

Department of Computer and Systems Sciences

Enhancing Medical Named Entity Recognition with Features Derived from Unsupervised Methods

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Creating the annotated corpus for training a named entity recognition model is expensive, particularly in specialised domains, such as medicine, which require expert annotators. Moreover, a model trained on text from one medical sub-domain often shows a drop in performance when applied on texts from another sub-domain, and annotated text from this other sub-domain might be required.

When incorporating features from unsupervised methods, to what extent is it possible to:

- Reduce the amount of annotated data needed to achieve a fixed level of performance?
- Reduce the amount of additional annotated data needed for adapting a model to a new sub-domain?

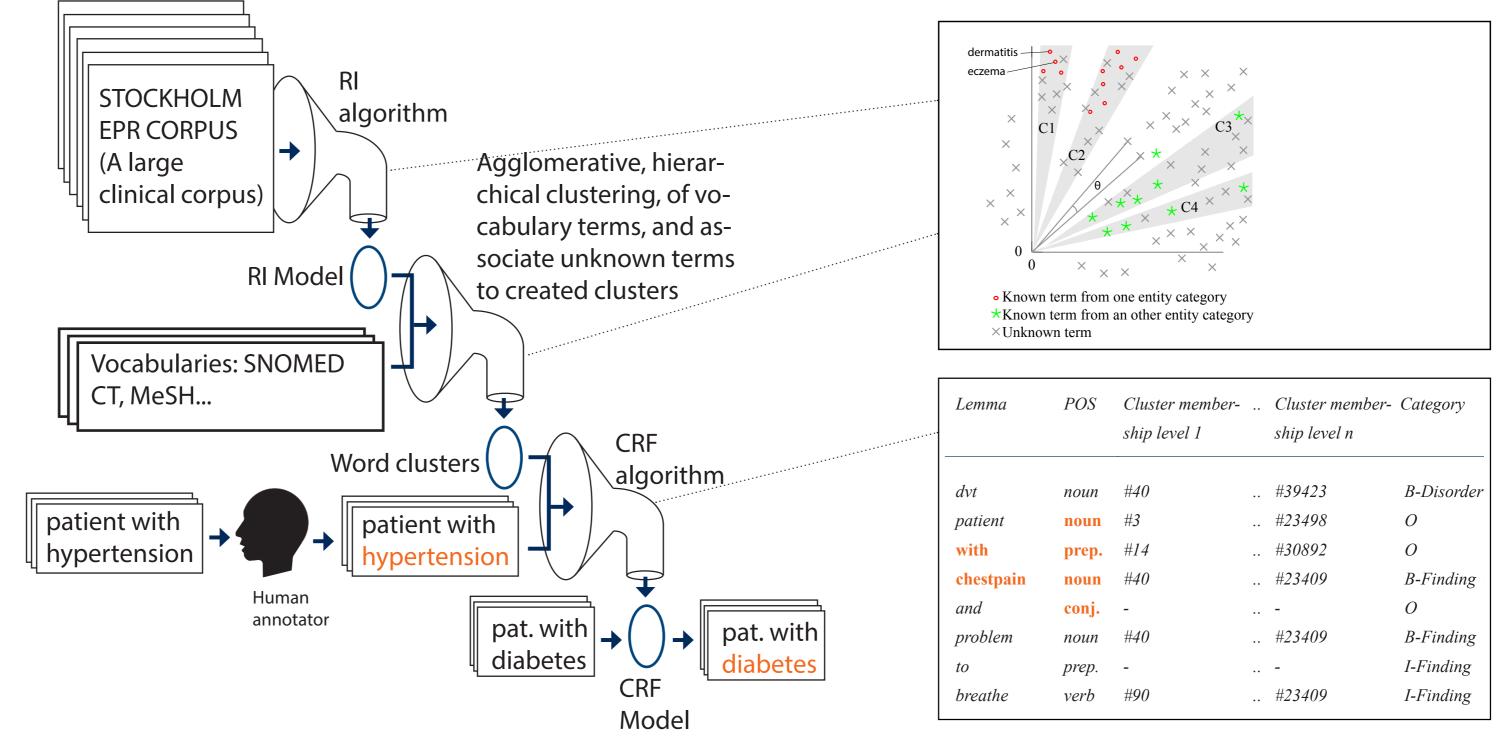
Data sets

Using clustering features for training a CRF model

The entity types disorder, finding, pharmaceutical and body structure annotated in texts from three medical subdomains:

- Internal medicine ER (i.m.)
- Cardiac ICU (Cardiac)
- Orthopaedic ER (Orthop.)

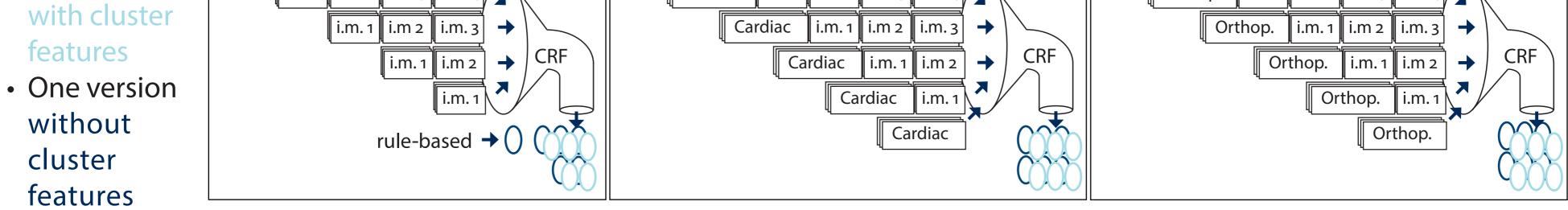
Internal medicine ER data is divided into training data and evaluation data.



Experimental setup

Internal medicine ER training data is divided into 5 partitions, and increasingly more data is used when training the model.

For each	Internal medicine ER (i.m.)	Internal medicine ER+Cardiac ICU	Internal medicine ER+Orthopaedic ER
created model	i.m. 1 i.m 2 i.m. 3 i.m. 4 i.m. 5	Cardiac i.m. 1 i.m 2 i.m. 3 i.m. 4 i.m. 5	Orthop. i.m. 1 i.m 2 i.m. 3 i.m. 4 i.m. 5
 One version 	$\lim_{n \to \infty} 1 \lim_{n \to \infty} 2 \lim_{n \to \infty} 3 \lim_{n \to \infty} 4 \lim_{n$	Cardiac im 1 im 2 im 3 im 4	Orthop lim 1 lim 2 lim 3 lim 4



The L1-norm is used for regularisation when setting the weights of the CRF model. n-fold cross-validation is used on the training data to determine the C-value which governs the strength of the regularisation.

The created models are evaluated against the Internal medicine ER evaluation data.

