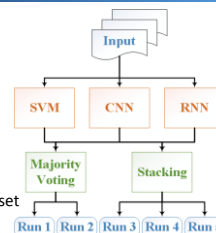
²Department of Computer Science, University of Kentucky

³Division of Biomedical Informatics, Department of Internal Medicine, University of Kentucky

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 1

Outline

- Individual models
 - SVM
 - CNN
 - RNN
- Ensembles of three models
 - Majority voting
 - Stacking
- Experiments
 - 5-fold cross validation on training + dev set
 - Results on test set



Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 2

Chemical-protein relations

- A multiclass classification problem
- The chemical-protein relations occurring in a single sentence

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 3

SVM with rich feature vector

SVM

- Linear kernel
- One-vs-rest scheme

Miwa, M.; Saito, R.; Miyao, Y. & Tsujii, J. A rich feature vector for protein-protein interaction extraction from multiple corpora. *Proceedings of the 2009 Conference on Empirical Methods in Natural Language Processing*, 2009, 1, 121-130

Rich Feature Vector

- Words/Part-of-speech tags surrounding the chemical and gene mentions
- Bag-of-words between the chemical and gene mentions
- Distance between two entity mentions
- Shortest path in a dependency graph

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 4

Shortest path in a dependency graph

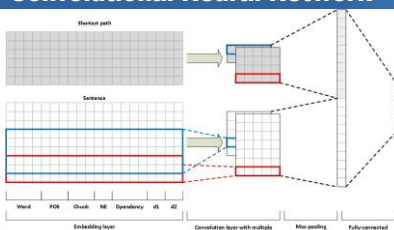
- Obtained using Bllip parser + Stanford dependencies converter

Genfibromide_{CHEMICAL}, a lipid-lowering drug, inhibits the induction of **miR-155**_{miRNA} in human astrocytes.

- Vertex walks
 - CHEMICAL – *nsubj* – inhibits
 - inhibits – *dobj* – induction
 - induction – *nmod:of* – GENE
- Edge walks
 - nsubj* – inhibits – *dobj*
 - dobj* – induction – *nmod:of*

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 5

Convolutional Neural Network



Peng, Y. & Lu, Z. Deep learning for extracting protein-protein interactions from biomedical literature. *Proceedings of BioNLP 2017*, 2017, 29-38

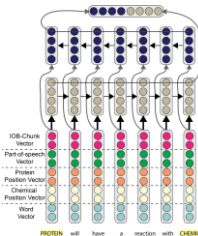
Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 6

Convolutional Neural Network

- Word embedding: 300
 - trained on PubMed using word2vec
- Part-of-speech, chunk and named entities: one-hot encoding
 - Obtained using Genia Tagger
- Convolutional window size: 3 and 5
- Filters: 300

Yan Peng, Anthony Rios, Ramakrish Kavukuru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 7

Recurrent Neural Network



Kavuluru, R., Rios, A. & Tran, T. Extracting Drug-Drug Interactions with Word and Character-Level Recurrent Neural Networks. *2017 IEEE International Conference on Healthcare Informatics (ICHI)*, 2017, 5-12

Yan Peng, Anthony Rios, Ramakrish Kavukuru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 8

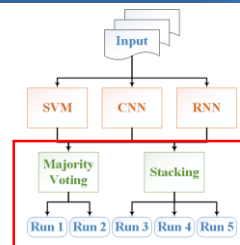
Recurrent Neural Network

- Pairwise ranking loss
 - The output layer has 5 positive classes
 - If all 5 class scores are negative, then we predict the negative class
- Preprocessing
 - Replace word occurs less than 5 times with an UNK token
- Word embedding: 300
 - Obtained from GloVe

Santos, C. N., Xiang, B. & Zhou, B. Classifying Relations by Ranking with Convolutional Neural Networks. *ACI*, 2015, 626-634

Yan Peng, Anthony Rios, Ramakrish Kavukuru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 9

Ensembles of SVM, CNN, and RNN models



Yan Peng, Anthony Rios, Ramakrish Kavukuru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 10

Ensembles of SVM, CNN, and RNN models

Majority voting

- Select the relations that are predicted by more than 2 models

Stacking

- Random Forest classifier
- 17 features:
 - 6 from SVM
 - 6 from CNN
 - 5 from RNN (pairwise ranking loss)

Yan Peng, Anthony Rios, Ramakrish Kavukuru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 11

Results for 5-fold cross validation

- Combine training and development sets
- 5-fold cross validation
 - 60% for training
 - 20% for development (also used to train the stacking systems)
 - 20% for test

Yan Peng, Anthony Rios, Ramakrish Kavukuru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 12

Results of 5-fold cross validation

Models	P	R	F
SVM	0.629	0.478	0.543
CNN	0.641	0.571	0.602
RNN	0.608	0.614	0.609
Majority voting	0.741	0.552	0.632
Stacking	0.755	0.552	0.638

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 13

Results on test set

Run	System	P	R	F
1	Majority Voting	0.7437	0.5529	0.6343
2	Majority Voting	0.7283	0.5503	0.6269
3	Stacking	0.7426	0.5382	0.6241
4	Stacking	0.7311	0.5685	0.6397
5	Stacking	0.7266	0.5735	0.6410

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 14

Results on test set

	System	P	R	F
5-fold CV	Majority voting	0.7408	0.5517	0.6319
	Stacking	0.7554	0.5524	0.6378
Testing	Majority Voting	0.7437	0.5529	0.6343
	Stacking	0.7266	0.5735	0.6410

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 15

Summary and future work

Summary

- Ensemble systems of three models: SVM, CNN, and RNN
- Results are consistent on training + development set and on the test set
- Ensemble methods improved the precisions
- Performance of CNN and RNN are comparable

Future work

- Error analysis
- Fair comparisons between CNN and RNN
- Effects of different parts of deep learning models

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 16

Acknowledgement

- The organizers of the BioCreative VI CHEMPROT task
- Members
 - Yifan Peng, NCBI
 - Anthony Rios, Department of Computer Science, University of Kentucky
 - Ramakanth Kavuluru, Department of Internal Medicine, University of Kentucky
 - Zhiyong Lu, NCBI

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 17

Thank You!
yifan.peng@nih.gov

Yifan Peng, Anthony Rios, Ramakanth Kavuluru, Zhiyong Lu | Chemical-protein relation extraction with ensembles of SVM, CNN, and RNN models | 18