

Incorporating Dependency Trees Improve Identification of Pregnant Women on Social Media Platforms

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Abstract

The increasing popularity of social media lead users to share enormous information on the internet. This information has various application like, it can be used to develop models to understand or predict user behavior on social media platforms. For example, few online retailers have studied the shopping patterns to predict shopper's pregnancy stage. Another interesting application is to use the social media platforms to analyze users' health-related information. In this study, we developed a tree kernel-based model to classify tweets conveying pregnancy related information using this corpus. The developed pregnancy classification model achieved an accuracy of 0.847 and an F-score of 0.565. A new corpus from popular social media platform Twitter was developed for the purpose of this study. In future, we would like to improve this corpus by reducing noise such as retweets.

1 Introduction

The web has become a powerful medium for disseminating information about diverse topics, people can share information anytime and anywhere. Real-time user generated information on the web, epitomized by social media and in particular microblogs, are becoming an important data source

to complement existing resources for disease surveillance (Brownstein, Freifeld, & Madoff, 2009), behavioral medicine (Ayers, Althouse, & Dredze, 2014), and public health (Dredze, 2012). Studies have shown that 26% of online adults discuss health information using social media (GE-Healthcare, 2012), with approximately 90% women using online media for health-care information, and 60% using pregnancy related apps for support. These statistics suggest that social media sources may contain key information regarding specific cohorts, such as pregnant women, and their drug usage habits. Twitter—a micro-blogging site which is actively used by over 328 million users¹—is a very popular social network currently being extensively used for public health monitoring tasks (Chandrashekar, Magge, Sarker, & Gonzalez, 2017; Jonnagaddala, Jue, & Dai). It is also an attractive resource for biosurveillance related shared tasks and competitions because it carries health-related knowledge expressed by various cohorts (Adam, Jonnagaddala, Chughtai, & Macintyre, 2017). However, the noisy nature of data on Twitter demands sophisticated models and techniques for mining the knowledge encapsulated.

The primary aim of this study is to detect whether a tweet convey pregnancy or not. This information further downstream can be used to study the safety of drugs in pregnancy are of paramount importance. Typically, pregnant woman in social media are detected using simple regular expressions and rules such as – matching the phrase

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¹ <https://about.twitter.com/company>

“*i am twenty weeks pregnant*” (Chandrashekar et al., 2017; Wang, Paul, & Dredze, 2014). However, employing rule based detection can lead to many false positives since it doesn’t consider context or sentiment embedded in the tweet. Often, the tweets recognised by rule based methods seem to be sarcastic. For example, consider the tweet “*I look like I’m 6 months pregnant*”. This tweet is a sarcastic tweet and the user actually is not pregnant. Thus, in order to overcome this issue, we propose to use a tree kernel-based approach to detect pregnant woman more effectively. Tree kernel-based approaches have been applied to many different researches, such as relation extraction (Culotta and Sorensen, 2004), question classification (Zhang and Lee, 2003) and protein interaction detection (Miwa et al., 2010). In recent years, tree kernel-based models were used to analyze Twitter data, but most of those studies were focused on opinion mining and sentiment classification (Agarwal et al., 2011; Alicante et al., 2016). In this study, we investigate the effectiveness of applying the approach on the task of determining whether a tweet is posted by a pregnant woman or not.

2 Related Work

Most of the studies in mining Twitter are focused on drug safety domain, e.g. drug abuse and adverse drug reaction (Dai, Touray, Wang, Jonnagaddala, & Syed-Abdul, 2016; Sarker et al., 2016). However, this information can also be used for health surveillance of pregnant women. The study most related to ours is presented by (Chandrashekar et al., 2017) in which they annotated 1,200 tweets with pregnancy announcements to allow the identification of pregnancy trimesters. Klein et. al, constructed an annotated corpus from Twitter focusing on personal medication intake (Klein, Sarker, Rouhizadeh, O’Connor, & Gonzalez, 2017). In an another related study an integrated corpus composing of 2,000 sentences from Twitter and PubMed called TwiMed was presented (Alvaro, Miyao, & Collier, 2017). The corpus contains the annotations of diseases, symptoms and drugs, and their relations.

Tree kernel-based approaches have been widely applied to text classification tasks. Zhang and Lee (2003) utilized tree kernel to question classification and demonstrated that syntactic structures is useful for questions classification. However, the space of tree fragments is too large to compute their inner products. In order to recursively and efficiently compute the common substructures similarity be-

tween two trees, Moschitti (2006) proposed convolution tree kernel and developed a toolkit for public. Wang et al. (2009) adopted convolution tree kernel to find out similar questions from Yahoo Answers dataset. They observed that the convolution tree kernel function can effectively utilize the syntactic structure of a sentence. On the other hand, Agarwal (2011) applied partial tree kernel (PTK) to classify sentiment polarity of twitter data and achieved remarkable performance. The kernel provides the ability to analyze additional semantic information by considering the contribution of shared subsequences containing all children of nodes. PTK compares words by the order of alphabets to determine word similarity for ameliorating the weak point of convolution tree kernel. Later on Croce et al. (2011) proposed smoothing partial tree kernel (SPTK) and improved the calculation method of word similarity by using singular value decomposition (SVD) to transform all words into vectors for determining the cosine similarity between them.

In this paper, we built models based on SPTK and three kinds of tree structures. Besides part-of-speech (POS) tags, the new tree structures incorporate dependency tree and grammatical relations for extracting more useful features. Furthermore, we used word embedding to substitute the vectors generated by SVD. Many studies have demonstrated that word embedding is really useful in many natural language processing tasks. With this in mind we also investigated the effectiveness of combining word embedding with SPTK and different tree structures on the task of identifying pregnancy women on Twitter.

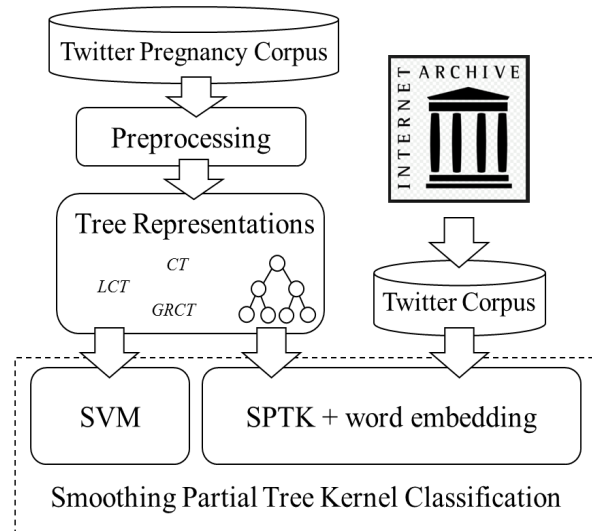


Figure 1: System architecture

3 Methods

The Figure 1 shows the architecture of the proposed pregnancy detection method. The architecture comprises three key components: preprocessing, tree representations, and smoothing partial tree kernel classification. Firstly, the preprocessing component processes a set of tweets that may convey pregnancy (called *candidate tweets* hereafter) through heuristic rules. Then, each candidate tweet is represented by three kinds of tree structures for capturing the information of syntactic, content, and semantic of the tweet. Finally, the smoothing partial tree-kernel classification component measures the similarity between tweet in terms of their tree structures, and the tree kernel is incorporated into support vector machine (SVM) for learning a classifier. We elaborate each component in the following sub-sections.

3.1 Preprocessing

Given a tweet, we first apply the Stanford parser² to generate the output of parse tree and PoS tagging. We further remove stop words and URLs from tweets. In addition, we observe that retweet is impossible to convey pregnancy since Twitter's retweet feature is to help users quickly share that tweet with all of users' followers. Therefore, we filter out tweets if the text start with "RT" (i.e. retweet). By filtering out retweets, the rest are the candidate tweets which may convey pregnancy.

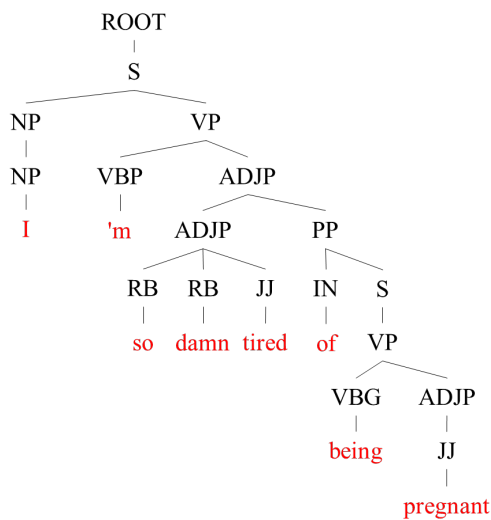


Figure 2: Constituency tree (CT)

3.2 Tree Representations

Different tree representations in tree kernel-based approach may lead to modeling more effective syntactic or semantic feature spaces. In this paper, three kinds tree structure are used to represent tweet, they are constituency tree (CT), lexical centered tree (LCT), and grammatical relation centered tree (GRCT). To facilitate comprehension of the different tree representations, we take a pregnant woman's Tweet "I'm so damn tired of being pregnant" as an example.

Figure 2 is CT, which is the basic tree representation generated by Stanford Parser. The parser works out the grammatical structure of sentences by grouping words together as phrases that could represent the subject or object of a verb. However, CT only contain information of the grammatical structure. Croce et al. (2011) proposed GRCT and LCT to complement CT. GRCT and LCT involve grammatical relations (GR), PoS tags and dependencies. GRCT adds tags of grammatical relations and lexical information as new nodes in CT to emphasize grammatical relationship information while LCT enhance the lexical information by adding grammatical relations and PoS-tags as the rightmost children. Figure 3 and Figure 4 show the same example sentence for GRCT and LCT.

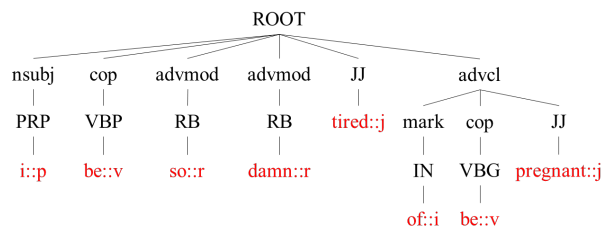


Figure 3: Grammatical relation centered tree (GRCT)

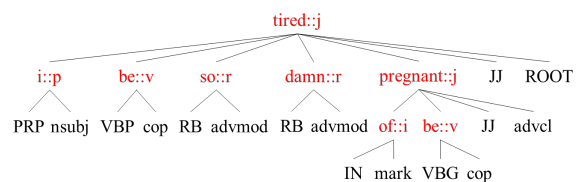


Figure 4: Lexical centered tree (LCT)

3.3 Smoothing Partial Tree Kernel Classification

In SVMs, a kernel function is employed to cleverly compute the similarity between two instances

² <http://nlp.stanford.edu/software/lex-parser.shtml>

without requiring the identification of the entire feature space. In the case of tree kernel, it represents tree in terms of their substructures and evaluates the number of common tree fragments between two trees T_1 and T_2 through the following equation:

$$K(T_1, T_2) = \sum_{n_1 \in N_{T_1}} \sum_{n_2 \in N_{T_2}} \Delta(n_1, n_2) \quad (1)$$

where N_{T_1} and N_{T_2} denote the sets of nodes in T_1 and T_2 , respectively. The function $\Delta(n_1, n_2)$ is equal to the number of common fragments rooted in the n_1 and n_2 nodes. Since the number of different sub-trees is exponential with the parse tree size, it is computationally infeasible to directly use the feature vector.

In recent years, multiple tree kernels have been proposed for resolving this computation issue, such as syntactic tree kernel (Collins and Duffy, 2002), partial tree kernel (Moschitti, 2006), and lexical semantic kernel (Basili et al., 2005). However, the lexical in these tree kernels must belong to the leaf nodes of exactly the same structures limits its applications. Trivially, it cannot work on dependency trees. Croce et al. (2011) proposed a much more general smoothed tree kernel (i.e. smoothing partial tree kernel, SPTK) that can be applied to any tree and exploit any combination of lexical similarities, respecting the syntax enforced by the tree. Therefore, we adopt SPTK to capture the syntactic similarity between the above high dimensional vectors implicitly, as well as partial lexical similarity of trees. The $\Delta_{SPTK}(n_1, n_2)$ can be defined as follows:

- (1) If nodes n_1 and n_2 are leaves, then $\Delta_{SPTK}(n_1, n_2) = \mu\lambda\sigma(n_1, n_2)$
- (2) Otherwise, calculate $\Delta_{SPTK}(n_1, n_2)$ recursively as:

$$\Delta_{\sigma}(n_1, n_2) = \mu\sigma(n_1, n_2) \times (\lambda^2 + \sum_{\vec{I}_1, \vec{I}_2, l(\vec{I}_1)=l(\vec{I}_2)} \lambda^{d(\vec{I}_1)+d(\vec{I}_2)} \times \prod_{j=1}^{l(\vec{I}_1)} \Delta_{\sigma}(c_{n_1}(\vec{I}_{1j}), c_{n_2}(\vec{I}_{2j}))), \quad (2)$$

where σ is any similarity between nodes, $\mu, \lambda \in [0, 1]$ are two decay factors, \vec{I}_1 and \vec{I}_2 are two

sequence of indices, which index subsequences of children u , $\vec{I} = (i_1, \dots, i_{|u|})$, in sequences of children s , $1 \leq i_1 < \dots < i_{|u|} \leq |s|$, i.e., such that $u = s_{i_1} \dots s_{i_{|u|}}$, and $d(\vec{I}) = i_{|u|} - i_1 + 1$ is the distance between the first and last child. c is one of the children of the node n , also in indexed by \vec{I} . This provides an advantage that tree fragments can be matched by applying word embedding similarity σ . Even those tree fragments are not identical but are semantically related.

3.4 Dataset

To the best of our knowledge, there is no openly available corpus for pregnant woman detection. Therefore, we compiled a dataset for the performance evaluation. We employed Tweetinvi³ to collect tweets mentioning pregnancy written in English from May 1, 2017 to May 29, 2017. To retrieve the tweets, we used a list of pregnancy-related query terms to search tweets online. For all 14,824 collected raw tweets, we pre-processed them by removing emoticons, line feeds, extra spaces and dots based on regular expression. The collected tweets contain duplications owing to the same tweets retrieved by different queries and the retweets shared by different users. We removed duplicated tweets or tweets contain similar descriptions by calculating Levenshtein distance among the collected tweets. If the similarity score of two tweets is larger than 70%, we discard the shorter one and reserve the longer one. Finally, we obtained 7,984 tweet sentences.

We randomly selected 3,000 tweets from the collected dataset to build an initial corpus. Five annotators were recruited to annotate the corpus by using MAE (Multi-document Annotation Environment) (Rim, 2016). They determined whether the tweet authors are pregnant or not based on the context information and gave ‘‘Yes’’ or ‘‘No’’ annotations indicating positive or negative cases. A preliminary consistency test was conducted on 500 tweets by having the first two of the annotators annotate the data, while the last one checked their annotations for consistency. The Fleiss' kappa coefficient value for the initial consistency test is 0.42 (moderate agreements). After examining the consistency, all annotators adjusted their annotations and re-annotated the entire data set. After finishing the annotation process, a voting method was employed to determine the

³ <https://github.com/linvi/tweetinvi>

structures together with two tree kernels that incorporate the dependency and grammatical relation information represented in the form of tree structures. We evaluated our models on manually annotated Twitter corpus specifically developed for the purpose of this study. The results demonstrate that employing dependency tree can improve the performance of pregnancy detection. We also observe that lexical features as central node is effective in representing pregnancy information in a tweet. The best performed model had an F-score of 0.5652 on our corpus. The SPTK model considers lexical similarities on the tree structure.

In future, we would like to explore different ways to integrate deeper semantics into tree structure for pregnancy detection on social media platforms. Moreover, we also would like to improve the corpus by ignoring retweets, avoid possible noise and obtain more meta data such as timestamp details. In addition, the pregnancy related tweets may reveal the other health behaviour related information of the of the pregnant authors like drug usage, food intake, any disease symptoms, and emotion. We would like to use this information in future to conduct syndromic surveillance on pregnant women using social media.

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