

Joint segmentation and named entity recognition using dual decomposition in Chinese discharge summaries

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ABSTRACT

Objective In this paper, we focus on three aspects: (1) to annotate a set of standard corpus in Chinese discharge summaries; (2) to perform word segmentation and named entity recognition in the above corpus; (3) to build a joint model that performs word segmentation and named entity recognition.

Design Two independent systems of word segmentation and named entity recognition were built based on conditional random field models. In the field of natural language processing, while most approaches use a single model to predict outputs, many works have proved that performance of many tasks can be improved by exploiting combined techniques. Therefore, in this paper, we proposed a joint model using dual decomposition to perform both the two tasks in order to exploit correlations between the two tasks. Three sets of features were designed to demonstrate the advantage of the joint model we proposed, compared with independent models, incremental models and a joint model trained on combined labels.

Measurements Micro-averaged precision (P), recall (R), and F-measure (F) were used to evaluate results.

Results The gold standard corpus is created using 336 Chinese discharge summaries of 71 355 words. The framework using dual decomposition achieved 0.2% improvement for segmentation and 1% improvement for recognition, compared with each of the two tasks alone.

Conclusions The joint model is efficient and effective in both segmentation and recognition compared with the two individual tasks. The model achieved encouraging results, demonstrating the feasibility of the two tasks.

INTRODUCTION

Electronic medical records have experienced a rapid growth in recent years.¹ While sophisticated algorithms have been developed for English medical records,^{2–8} rare algorithms have been introduced to process Chinese medical records. On the other hand, China is the largest country in terms of the total number of population. Therefore, the large amount of information contained in Chinese medical texts, if processed properly, will certainly benefit the field of biomedical informatics.

In this paper, we mainly focus on two important tasks for Chinese medical text processing: word segmentation^{9–10} and named entity recognition.^{10–11} Due to the nature of the Chinese language, there is no space separating two different words. As a result, the first step for a Chinese language

processor would be to separate Chinese characters into semantic words. For instance, the sentence ‘患者夜间无关节疼痛’ (the patient does not have joint pain at night) should be segmented as ‘患者 (patient)/夜间 (at night)/无 (no)/关节 (joint) 疼痛 (pain)’ (see figure 1). The second task (named entity recognition) involves detecting named entities from discharge summaries. In this paper, we consider the following four categories of named entities: (1) problems and symptoms like ‘咳嗽 (cough)’ and ‘肺炎 (pneumonia)’; (2) medical tests and assays like ‘血压 (blood pressure)’ and ‘CT’; (3) medications like ‘阿司匹林 (aspirin)’; and (4) treatments like ‘化疗 (chemotherapy)’. In the example sentence above, the phrase ‘关节疼痛 (joint pain)’ should be extracted as an entity of type ‘problem’ (figure 1).

Noticeably, there exists a strong correlation between the two tasks. As words represent a group of characters that convey a certain semantic meaning, no named entities should be allowed to contain a fraction of a word, otherwise such a named entity will not have a clear meaning. On the other hand, a named entity is allowed to contain an arbitrary number of words (except for named entities of type ‘medication’, which will be explained in the Methods section). For instance, the named entity ‘失血性休克 (hemorrhagic shock)’ can be separated into two words: ‘失血性 (hemorrhagic)’ and ‘休克 (shock)’.

Out of vocabulary is another big challenge for word segmentation. In the medical domain, there is an enormous number of medical terms that cannot be found in current Chinese word dictionaries, especially medication names like ‘阿乐欣 (azlocillin sodium)’. According to the statistics of our data corpus, out-of-vocabulary words account for nearly 80% in all medication names. This large amount of out-of-vocabulary words make the word segmentation task especially more difficult, compared to word segmentation in other domains. For example, the standard segmentation for ‘肝内胆管多发结石 (many stones in the intrahepatic bile duct)’ is ‘肝|内|胆|管|多|发|结|石 (in the intrahepatic bile duct|many|stones)’, while the segmentation result from the state-of-art segmentation tool is ‘肝|内|胆|管|多|发|结|石 (in the intrahepatic|bile|duct|many|stones)’.

Resolving ambiguities is also a difficult problem for our tasks. For example, the standard results of segmentation for the phrase ‘无畏寒 (no chills)’ is ‘无|畏|寒 (no | chills)’ for segmentation and ‘畏寒 (chills)’ should be identified as a named entity of

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Figure 1 A sample for the two tasks of segmentation and recognition; white space: segmentation tag; blue color: name entity.



type ‘problem’. However, the group of characters ‘无畏’ also stands for ‘fearless’ in Chinese. The segmentation system should be able to resolve these types of ambiguity in order to achieve a good performance.

Due to the complex challenges discussed in the previous paragraphs, existing segmentation tools had an undesirable performance on our data corpus. Our experiments showed that one of the state-of-art Chinese word segmentation tools, MSRSeg,⁹ obtained an F-measure of only 82.58%.

In our work, we first used two independently trained conditional random field (CRF) models to perform the two tasks separately. However, as we have mentioned, this method does not take the inherent correlation between segmentation and named entity recognition into account. To tackle these challenges, we designed joint models for both word segmentation and named entity recognition. We also attempted to exploit the relationship between the two tasks in order to boost the performance of our system. Our new framework uses the dual decomposition algorithm for inference, which solves a linear programming relaxation of the global inference problem. Experiments showed that the new framework achieved a better performance than baseline methods using only independent or incremental models. We also demonstrated that compared to other joint models with similar performance, our joint model using dual decomposition has a much smaller inference complexity and thus results in a much shorter running time.

The data corpus we used is Chinese discharge summaries provided by a Chinese hospital. It includes 336 labeled Chinese discharge summaries. After manual annotation, we labeled a total of 8881 medical problems, 1188 treatments, 782 medications, 1299 tests, and 71 355 words in the data corpus.

Our contributions in this paper are fourfold. First, we created a high-quality corpus with named entity and segmentation annotation in Chinese discharge summaries. Second, two independent models trained on combined labels were introduced to perform the segmentation and entity recognition tasks. Third, we furthermore proposed a new method using dual decomposition to improve the performance of the two tasks based on two independently trained CRF models. Finally, to the best of our knowledge, our work is the first to focus on Chinese discharge summaries.

RELATED WORK

Word segmentation¹² and named entity recognition^{13–14} are traditional topics in natural language processing. For word segmentation, there are three major difficulties in the Chinese language: the construction of language resources, segmentation ambiguity, and out of vocabulary. For named entity recognition, most methods used the information of word segmentation and part-of-speech (POS) tags as features of CRF models. Almost all of these methods solved the two tasks separately or used a pipeline process. In addition, to the best of our knowledge, there is no related work for segmentation and named entity recognition in Chinese discharge summaries.

In natural language processing, some tasks are correlated such as segmentation and named entity recognition, POS tagging and

dependency parsing, etc. Kruengkrai *et al*¹⁵ proposed an error-driven word-character hybrid joint model for Chinese word segmentation and POS tagging. The method achieved a superior performance. Hatori *et al*¹⁶ proposed a joint model for POS tagging and dependency parsing in Chinese. Their performance is considered as a new state-of-the-art performance on this joint task. Srikumar and Roth¹⁷ proposed a joint model that captured the interdependencies between verb semantic role labeling and relations expressed using prepositions, for extended semantic role labeling. In particular, some joint models were related to the dual decomposition algorithm, which was introduced by Dantzig and Wolfe in 1960.¹⁸ Dual decomposition was applied in natural language processing by Rush *et al*¹⁹ and Koo *et al*²⁰ in 2010. They mainly applied the method for two types of problems: combining results from two lexicalized parsing models, and combination of results of a lexicalized parsing model and a trigram POS tagger. The dual decomposition approach has a major contribution by a 0.5% improvement. They also applied the decoding algorithm to phrase-based translation models.^{21–22} In addition, McClosky *et al*²³ applied the dual decomposition algorithm in biomedical event extraction. Three models of joint trigger and argument extraction, capturing correlations between events, and consistency between arguments of the same event were presented. The dual decomposition was used as inference. Their group achieved the first place on the BioNLP 2009 shared task, the BioNLP 2011 Genia task and the BioNLP 2011 infectious diseases task.^{23–26} Luo *et al*²⁷ proposed a dual decomposition method for Chinese predicate-argument structure analysis. Compared with state-of-the-art methods, the F-measure of the proposed method (85.97%) is better than that of state-of-the-art methods (85.3%). Hanamoto *et al*²⁸ proposed a dual decomposition method for coordination structure. They combined head-driven phrase structure grammar (HPSG) parsing and coordinate structure analysis with alignment-based local features. The experiment shows that the joint model is better compared with each of the two algorithms alone. In this work, our joint models focus on segmentation and name entity in Chinese discharge summaries. Up to now, we cannot find a similar study with the two tasks in Chinese discharge summaries. Based on this, we proposed a joint model using dual decomposition for the two tasks in the corpus.

MATERIALS AND ANNOTATION

Dataset

The data, 336 discharge summaries, were collected by random sampling from diverse departments in a hospital in China. We list some statistics, including the average number of characters, sentences, entities, etc, in supplementary material A, tables S1 and S2 (available online only). The annotation guidelines and the annotated corpus are available online at <http://research.microsoft.com/en-us/projects/ehuatu/>.

Annotation guidelines

The annotation for named entities is more involved and we list the detailed annotation guideline for named entity

annotation in the appendix file (see supplementary material B, available online only).

Our segmentation annotation is similar to that of Gao *et al.*⁹ However, in the medical domain, we experienced several issues that were not included in Gao *et al.*⁹ Therefore, aside from rules in Gao *et al.*⁹ we also employ a general rule in order to guarantee a consistent segmentation labeling: if the English translation of a group of Chinese characters is only one word, then the corresponding group of characters is labeled as a single word. For instance, ‘癌转移 (cancerometastasis)’ and ‘球结膜水肿 (chemosis)’ are labeled as single words, although according to Gao *et al.*⁹ they should be labeled as separated words. We also employed several additional rules to make the segmentation annotation more consistent, which we list in the appendix (see supplementary material C, available online only).

Annotation flow

To find a gold standard, annotations were made by annotators with relevant domain backgrounds. Three doctors (A1, A2, and A3) were asked to make annotations. Each record was annotated by two doctors (A1 and A2) independently, and a third doctor (A3) would judge the results as correct or incorrect. The results from the doctors were merged as follows. If the two doctors hold the same opinion, it is decided. If the opinions of the two are different, then the annotation will be determined by the third doctor. However, if the third doctor still cannot decide on the annotation, further discussion must be conducted until an agreement on the annotation is reached. Doctors proficient in medical terminologies are able to find the content that should be annotated accurately; however, they are inconsistent in identifying the boundaries of records due to the diversity of language.

Most of the inconsistency comes from the doctors' dealing with word boundaries. For example, in the phrase ‘偶发结石 (stones occasionally found)’, the doctors were inconsistent on whether to include ‘偶发 (occasionally found)’ in the named entity. Similarly, when annotating drug names like ‘阿莫西林胶囊 (amoxicillin capsules)’, the doctors were inconsistent with whether ‘胶囊 (capsules)’ was part of the medication entity. Occasionally, doctors may annotate two different entities as a single entity because the two entities often occur simultaneously, such as ‘咳嗽咳痰 (cough and expectoration)’.

Inconsistency in annotation may affect the learning model and subsequently lower the performance of our system. Therefore, to obtain a more consistently labeled corpus, a second round of annotation is required. In the second round, three annotators with backgrounds in computer linguistics (B1, B2, and B3) were asked to annotate the record following the same procedures as in the first round. They were given the annotation produced by doctors and were asked to annotate on top of them so that no entities were mislabeled due to a lack of medical expertise. Refined results and a final gold standard were obtained by combining the results from the above two rounds of annotation.

Inter-annotator agreement

Disagreements exist between annotators with different domain backgrounds, but ultimately benefit us in building a more completely annotated corpus, showing the importance of a mixture of both medical background and computer linguistic background in achieving an optimized result. The Kappa statistics were used in this paper to assess the agreements of different annotators.

As shown in supplementary material D, table S3 (available online only), there is a great difference in inter-annotator agreements (IAA) between annotators and the gold standard. This

difference substantially demonstrates the complementary nature of the annotation from different domains. Therefore, it is reasonable to state that IAA between annotators and the final gold standard is of great importance as it denotes the accuracy and the completeness of the annotations.

With high values of k in B1 and B2, the annotation result is very satisfying. Shown in table S3, the k for doctor annotators and the gold standard is rather rough ($k=71.94\%$ and 71.87%), and the k for computer linguistic annotators and the gold standard is higher ($k=91.59\%$ and 90.98%). The experiment shows the substantial agreement between them, which represents the completeness of the annotations.

We also computed the IAA for segmentation annotation, which is shown in supplementary material D, table S4 (available online only).

METHODS

In this section, we first briefly describe the CRF,²⁹ which are widely used to solve sequential labeling problems in the literature of natural language processing.³⁰ We then introduce baseline methods, including independent models and incremental models based on two CRF models. We also introduce a joint model trained on combined labels. Next, we describe the concept of dual decomposition.¹⁹ Finally, we present the joint model using a dual decomposition algorithm for the two tasks. Figure 2 shows the flow charts of the four methods (independent model, incremental model, joint model trained on combined labels (joint_CRF) and joint model using dual decomposition (joint_DD)).

Conditional random fields

We give a brief description of CRF models²⁹ used for sequential labeling. In natural language processing, we consider CRF whose dependency graph is a chain. Let $\vec{o} = (o_1, o_2, \dots, o_n)$ be an observation sequence. Our goal is to assign sequential labels $\vec{x} = (x_1, x_2, \dots, x_n)$ where $x_i \in X$ is the label assigned for the i th instance. For instance, in our segmentation task \vec{o} is the sequence of all Chinese characters of a sentence and \vec{x} can be used to represent a segmentation solution of the sentence \vec{o} . When using CRF models to solve this problem, we first define classes of features $f_{a,i}(x_{i-1}, x_i, \vec{o}, i)$, where $a \in A$ is the type of feature (eg, punctuation, capital letters, etc.) and o_i is the instance to be labeled. A set of parameters λ_a are then trained on the training data and the loss function for a label sequence \vec{x} on an observation sequence \vec{o} can then be expressed as

$$p(\vec{x}|\vec{o}; \vec{\lambda}) = \frac{1}{Z(\vec{o}, \vec{\lambda})} \cdot \exp\left(\sum_{a \in A, x_1, x_2 \in X} \lambda_{a, x_1, x_2} \sum_{i=1}^n f_{a,i, x_1, x_2}(x_{i-1}, x_i, \vec{o}, i)\right), \quad (1)$$

where $Z(\vec{o}, \vec{\lambda})$ is a normalization factor and can be omitted when we only care about the best label sequence. Equation (2) is then used to find the optimal label sequence \vec{x}^* .

$$\vec{x}^* \stackrel{\text{def}}{=} \operatorname{argmax}_{\vec{x} \in X^n} p(\vec{x}|\vec{o}; \vec{\lambda}). \quad (2)$$

Baseline methods: independent models and incremental models

Baseline methods are to view the two tasks as completely independent tasks. Both word segmentation and named entity

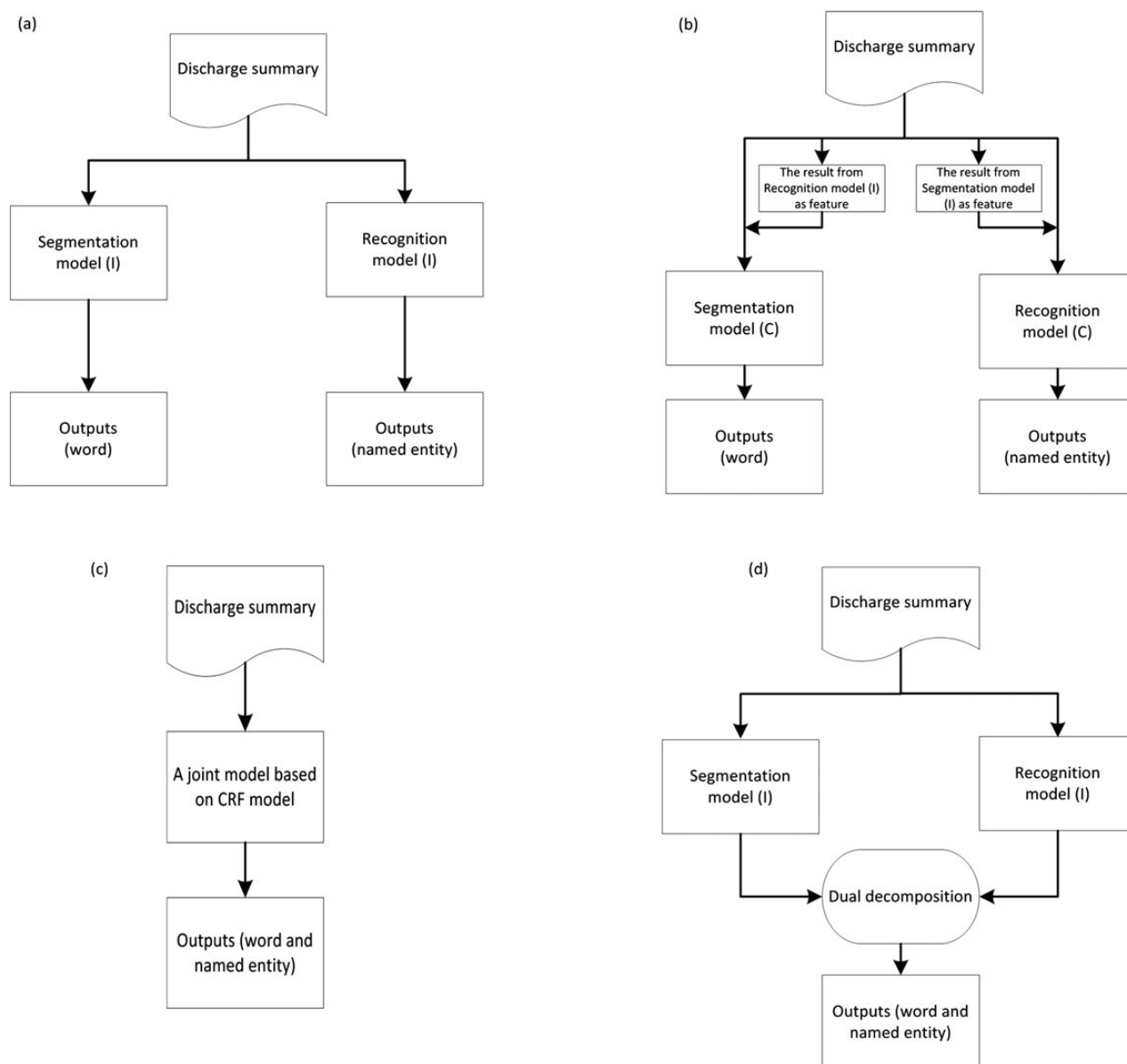


Figure 2 The flow charts of the four methods (independent model, incremental model, joint_CRF and joint_DD). CRF, conditional random field; DD, dual decomposition.

recognition can be viewed as sequential labeling tasks, which are often solved using CRF models in the literature of natural language processing. To perform the two tasks, two sets of different training data (one is for word segmentation and the other is for named entity recognition) were used in two different CRF models to perform them.

Needless to say, independent CRF models do not give satisfactory performance on the two tasks because this method does not take the inherent correlation between the two tasks into account. For example, almost all medication named entities serve as an isolated single word in word segmentation (such as ‘阿司匹林 (aspirin)’) and no named entities will contain a fraction of a segmented word. Therefore, knowing the result of one task can significantly benefit the inference of another task. To this end, we developed incremental models, in which the labeled results of word segmentation of named entity recognition were used as features to train new (and better performed)

models incrementally. Note that the results used as features of segmentation and entity recognition were produced by two independent models. Please see figure 2 for a graphical depiction of independent and incremental models.

Although the incremental model can capture the correlation between word segmentation and named entity recognition to some extent, it has two major fallacies: (1) Segmentation and entity recognition results produced by independent CRF models may not be accurate. As a result, using these inaccurate results as features may lead to the problem of error propagation and lower the performance of the final incremental models. (2) Due to the nature of the CRF algorithm, the incremental models are unable to capture complex correlations between segmentation and entity recognition. For example, the models cannot capture the fact that all medication entities will be segmented as a single word, because these entities may contain many words and such inference is beyond the capability of chain CRF models.

A joint model using a CRF model

Aside from independent and incremental models, we proposed a joint model that was trained on combined labels. That is, instead of training two separate CRF models for word segmentation and named entity recognition, we combined the sequential labels of the two tasks and trained a joint CRF model that performs the two tasks simultaneously. Mathematically, we construct a new label set $Z = X \times Y = \{x + y | x \in X, y \in Y\}$, where '+' means the concatenation of two strings. Subsequently, the loss function of two sequential labels (ie, segmentation results and named entity recognition results) can then be expressed as

$$p(\vec{z}|\vec{o}; \vec{\lambda}) = \frac{1}{Z(\vec{o}, \vec{\lambda})} \cdot \exp \left(\sum_{a \in A, z_1, z_2 \in Z} \lambda_{a, z_1, z_2} \sum_{i=1}^n f_{a, i, z_1, z_2}(z_{i-1}, z_i, \vec{o}, i) \right), \quad (3)$$

which can also be written as

$$p(\vec{x}, \vec{y}|\vec{o}; \vec{\lambda}) = \frac{1}{Z(\vec{o}, \vec{\lambda})} \cdot \exp \left(\sum_{a \in A, x_1, x_2, y_1, y_2} \lambda_{a, x_1, x_2, y_1, y_2} \sum_{i=1}^n f_{a, i}(x_{i-1}, x_i, y_{i-1}, y_i, \vec{o}, i) \right). \quad (4)$$

Clearly, this method not only utilizes the correlation between segmentation and entity recognition tasks, but it also avoids the error propagation problem in the incremental model because the joint model performs the two tasks at the same time. Our experiments also show that the joint model trained on combined labels outperforms both independent and incremental models in various settings.

Nevertheless, there is a major drawback with this joint method: the number of parameters in the joint model is much larger than the number in independent and incremental models. To be more specific, let F , X , Y denote the feature set, label sets for word segmentation and named entity recognition, respectively. Assuming we are using a chain CRF, the joint model will require a total of $|F| \cdot |X|^2 \cdot |Y|^2$ parameters (since $|Z| = |X| \cdot |Y|$), while the independent and incremental models only require $|F| \cdot (|X|^2 + |Y|^2)$ parameters. As a result, the training time required by the joint model is significantly longer than required by independent and incremental models. The large amount of parameters also makes the model perform worse on a smaller corpus, because there might not be enough data to infer the joint model's parameters accurately.

To overcome the shortcomings of this method, we proposed another joint model based on the technique of dual decomposition. Our proposed method provides a flexible way to represent the correlation between segmentation and named entity recognition tasks. Furthermore, the number of parameters in our new joint model is much smaller compared to the joint model trained on combined labels. To begin with, we first introduce the concept of dual decomposition.

Dual decomposition

Dual decomposition, proposed by Rush *et al.*¹⁹ is a classic method to solve optimization problems whose objective

function can be divided into subproblems that can be solved easily.²²⁻³¹ A typical optimization problem that can be solved using dual decomposition is presented as follows.

$$\begin{aligned} \min_{x \in X, y \in Y} & f(x) + w \cdot g(y), \\ \text{s.t. } & \forall i \in [m], T_i(x) = S_i(y), \\ & \forall j \in [k], t_j(x) \leq s_j(y). \end{aligned} \quad (5)$$

where w is the weight of the second task and $T_i(x), S_i(x), t_i(x), s_i(y)$ are usually predicates and the constraints require the two variables (ie, x and y) to satisfy properties T_i, S_i and t_i, s_j simultaneously for all i and j . While the primal optimization problem is often hard and cannot be solved efficiently, its Lagrange dual problem can be solved (or approximated) efficiently as the two subproblems $\min_{x \in X} f(x)$ and $\min_{y \in Y} g(y)$ can be solved efficiently when considered separately as two independent optimization problems.

$$\begin{aligned} \min_{u, v} \max_{x \in X, y \in Y} & f(x) + w \cdot g(y) + \sum_{i=1}^m u_i(T_i(x) - S_i(y)) \\ & + \sum_{j=1}^k v_j(t_j(x) - s_j(y)), \\ \text{s.t. } & \forall j \in [k], v_j \geq 0. \end{aligned} \quad (6)$$

To find the optimal u and v , we used a subgradient method,¹⁹⁻³² which involves iteratively solving the easy subproblems and updates weights u accordingly.

Proposed method: a joint model using dual decomposition

Let M_{SEG} and M_{NER} be two CRF models trained separately for word segmentation and named entity recognition. Let also $p(\vec{x}|\vec{o}; M_{\text{SEG}})$ and $p(\vec{y}|\vec{o}; M_{\text{NER}})$ be conditional probabilities of segmentation \vec{y} and named entity recognition \vec{z} . Clearly, given the CRF models M_{SEG} , M_{NER} and the observation (ie, feature) sequence \vec{o} , the optimal x^{opt} and y^{opt} can be computed efficiently.

$$\begin{aligned} \vec{x}^{\text{opt}} & \stackrel{\text{def}}{=} \arg\max_{\vec{x} \in X} p(\vec{x}|\vec{o}; M_{\text{SEG}}), \\ \vec{y}^{\text{opt}} & \stackrel{\text{def}}{=} \arg\max_{\vec{y} \in Y} p(\vec{y}|\vec{o}; M_{\text{NER}}). \end{aligned} \quad (7)$$

Constraints in the primal optimization problem are correlations between word segmentation and named entity recognition. In our system, we implement two classes of constraints:

1. A named entity shall never contain a fraction of a segmented word. In other words, no group of Chinese characters can be segmented as a single word if these characters contain the border of a named entity.
2. A medication named entity should always be segmented as a single word.

The above constraints are natural rules for word segmentation and named entity recognition and were also carried out when annotating the corpus. Please refer to our description of data corpus to find detailed rules guiding our annotation. Mathematically, for an observation sequence \vec{o} , $t_i(\vec{y}, \vec{o})$ were used to represent whether the i th character is inside a named entity of type 'medication'; $l_i(\vec{y}, \vec{o})$ and $s_i(\vec{x}, \vec{o})$ was used to represent whether the i th character and the $(i+1)$ th

character belong to different named entities or segmented words. A solution (\vec{x}, \vec{y}) is considered feasible if $t_i(\vec{y}, \vec{o}) \leq 1 - s_i(\vec{x}, \vec{o})$ and $l_i(\vec{y}, \vec{o}) \leq s_i(\vec{y}, \vec{o})$ for all i . We then used the algorithm of dual decomposition, which is detailed in Box 1 to find the optimal \vec{x}^*, \vec{y}^* given the two CRF models M_{SEG} and M_{NER} . In our experiment, the maximum number of iterations K is set to 50.

EXPERIMENT

Experiment settings

We now describe the settings of our experiments. The CRF toolkit we used is CRF++.³³ Four methods (independent models, incremental models, a joint model trained on combined labels and a joint model using dual decomposition) were compared for both word segmentation and named entity recognition under three different sets of features. For feature set 1 (Ftr. Set 1), only trigrams of Chinese characters were used as features; for feature set 2 (Ftr. Set 2), we added some other basic features, such as punctuations, digits, English characters, conjunctions (like ‘和 (and)’, ‘伴 (with)’), positions (like ‘左 (left)’, ‘右 (right)’), etc.; for feature set 3 (Ftr. Set 3), apart from features from the two previous feature sets, we added special dictionaries for problem, test and medication entities as features. Please refer to Xu *et al.*³⁴ for a detailed description of the method used to construct these special dictionaries. We also gave more

detailed descriptions of the features we used in supplementary material E (available online only).

Note that we did not integrate any POS features into our recognition systems because mainstream Chinese POS tagging systems perform considerably poorer on Chinese texts in the clinical domain. There are at least two reasons for the poor performance: first, most drug names consist of several Chinese characters and most Chinese POS tagging systems tend to mislabel these characters completely. Take the word ‘氯硝西洋’ (clonazepam, literally ‘chlorine saltpeter west pam’) as an example. The Stanford Chinese POS tagger³⁵ tags this word as ‘氯/CD 硝/M 西/NN 洋/NN’, which is completely wrong. Second, most disease names have a long list of modifying words before the actual disease, which is a syntactical pattern rarely found in ordinary texts like newspapers. For example, the word ‘慢性阻塞性肺炎 (chronic obstructive pulmonary disease)’ has two modifiers ‘慢性 (chronic)’ and ‘阻塞性 (obstructive)’ that modifies the central word ‘肺炎 (pneumonia)’. The second modifier ‘阻塞性 (obstructive)’ is tagged by the Stanford Chinese POS tagger as a verb and is clearly wrong. In summary, as mainstream Chinese POS taggers performs considerably worse on our corpus, we did not use POS tags as features for fear that these features will lower the performance of our system.

Box 1 Subgradient algorithm for dual decomposition

Input Two CRF models $M_{\text{SEG}}, M_{\text{NER}}$; the observation sequence \vec{o} and its length n .

Output A pair of solution (\vec{x}^*, \vec{y}^*) .

Parameters K , the maximum number of iterations; $\alpha_k \geq 0$, the step sizes.

Initialize Weight of each constraint $u_i, v_i \leftarrow 0$.

for $k = 1$ to K **do**

$$\vec{y}^{(k)} \leftarrow \operatorname{argmax}_{\vec{y} \in \mathcal{Y}^n} \left(\vec{y} | \vec{o}; M_{\text{NER}} \right) + \left(\sum_{i=1}^n u_i^{(k)} t_i(\vec{y}, \vec{o}) + \sum_{j=1}^n v_j^{(k)} l_j(\vec{y}, \vec{o}) \right).$$

$$\vec{x}^{(k)} \leftarrow \operatorname{argmax}_{\vec{x} \in \mathcal{X}^n} \left(\vec{x} | \vec{o}; M_{\text{SEG}} \right) - \left(\sum_{i=1}^n u_i^{(k)} (1 - s_i(\vec{x}, \vec{o})) + \sum_{j=1}^n v_j^{(k)} s_j(\vec{x}, \vec{o}) \right).$$

if $t_i(\vec{y}^{(k)}, i) \leq 1 - s_i(\vec{x}^{(k)}, i)$ and $l_i(\vec{y}^{(k)}, i) \leq s_i(\vec{x}^{(k)}, i)$ **for all** i **then**

return $(\vec{x}^{(k)}, \vec{y}^{(k)})$.

for all $i = 1$ to n **do**

$$u_i^{(k+1)} \leftarrow \max(0, u_i^{(k)} + \alpha_k (t_i(\vec{y}^{(k)}, \vec{o}) - 1 + s_i(\vec{x}^{(k)}, \vec{o}))).$$

$$v_i^{(k+1)} \leftarrow \max(0, v_i^{(k)} + \alpha_k (l_i(\vec{y}^{(k)}, \vec{o}) - s_i(\vec{x}^{(k)}, \vec{o}))).$$

return $(\vec{x}^{(K)}, \vec{y}^{(K)})$.

CRF, conditional random field.

Evaluation metric

The three standard performance metrics, precision (P), recall (R) and F-measure (F), are used as evaluation metrics in our task.

In the experiment, we followed the standard leave-one-out method and the cross-validation method. In this paper, fivefold cross-validation and averaged metrics were used.

RESULTS

The experimental results are shown in table 1. In general, joint models achieved better results than independent and incremental models on both tasks. The joint model using dual decomposition achieved an increase of 0.30%, 0.58% and 0.17% on the named entity recognition task under three different sets of features, compared to the joint model using combined task labels. The two joint models also achieved comparable performance on the word segmentation task. Furthermore, as we have mentioned in the previous section, the joint model using dual decomposition inference has a much smaller number of parameters than the joint model using combined task labels, which means the training time required by the former model is much less than required by the latter one. In our experiment, the joint model using dual decomposition used nearly 2 h to finish the fivefold cross-validation while the joint model trained on combined labels used more than 10 h to finish the same task (both models were trained using the third set of features). The experimental results clearly demonstrate the efficiency and effectiveness of our proposed joint model using dual decomposition inference.

DISCUSSION

Annotation of segmentation

In segmentation annotation, the general rule is ‘if the English translation of a group of Chinese characters is only one word, then the corresponding group of characters is labeled as a single word’. We employed this rule mainly to keep a one-to-one correspondence between Chinese and English medical terminologies (eg, ‘球结膜水肿’ and ‘chemosis’), which lays the

Table 1 Performances of four models for two tasks

| | | Named entity recognition | | | | Word segmentation | | | |
|------------|-------------|--------------------------|-------|----------------|-----------------|-------------------|-------|----------------|-----------------|
| | | P | R | F ₁ | ΔF ₁ | P | R | F ₁ | ΔF ₁ |
| Ftr. Set 1 | MSRSeg | – | – | – | – | 79.42 | 86.00 | 82.58 | |
| | Independent | 89.61 | 84.19 | 86.82 | | 95.20 | 94.75 | 94.97 | |
| | Incremental | 89.84 | 85.69 | 87.72 | 0.90 | 95.25 | 94.92 | 95.09 | 0.12 |
| | Joint_CRF | 90.26 | 86.66 | 88.42 | 1.60 | 95.27 | 95.13 | 95.20 | 0.23 |
| Ftr. Set 2 | Joint_DD | 91.50 | 86.11 | 88.72 | 1.90 | 95.30 | 95.25 | 95.27 | 0.30 |
| | Independent | 89.48 | 85.18 | 87.28 | | 95.31 | 95.05 | 95.18 | |
| | Incremental | 89.59 | 86.03 | 87.77 | 0.49 | 95.26 | 95.10 | 95.18 | 0.00 |
| | Joint_CRF | 89.84 | 87.11 | 88.45 | 1.17 | 95.31 | 95.31 | 95.31 | 0.13 |
| Ftr. Set 3 | Joint_DD | 91.25 | 86.92 | 89.03 | 1.75 | 95.31 | 95.38 | 95.35 | 0.17 |
| | Independent | 91.03 | 87.50 | 89.23 | | 95.38 | 95.30 | 95.34 | |
| | Incremental | 90.96 | 88.10 | 89.51 | 0.28 | 95.48 | 95.56 | 95.55 | 0.21 |
| | Joint_CRF | 91.52 | 88.67 | 90.07 | 0.84 | 95.52 | 95.57 | 95.59 | 0.25 |
| | Joint_DD | 92.15 | 88.41 | 90.24 | 1.01 | 95.64 | 95.53 | 95.55 | 0.21 |

| | | Problem | | | Test | | | Treatment | | | Medication | | |
|------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|
| | | P | R | F | P | R | F | P | R | F | P | R | F |
| Ftr. Set 1 | Independent | 90.4 | 87.9 | 89.1 | 90.5 | 82.8 | 86.5 | 79.9 | 66.0 | 72.3 | 91.2 | 71.1 | 79.9 |
| | Incremental | 90.6 | 89.1 | 89.9 | 90.3 | 84.7 | 87.4 | 79.3 | 66.8 | 72.5 | 93.8 | 75.5 | 83.7 |
| | Joint_CRF | 91.2 | 89.8 | 90.5 | 90.9 | 85.6 | 88.1 | 81.0 | 67.9 | 73.9 | 89.9 | 80.4 | 84.9 |
| | Joint_DD | 92.3 | 89.8 | 91.0 | 92.4 | 84.1 | 88.0 | 81.9 | 69.3 | 75.1 | 93.7 | 72.5 | 81.8 |
| Ftr. Set 2 | Independent | 90.4 | 88.7 | 89.5 | 89.0 | 82.6 | 85.7 | 79.9 | 67.1 | 72.9 | 92.5 | 75.9 | 83.4 |
| | Incremental | 90.4 | 89.3 | 89.9 | 89.3 | 84.4 | 86.8 | 79.8 | 68.0 | 73.5 | 93.8 | 77.2 | 84.7 |
| | Joint_CRF | 90.8 | 90.0 | 90.4 | 89.6 | 85.6 | 87.6 | 81.2 | 69.3 | 74.8 | 89.5 | 82.8 | 86.0 |
| | Joint_DD | 92.0 | 90.5 | 91.3 | 92.1 | 84.1 | 87.9 | 81.5 | 69.7 | 75.1 | 94.0 | 75.9 | 84.0 |
| Ftr. Set 3 | Independent | 92.0 | 90.5 | 91.2 | 91.1 | 85.5 | 88.2 | 80.0 | 67.1 | 73.0 | 93.6 | 86.4 | 89.9 |
| | Incremental | 92.0 | 91.0 | 91.5 | 91.0 | 86.0 | 88.5 | 79.7 | 68.3 | 73.6 | 92.8 | 86.8 | 89.7 |
| | Joint_CRF | 92.6 | 91.3 | 91.9 | 91.5 | 87.0 | 89.2 | 81.3 | 70.1 | 75.3 | 92.6 | 88.7 | 90.6 |
| | Joint_DD | 93.1 | 91.2 | 92.1 | 92.2 | 86.3 | 89.2 | 82.4 | 69.7 | 75.5 | 93.7 | 87.6 | 90.6 |

CRF, conditional random field; DD, dual decomposition. Bold indicates the highest values.

groundwork for detecting pair links of the same objects between English and Chinese medical terminologies.

In addition, we labeled body parts as independent words and separate location expressions as independent words. This is because body parts and location expressions are very important information in Chinese discharge summaries: even if information related to body parts and locations is slightly wrong, doctors will have trouble curing patients' diseases. Besides, this rule is useful and effective for building a mapping from expressions of body parts to actual anatomies from each discharge summary. Such a mapping can greatly help physicians to combine all the anatomical information instead of being limited to independent organs or tissues to acquire a comprehensive knowledge of patients' diseases.

Annotation of named entities

Slightly different from the method in I2B2 Challenge 2010 tasks, our method includes two more categories (medication and anatomy) in addition to the three used in the I2B2 Challenge tasks (medical problem, treatment, and test). Dividing medications from treatments will benefit future research on the usage and effectiveness of medications. Marking anatomy in problem and test phrases is also beneficial because it enables us to locate the exact positions of symptoms or medical tests.

However, our annotation method has limitations. For instance, allowing verb phrases and post-entity modifiers in an entity may cause difficulties in the design of automatic named entity recognition systems. Furthermore, there is some important information in clinical narratives that will be missed by our annotation guideline. For example, words indicating the changes in patients'

conditions are often missed, such as '恶化 (worsen)', because these words are often unnecessary verb phrases.

Comparison between Chinese and English corpora

The most significant difference between Chinese and English corpora is the lack of a space between two Chinese words, which requires word segmentation for Chinese text. It is clear that the performance of word segmentation will greatly affect the accuracy of a named entity recognition system, as in the example phrase '停用新赛斯平 (stop using ciclosporin soft capsules)'. If the drug name '新赛斯平 (ciclosporin)' is not segmented as a single word, then it is very difficult for the named entity recognition system to identify the word boundary correctly and the character '新 (new)' may even be identified as an adjective by mistake.

Chinese word segmentation is not an easy task in general, but some properties of Chinese clinical reports make the segmentation and named entity recognition task even harder. For example, most drug names used in Chinese clinical reports are translations of their English counterparts and as Chinese is not an alphabetic language, there are a number of ways to translate one particular medicine, like '阿司匹林', '阿斯匹林', and '阿思匹林' (all referring to aspirin). This additional ambiguity makes the Chinese named entity recognition task much harder than in English. Furthermore, there is no capitalization in the Chinese writing system, which is another challenge because we cannot recognize medication entities based on capitalized letters.

Another interesting difference between Chinese and English named entities in the medical domain is the formation of Chinese words for diseases. Many Chinese disease names contain anatomies, like '肺炎 (pneumonia, lung inflammation

literally)' and '肝炎 (hepatitis, liver inflammation literally)'. This shows that the identification of anatomy entities can help the recognition of some Chinese disease entities, whereas in English such an improvement is rare because the English words for pneumonia and hepatitis do not have any part for the corresponding anatomies (ie, lung and liver).

Methods

We first use an example to demonstrate why the method using independent models suffers from its lack of correlation information between word segmentation and named entity recognition, which eventually lowers its performance on both tasks. In the sentence '患者 在医院 输液 治疗 数天 (the patient was given infusion therapy in the hospital for several days)', the CRF model for word segmentation correctly segmented the above sentence into '患者 (patient)/在 (in)/医院 (hospital)/输液 (infusion)/治疗 (therapy)/数天 (several days)'. However, the CRF model for named entity recognition gave an incorrect named entity '输液 治疗数', which is mistakenly considered by the model to mean 'the number of infusion therapies'. Clearly, this mistake can be avoided if we consider word segmentation and named entity recognition jointly, because '数天 (several days)' is correctly segmented as a single word and hence the phrase '输液 治疗数' should never be allowed to be a named entity because it contains a fraction of a word. As expected, incremental models and the two joint models gave correct answers for both word segmentation and named entity recognition.

Although incremental models can capture some correlation between the segmentation and entity recognition tasks, they suffered from the problem of error propagation because incorrect answers of one task will be likely to misguide the CRF model to give incorrect answers for the other task. We have observed this phenomenon when we were trying to process the phrase '还原性谷胱甘肽 (reduced glutathione)'. The independent models gave incorrect segmentation results for this phrase but gave correct named entity segmentation results, mainly because this phrase is enclosed with commas at both ends and the CRF model for named entity recognition correctly identified the named entity. However, when the (incorrect) initial segmentation results were used as features to train a CRF model incrementally for entity recognition, the newly trained model suffers from the incorrect segmentation and extracted an incorrect entity ('原性谷胱甘肽', missing the leading character '还').

In contrast to incremental models, our proposed joint model uses dual decomposition to coordinate results from the two tasks and avoid being misled by incorrect answers of one task. In fact, because the named entity recognition system correctly identified the phrase as a medication name and we have specified constraints requiring all named entities of type medication must be labeled as a single word when segmenting the text, the dual decomposition algorithm automatically changes the segmentation labels to the correct ones. This example clearly demonstrates the effectiveness and fitness of our proposed method for word segmentation and named entity recognition tasks.

Another important advantage of our proposed joint model based on dual decomposition is its smaller inference complexity. Unlike a complete joint CRF model, which combines labels for the two tasks altogether, our proposed method trains models for each task separately, which significantly reduces the amount of inference time. In our experiment, the joint CRF model requires an average of 10 h to finish training while our proposed dual decomposition method completes training in less than 3 h. Note that the number of labels used in the joint CRF model is

proportional to the product of the number of labels used in each task; we expect the joint CRF model runs much slower than our proposed dual decomposition method as the number of tasks and their labels grow larger.

Errors in the treatment category

Noticeably, in all the four semantic categories (ie, problems, tests, medication and treatments), the performance of treatment identification (especially the recall) is significantly lower than the performance of the other three categories. This is due to the complicated structure of treatment entities: unlike the other three categories whose entities are mostly noun phrases (eg, '咳嗽 (cough; problem)', '血压 (blood pressure; test)', '阿司匹林 (aspirin; medication)', etc.), many treatment entities involve verb phrases (eg, '扩张支气管' (expansion of bronchus, literally 'expand (扩张) bronchus (支气管)') and '祛痰' (expectorant, literally 'remove (祛) sputum (痰)'). As most current named entity recognition (NER) algorithms focus on entities that are noun phrases, the structure of treatment entities makes its recognition understandably harder.

Domain difference

The performance of state-of-the-art segmentation methods in common corpus is 96%/F or so³⁶ while our best performance is 95.5% or so in Chinese discharge summaries. This clearly demonstrates that our annotation guideline is a practical one because both methods achieved comparable performance.

CONCLUSION AND FUTURE WORK

In this paper, we created a new corpus of word segmentation and name entity recognition based on Chinese discharge summaries. Named entities are divided into four categories: problem, test, treatment, and medication. We described the annotation guideline of word segmentation in detail. Combining the correlation between named entity recognition and word segmentation, a new joint model using dual decomposition is presented to perform the two tasks. Experiments demonstrate that the joint model has evident advantages in terms of performance as well as running time, compared with other baseline models.

In future, dual decomposition as a framework will be designed for several primary and essential tasks in Chinese discharge summaries, such as POS tagging and dependency parsing.

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