Augmenting a De-identification System for Swedish Clinical Text Using Open Resources and Deep Learning

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Abstract

Electronic patient records are produced in abundance every day and there is a demand to use them for research or management purposes. The records, however, contain information in the free text that can identify the patient and therefore tools are needed to identify this sensitive information.

The aim is to compare two machine learning algorithms, Long Short-Term Memory (LSTM) and Conditional Random Fields (CRF) applied to a Swedish clinical data set annotated for de-identification. The results show that CRF performs better than deep learning with LSTM, with CRF giving the best results with an F1 score of 0.91 when adding more data from within the same domain. Adding general open data did, on the other hand, not improve the results.

1 Introduction

Electronic health records (EHR) are today produced in abundance and consist of information valuable to improve the medical care of future patients. They are, however, seldom reused for research as free text in patient records often contain possibly identifiable information about patients. To enable access to electronic health records while preserving patient privacy there is a need for automatic de-identification.

The US Health Insurance Portability and Accountability Act (HIPAA) defines 18 categories of Protected Health Information (PHI) which has to be concealed for EHRs to be considered de-identified in the US (Health Insurance Portability and Accountability Act (HIPAA), 2003). The categories include names, geographic divisions smaller than state, dates related to an individual, contact information and other data that can uniquely identify the individual.

Modules built to identify PHI primarily rely on two methods: Rule-based methods and supervised machine-learning methods (Meystre et al., 2010). The two methods are often used together in hybrid systems (Stubbs et al., 2017). Rule-based methods do not require annotated data for training, are easy to modify and the results are easy to interpret, but they lack robustness and designing rules is a complex task (Meystre et al., 2010). Machine learning methods may provide greater robustness, but require an abundant amount of annotated data. According to Dernoncourt et al. (2017), statistical machine learning models require feature engineering, while artificial neural networks (ANN) does not. The latter does, however, require more data.

Lee et al. (2017) show that training a model on a large source dataset and then fine-tuning by retraining it on the smaller target data set can improve the results in comparison to only using the smallest data set. While the data sets used by Lee et al. (2017) consisted of 29,000 PHI instances in the smaller target data set and 61,000 PHI instances in the larger source data set the largest available Swedish data set, the Stockholm EPR PHI Corpus, has only 4,421 instances of PHI (Velupillai et al., 2009; Dalianis and Velupillai, 2010). It does exist a smaller related corpus with Electronic Health Records with annotations for de-identification, the Stockholm EPR PHI Domain Corpus (Henriksson et al., 2017b). For a larger data set with general Swedish text annotated for named entity recognition, Stockholm Umeå Corpus exists (Östling, 2012).

This study investigates the possibilities of augmenting the quality of de-identification by adding a general Swedish data set for named entity recognition such as Stockholm Umeå Corpus to already existing annotated PHI data sets and secondly the
The use of deep learning methods such as LSTM.

2 Previous research

The state-of-the-art de-identification systems have for a long time been hybrid systems, where a machine learning approach, typically Conditional Random Fields (CRF) is used to identify classes including names, professions, and locations and a rule-based approach is used to identify rarely occurring or regular classes as zip codes, phone numbers and e-mail addresses (Uzuner et al., 2007; Stubbs et al., 2015). The best result during the i2b2 de-identification challenge 2014 (Stubbs et al., 2015) has a micro-averaged entity-based recall of 93.90%, a precision of 97.63% and an F₁ score of 0.96 on i2b2 PHI-categories.

The first neural network de-identification system was introduced in 2016 (Dernoncourt et al., 2017). This system used a type of deep learning with recurrent neural networks (RNN) called long short-term memory (LSTM) with three layers: A character enhanced token-embedding layer, a label prediction layer and a label sequence optimisation layer. The model is bidirectional to better handle long term dependencies. The ANN model presented, performed better than the best system from the i2b2 2014 challenge. Combining Bi-LSTM and CRF further improved the system. Similar systems based on LSTM and CRF have been successful for de-identification (Liu et al., 2017), and during the i2b2 de-identification challenge of 2016 a model combining an LSTM, a CRF and rules won the challenge with an entity-based micro-averaged F₁ score of 0.91 for HIPAA classes (Stubbs et al., 2017).

The largest Swedish dataset with health records annotated for de-identification is the Stockholm EPR PHI Corpus, which is a part of Health Bank - Swedish Health Record Research Bank. Health Bank encompasses structured and unstructured data from 512 clinical units from Karolinska University Hospital collected from 2006 to 2014 (Dalianis et al., 2015). The Stockholm EPR PHI Corpus was first manually annotated by three annotators into 28 PHI classes based on HIPAA and enriched with further classes (Velupillai et al., 2009). The annotations were later on merged into conceptually similar classes while removing classes with few instances, creating a gold standard with eight PHI annotation classes: Age, numeric and non-numeric full dates and McMurtry et al. (2013) have trained decision tree classifiers using 28 features based on part of speech tags, term frequencies, and dictionaries in open journal publications and confidential physician notes to recognise non-PHI words. According to the study, distributional differences between private and open medical texts can be used to classify PHI.

3 Data and method

3.1 Data

Three data sets for de-identification are used: The Stockholm EPR PHI Corpus, the Stockholm EPR PHI Domain Corpus and Stockholm Umeå Corpus 3.0 (SUC). The data consists of both clinical data¹ and open-source data. The Stockholm EPR PHI Corpus is used both for development, training, and testing, while Stockholm EPR PHI Domain Corpus and SUC are only used for training.

All data is encoded using BIOES-encoding, indicating the position of the token within the PHI entity. It encoded whether the token was in the Beginning, Inside or Ending of a multi-token entity, a Single entity or Outside an entity (Reimers and Gurevych, 2017).

Stockholm EPR PHI Corpus consists of 100 patient records from five clinical units: Neurology, orthopaedia, infection, dental surgery and nutrition at Karolinska University Hospital (Dalianis and Velupillai, 2010) and has approximately 200,000 tokens. The Stockholm EPR PHI Corpus was first manually annotated by three annotators into 28 PHI classes based on HIPAA and enriched with further classes (Velupillai et al., 2009). The annotations were later on merged into conceptually similar classes while removing classes with few instances, creating a gold standard with eight PHI annotation classes: Age, numeric and non-numeric full dates and

¹This research has been approved by the Regional Ethical Review Board in Stockholm (2012/834-31/5).
Table 1: Results from Dalianis and Velupillai (2010)

| Class               | Annotated | Retrieved | Relevant | Exact matches | Precision | Recall | F-score | Partial matches | Precision | Recall | F-score |
|---------------------|-----------|-----------|----------|---------------|-----------|--------|---------|----------------|-----------|--------|---------|
| Age                 | 56        | 45        | 37       | 0.822222      | 0.680917  | 0.739763| 0.904762| 0.777061       | 0.836142  |
| Date_Part           | 710       | 654       | 617      | 0.943425      | 0.868051  | 0.900189| 0.946198| 0.917730       | 0.907438  |
| Full_Date           | 500       | 426       | 342      | 0.802817      | 0.684000  | 0.738661| 0.931665| 0.802106       | 0.862045  |
| First_Name          | 923       | 749       | 713      | 0.951936      | 0.739726  | 0.909546| 0.954606| 0.773772       | 0.854729  |
| Last_Name           | 928       | 816       | 777      | 0.952206      | 0.837284  | 0.891055| 0.961653| 0.845848       | 0.896835  |
| Health_Care_Unit    | 1021      | 689       | 589      | 0.811321      | 0.657502  | 0.723982| 0.921497| 0.808116       | 0.732705  |
| Location            | 148       | 73        | 54       | 0.739726      | 0.996861  | 0.886888| 0.778959| 0.379129       | 0.509933  |
| Phone_Number        | 135       | 88        | 80       | 0.930233      | 0.592593  | 0.723982| 0.954198| 0.631105       | 0.746535  |
| Total               | 4,421     | 2,886     | 3,179    | 0.896530      | 0.719068  | 0.798844| 0.941190| 0.751441       | 0.835680  |

The myth annotation consists of names of mythical creatures as a subclass, event and other. The annotations for person, location and age were used in this study, further the person annotation was semi-manually divided into first names and last names. The entire corpus is used.

The Stockholm EPR PHI Corpus was first divided into two sets: One small for development and validation with 10% of the patient records and one for training and testing by cross-validation with 90% of the patient records. For the CRF, tenfold cross-validation was used. The patient records from the Stockholm EPR PHI Corpus were divided into ten folds. The Stockholm EPR Domain Corpus and SUC Corpus were divided into ten folds, where for each fold 90% of the sentences were used for training. Only the folds from the Stockholm EPR PHI Corpus were used for testing. A similar approach was done for LSTM, ates and places and the animal annotation consists of names of animals.
but used validation data for early stopping. The LSTM has only been evaluated on the three first folds due to time constraints.

3.2 Method

This study compares the predictive powers for three models based on the data described above. The first model is only trained on data from the Stockholm EPR PHI Corpus, the second model is trained on data from the Stockholm EPR PHI Corpus and the Stockholm EPR Domain Corpus. The last model is trained on the Stockholm EPR PHI Corpus and SUC. All models are evaluated on data from the Stockholm EPR PHI Corpus using ten cross fold-validation.

The result is evaluated with micro averaged entity-based precision, recall and F$_1$ score, which is the standard for evaluating named entity recognition (Stubbs et al., 2015).

3.2.1 LSTM

Recurrent neural network (RRN) is a type of deep learning artificial neural network designed for processing sequential data (Dernoncourt et al., 2017). The bidirectional LSTM architecture is designed to access long-range dependencies in both forward and backward directions (Dernoncourt et al., 2017). The experiment uses the architecture described in Lample et al. (2016) based on an open-source implementation with Tensorflow.

As stated by Lample et al. (2016), character-based representations can be used to capture both morphological and orthographic information. The character-representations are learned from the used training set for each experiment. Pre-trained word representation is used, based on a subset of clinical text from Health Bank of 200 millions tokens producing 300,824 vectors with a dimension of 300.

The implementation uses the adaptive learning rate method Adam, an algorithm for optimisation of stochastic objective functions (Kingma and Ba, 2014). It computes different learning rates for each parameter based on estimates from the first and second moments of the gradients. The learning rate was set to 0.001 with a decay of 0.9.

Dropout was used with a dropout rate of 0.5. This was used with a batch size of 64. The training is done in a maximum of 20 epochs, with early stopping if no improvement three times in a row in the development set. The model was then evaluated on the test set. The CRF layer (Lample et al., 2016) was not used as it did not show any benefits for the validation set compared to using only LSTM.

3.2.2 CRF

Conditional Random Fields (CRFs) with linear chain is a statistical machine learning method first introduced by Lafferty et al. (2001) that predicts sequences of labels based on sequences in the input. A set of features is typically defined to extract features for each word in a sentence. The CRF tries to determine weights that will maximise the likelihood of leading to the labels in the training data.

In this study, CRFSuite (Okazaki, 2007) is used with a the sklearn-crfsuite wrapper. The features used are: Word as lower case, the first and last four and eight letters, lemma, part of speech tag, if the word is in lower case, upper case or title case, if there are only numbers in the word or only letters in the word or if it has special characters, how many letters, numbers or other characters the word has. This is carried out with a window size of 5. Information about which heading a word comes from is included for texts from the Stockholm EPR PHI Corpus. Furthermore, the CRF uses gazetteers for first names, last names, locations, honorifics or medical profession titles, hospitals in the Stockholm region and regular expressions for identifying date parts, full dates and telephone numbers.

The CRF uses gradient descent with Limited-memory BFGS (L-BFGS) for optimization. LBFGS is an optimization algorithm (Koller et al., 2007).

Lemma and part of speech tagging for each word was performed with Stagger (Östling, 2012) for the Stockholm EPR PHI Corpus and the Stockholm EPR Domain Corpus. SUC is already manually annotated with lemma and part of speech.

4 Results

4.1 LSTM - Results

As seen in Table 3 presenting the results for the LSTM, the systems handle first names, last names, date parts, full dates better than ages, health care units and location.

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3https://github.com/guillaumegenthial/sequence_tagging

4https://sklearn-crfsuite.readthedocs.io
Table 3: Entity-based evaluation for LSTM for the first three folds. The mean is presented for each label. The highest $F_1$ scores are highlighted for each class.

The only two types of PHI improved when adding SUC is age and full date and no improvements can be seen in any other classes. Rather a drop of performance can be seen for location, last names and first names. There is a small increase of recall for first names and locations, but with lower precision.

Stockholm EPR PHI Corpus alone performs considerably better for identifying phone numbers, locations and health care units, while first names and full dates seem to be identified correctly to a greater extent with additional data from the Stockholm EPR PHI Domain Corpus.

Overall, there is no improvement when adding another corpus to the training, but rather a drop in performance.

4.2 CRF - Results

Overall the CRF systems perform well, particularly for finding dates and names. The recall is lower for Phone Number and both the precision and recall is lower for Health Care Unit, Location and Age. As seen in Table 4 with results for the CRF, compared to LSTM results in Table 3, the CRF performs better overall with greater precision, but the LSTM has a higher recall.

Adding the Domain Data increases the $F_1$ score marginally. Some small, likely insignificant, improvements can be seen for Health Care Unit, Age and Phone Number. There is not the same drop of performance as for the LSTM systems.

Adding SUC does not improve the ability to predict, and instead both precision and recall is lower for all classes except last names and ages. The drop of performance is however less severe than for the LSTM.

5 Analysis

Health Care Unit and Location are the most commingled PHI classes. Health care units are often named by their geographic location. Huddinge can for example refer to the hospital Karolinska University Hospital Huddinge but also the municipality Huddinge. In the gold standard, locations are annotated as a part of the health care unit occasionally depending if it is an actual part of the name and whether it is directly adjacent to an identifiable health care unit. Errors are partly caused by the difficulty to distinguish these cases. Furthermore, some health care units are only occasionally annotated as PHI, which also makes it more difficult for the system to learn the structure. ASIH, which stands for Advanced Care At Home in Swedish, is for example in 8 of 20 cases annotated as a singular health care unit entity.

Location is a class with generally low $F_1$ score. One reason for this may be that the test data includes companies as locations. Location has relatively few annotations, and almost one-quarter of these are company annotations. Companies are overall rarely occurring, but frequently mentioned in one patient record. In the record with the most company annotations, none of the seven mentioned companies is found by any system.

When identifying age, the numeral in the age entity is often correctly identified, but the upcoming word is either incorrectly included or missed. The unit following the numeral, often 'years', is occasionally annotated within the PHI and occasionally not, which is one reason for these errors. Age annotations where the numeral is followed by 'årig' (year-old) are found to a greater extent than those followed by 'år' (years).
Uncommon names, common words that are also names, misspelt names and names in lower case are less often identified. This especially happens in contexts where there are no other words, either to the left or right. This is common in sections similar to 'Assigned nurse'. There are some cases where first names are annotated as last names and vice versa. The first names Carina, Riita and Abdul are annotated as last names in the gold standard, leading to errors.

Non-PHI adjacent to a PHI entity is annotated as PHI and more general entities, similar to annotated PHI, for example "the summer of 2007", are more often mistaken as PHI. There are also some cases where inconsistent annotation leads to false positives for especially Health Care Units, they also lead to false negatives. The systems also manage to find some PHI not previously annotated in the gold standard.

### 6 Discussion

In comparison to other work where the Stockholm EPR PHI Corpus is used to classify PHI this set of features and CRF implementation works well for identifying PHI. The CRF also performed better by itself than the LSTM when focusing on the overall $F_1$ score. This the highest recall overall of 88.96% is nonetheless achieved with the LSTM system and without any additional corpus.

There is an overall drop of recall and $F_1$ when adding other corpora to the LSTM version, while the CRF version is slightly improved by adding the EPR PHI Domain corpus, with an $F_1$ score of 0.91. On the other hand the highest recall for *Last name* is achieved using LSTM and SUC and the recall of *First name* and *Last Name* is also improved with additional EPR data in. It could be argued that recall is more important than precision and that *Last name* and *First name* are two of the most sensitive classes.

In SUC, organisations and places are annotated separately. Company names tend to sound like names and occur in similar contexts like health care units. A distinction between locations and companies may enable the usage of the organisation annotation from SUC, with possible improvements on similar labels as well as reduce the heterogeneity.

Differences between the annotation quality or guidelines may also affect the result. Inaccuracies within the Stockholm EPR PHI Corpus is mentioned in the analysis. The Stockholm EPR PHI Corpus was annotated by three annotators and further examined by others. The Stockholm EPR Domain Corpus was, however, originally annotated by only one person and re-annotated for this study by one of the authors to comply with the annotation guidelines of the Stockholm EPR PHI Corpus. This corpus is likely to have more inaccuracies than the Stockholm EPR PHI Corpus.

There is generally a drop in performance between domains and within cross clinical or cross hospital settings. Therefore, it may not come as a surprise that training partially on another domain does not benefit the classifier regardless of the data size. Open text within the medical domain may be more beneficial due to higher domain similarities. A selection of specific documents within SUC is unlikely to benefit the classifier as only a minority of SUC includes medical text.

Using partial match may improve the results for multi-token entity expressions, such as phone

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Table 4: Entity-based evaluation for CRF with tenfold cross-validation. The mean is presented for each label. The highest $F_1$ scores are highlighted for each class.

|                  | EPR PHI          | EPR PHI + Domain | EPR PHI + SUC |
|------------------|------------------|------------------|---------------|
|                  | P %  | R %  | $F_1$ | P %  | R %  | $F_1$ | P %  | R %  | $F_1$ |
| First Name       | 95.05| 92.78| 0.94  | 95.50| 91.41| 0.93  | 94.51| 92.24| 0.93  |
| Last Name        | 97.02| 92.20| 0.94  | 96.39| 90.36| 0.93  | 96.93| 90.08| 0.93  |
| Phone Number     | 92.81| 81.52| 0.87  | 94.58| 84.32| 0.89  | 96.14| 72.14| 0.82  |
| Age              | 79.29| 60.95| 0.68  | 85.09| 71.27| 0.77  | 89.67| 76.21| 0.82  |
| Full Date        | 98.62| 99.15| 0.99  | 96.34| 94.74| 0.96  | 95.82| 93.28| 0.94  |
| Date Part        | 97.06| 95.68| 0.96  | 97.45| 94.73| 0.96  | 95.46| 90.08| 0.93  |
| Health Care Unit | 86.11| 66.40| 0.75  | 88.62| 72.79| 0.80  | 85.45| 67.38| 0.75  |
| Location         | 74.07| 73.70| 0.72  | 76.05| 59.89| 0.66  | 62.40| 70.92| 0.65  |
| Overall          | 93.76| 86.53| 0.90  | 94.66| 86.72| 0.91  | 92.31| 84.97| 0.88  |
numbers, locations, dates and health care units, see Figure 1.

7 Conclusion and future directions

This study aimed to investigate the possibilities of augmenting the quality of de-identification by using annotated data sets for named entities or the use of deep learning methods such as LSTM. The findings suggest that adding data from a general corpus for named entities is not a viable option, but perhaps for individual classes. LSTM performs reasonably well by itself, even if the CRF models seem to perform better. It is worth noting that the LSTM is not yet evaluated on all folds, and considering the increase of recall, it is still warranted to see if a hybrid version of this CRF and LSTM can improve the results further. One possible approach would be to use a LSTM system to de-identify personal names and a CRF system to de-identify phone numbers, locations, dates and health care units.

The current study only examined the effects of using two corpora together as training data, and not the performance when training on one data set, the Stockholm EPR PHI Corpus or SUC, and then using domain adaptation to the target data set, the Stockholm EPR PHI Corpus. While the identification of some PHI classes benefit from added data, there are also classes where no improvements are seen despite data being added.

The analysis has shown that there is a need to revise the old gold standard for the Stockholm EPR PHI by adding previously overlooked PHI, changing PHI accidentally annotated as another PHI, and possibly review the guidelines for the manual annotation of health care units, locations and ages.

Our best performing de-identification system surpasses previous systems based on Stockholm EPR PHI Corpus. It performs in line with the best performing de-identification systems from the latest i2b2 de-identification challenge (Stubbs et al., 2017) but lower than the best from earlier challenge (Stubbs et al., 2015). One observation, however, is that data set in Stubbs et al. (2015) is seven times larger than the Stockholm EPR PHI Corpus in terms of both tokens and PHI instances (Dernoncourt et al., 2017).

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