

GREED: Graph Learning Based Relation Extraction with Entity and Dependency Relations

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Abstract: A large number of electronic medical documents are generated by specialists, containing valuable information for various medical tasks such as medical prescriptions. Extracting this information from extensive natural language text can be challenging. Named Entity Recognition (NER) and Relation Extraction (RE) are key tasks in clinical information extraction. Systems often rely on machine learning and rule-based techniques. Modern methods involve dependency parsing and graph-based deep learning algorithms. However, the effectiveness of these techniques and certain features is not thoroughly studied. Additionally, it would be advantageous to properly integrate rules with deep learning models. In this paper, we introduce GREED (Graph learning based Relation Extraction with Entity and Dependency relations). GREED is based on graph classification using Graph Convolutional Network (GCN). We transform each sentence into a weighted graph via dependency parsing. Words are represented with features that capture co-occurrence, dependency type, entities, and relation verbs, with focus on the entity pair. Experiments on clinical records (i2b2/VA 2010) show that relevant features efficiently integrated with GCN achieve higher performance.


1 INTRODUCTION

There is a large amount of unstructured text in Electronic Health Records (EHR) which contain rich information about patients and clinical events. In healthcare, there is more than 80% of the total data is unstructured and it is created by hospitals, healthcare clinics, or biomedical labs (Kong, 2019). The text in this data is written in natural language and must be read carefully by a medical expert to extract the desired information. However, this operation is costly and time-consuming given the huge amount of available documents.

In fact, clinical records, such as discharge summaries and progress reports, are useful for many computerized clinical applications such as decision support systems. Accordingly, the application of natural language processing (NLP) (Quevedo and Chicaiza, 2023) technologies in the automatic extraction of information from narratives is becoming increasingly interesting. Named entity recognition (Landolsi et al., 2022b) and Relation Extraction (RE) are key components of information extraction tasks in the clinical

domain (Landolsi et al., 2022a). Detecting and classifying the annotated semantic relationships between medical named entities mentioned in biomedical texts has numerous applications that range from advancing basic science to improving clinical practice (Luo et al., 2017).

Significant research on RE has been carried out on unstructured text. Most of the systems are relied on rule-based and machine learning-based approaches (Yang et al., 2021). Machine learning-based methods (Yang et al., 2021; Mahendran et al., 2022; Ed-drissiya El-allaly et al., 2022) usually generate high-quality features for sentences or words based on NLP to train a deep learning classifier. The rule-based methods (Chikka and Karlapalem, 2018; Ben Abdesslem Karaa et al., 2021; Kim et al., 2021) consist in constructing and applying rules which often rely on dependency parsing, entity co-occurrence detection, and pattern matching. Some recent methods use graph-based deep learning algorithms to incorporate dependency parsing and other features (Ed-drissiya El-allaly et al., 2022). However, this technique and the use of some types of features are not well explored although they can be of greater benefit for this task. The preparation of rules often requires a man-

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ual effort by medical experts (Ben Abdesslem Karaa et al., 2021). Also, better integration of the rules with the deep learning models would be of better benefit (Chikka and Karlapalem, 2018).

In our work, we have trained a Graph Convolutional Network (GCN) model (Yuan et al., 2022) to perform the RE based on graph classification. For that, we used dependency parsing to transform each sentence into a weighted graph.

The remainder of this paper is organized as follows. Section 2 discusses state-of-the-art methods for relation extraction. Section 3 presents our model named GREED; whereas section 4 conducts an extensive experimentation of GREED using standard criteria. The final section concludes this paper and discusses future research directions.

2 RELATED WORK

We can categorize the relation extraction methods of the literature into two main categories according to the used techniques: rule-based methods and machine learning-based methods (Yang et al., 2021).

The rule-based approach consists of preparing rules to be inferred from the text to precisely extract information about relations. Usually, supplemental knowledge data can be used with these rules. As an advantage, useful patterns for precise extraction can be explored by the syntactical analysis of the sentence. In addition, pertinent knowledge data can be efficiently exploited by rules. However, a huge effort from medical experts is usually required to define the rules and the knowledge data to cover most of the possible cases. Also, these rules are usually domain-specific and not adaptable to other types of data. For example, Chikka and Karlapalem (2018) use some post-processing rules based on dependency parsing to improve the results of a BiLSTM model trained on word and sentence features. Ben Abdesslem Karaa et al. (2021) relied on the matching tool MetaMap (Aronson and Lang, 2010) to search semantic types and concepts from the UMLS meta-thesaurus (Bodenreider, 2004). Thus, they have applied some NLP techniques to generate sentence features and train an SVM (Hearst et al., 1998) model. The method of Kim et al. (2021) applies some syntactical rules on the entities to generate lexical features to train an online gradient descent model. Then, this method uses some post-processing rules based on some trigger phrases and terms prepared manually.

The principle of machine learning-based methods is to learn to classify each sentence with an entity pair into the corresponding relation class. As an advan-

tage, this approach can efficiently incorporate several types of features, especially by using deep learning models such as GCN. In addition, it can capture the contextual dependency between words during classification. However, it requires a big number of training samples which are manually annotated by medical experts. Also, relevant features and suitable techniques must be chosen carefully for high-quality learning. For example, the method of Li et al. (2019) is based on a BiLSTM model with an embedding of the shortest dependency path between the entities generated by CNN in addition to dependency type features. Zhou et al. (2020) have generated weighted dependency graphs based on shortest path edges to extract generic relations by using a GCN model. Patel et al. (2021) use a CNN model to generate lexical and sentence features. Ed-drissiya El-allaly et al. (2022) have combined BERT with GCN in a joint model to perform an N-level sequence labeling task.

In this paper, we propose a hybrid method named GREED (Graph learning based Relation Extraction with Entity and Dependency relations) which is based on GCN to classify each sentence into relation classes. For that, the sentence is transformed into a weighted dependency graph which is well pre-processed to be suitable for GCN. Also, we have extracted several crucial features for each word which take into account: word co-occurrence, dependency types, entities, and relation verbs. The relation verbs feature represents relevant side information. For that, we have used a simple rule to match the verbs after collecting them automatically from the training set. The features, edge weights, and the GCN model pay attention to the selected entity pair to focus on the most important information. Our model is detailed in the next section.

3 OUR PROPOSAL

3.1 Motivation

In our work, we have chosen the GCN model to process a sentence weighted dependency graph and incorporate several word features. This model was used by some recent methods (Mahendran et al., 2022; Li et al., 2022; Yuan et al., 2022) and is known to be suitable for this task. This approach is better than analyzing the words sequentially since predicates are significant for our task (Zhou et al., 2020). Usually, most methods give an uniform weight for the dependency edges. However, we have added weighted edges based on the shortest paths to be more adequate for the GCN layer (Zhou et al., 2020). In addition,

we were inspired by the work of Yuan et al. (2022) by paying more attention to the edges that are closer to the entity pair. Therefore, we have assigned higher weights to these edges. To focus on the entity pair-related information, we have added an entity-attention mechanism after the GCN layers based on the mask pooling (Zhou et al., 2020). Moreover, we have used the relative positions of the entities as word features (Li et al., 2022). While the word co-occurrence is important information for our task (Shahab, 2017; Perera et al., 2020; Mahendran et al., 2022), we have used a clinical GloVe model (Flamholz et al., 2022) to generate word embedding. Usually, the difference in the dependency relation types is usually ignored. However, we have added the dependency relation type to the word features. Relevant side information is usually ignored while it add knowledge beyond the sentence (Yuan et al., 2022; Mahendran et al., 2022). Thus, we have automatically constructed a dictionary of relation verbs. These verbs are matched based on a syntactic dependency parsing to generate features (Chikka and Karlapalem, 2018). In addition, we have used other essential features such as the part of speech and the entity types. To make a contextual word representation and take into account the sequential dependency, we have added a BiLSTM layer after the input features layer (Yuan et al., 2022).

3.2 Architecture

The principle of our method GREED is to classify the weighted dependency graph of the sentence by using a GCN model with the incorporation of several word features. Thus, GREED has to detect which relation can exist between a given entity pair. Firstly, the sentence is transformed into a dependency graph where the edges between words are weighted according to the shortest paths and the proximity to the entity pair. Furthermore, each word is represented by a features vector which includes: clinical GloVe, PoS, entity type, dependency type, relation verb matching, and distance to the entity pair. To make the features capture the contextual and sequential information, a BiLSTM layer is added after the input features. Thus, the graph and the features are fed into the GCN model with an entity-attention mechanism based on mask pooling. Thus, the output is the corresponding relation class of the input sentence and entity pair. The main steps of our proposal are depicted in Figure 1.

3.3 Graph Construction

The sentence is transformed into a graph $G = (V, E)$ where V is the number of nodes (words) and E is the

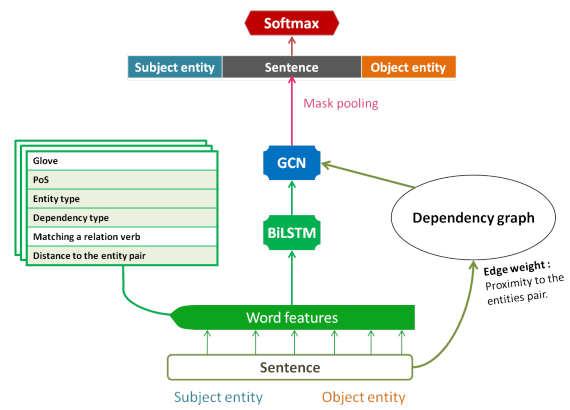


Figure 1: The general architecture of our proposal.

number of edges that represent the dependency relations. Each edge has a weight and each word has a feature vector. The weights are based on the distance between words with the distance to the entity pair. The features are based on sentence-level and global information and take into account the entity pair.

3.4 Edge Weighting

We have syntactically parsed the sentence to get its dependency tree and define the edges between words. This parsing is very important for the relation extraction that is essentially based on the predicates (Mahendran et al., 2022; Li et al., 2022; Yuan et al., 2022). Most of the methods based on this idea use unweighted edges. Indeed, we have followed the method of Zhou et al. (2020) by adding weighted edges based on the shortest paths to enhance the dependency propagation made by the GCN layers. The method of Yuan et al. (2022) focuses on some parts of edges that are close to the entity pair to get the most relevant information. Based on this idea, we have modified the weights to take into account the proximity to the entity pair. In addition, we have taken into account the sequential distance between words by exploiting a standard metric (Liu, 2008). In order to compute the edge weights between nodes, we proceed as follows. First, we use the phrase to compute the distance between two words using their order of appearance in the same phrase (say i and j) using the Manhattan distance:

$$dSeq_{ij} = |i - j| \quad (1)$$

Thereafter, we use the dependency graph to compute another form of distance between nodes i and j by summing the distances $dSeq$ of the edges along the shortest path between i and j as follows:

$$d_{ij} = \sum_{k \in \Pi_{ij}} dSeq(src_k, dst_k) \quad (2)$$

where Π_{ij} is the shortest between nodes i and j ; src_k, dst_k are the source and destination for each edge $k \in \Pi_{ij}$; and $dSeq(\cdot)$ is the distance computed by equation 1. Doing this, we fuse between two important informations: sequence distance and dependency distance. Hence, given a node i , we compute the distance between the corresponding word and the existing entities in the relation as:

$$dEnt_i = \min_e d_{ie} \quad (3)$$

where e is the set of entity words. Now, we compute the weight between nodes i and j by the following formula:

$$w_{ij} = \frac{1}{e^{d_{ij} + \min(dEnt_i, dEnt_j) - 2}} \quad (4)$$

Note that we use the exponential to amplify the distance between nodes. We will describe in the next section, the features used for graph nodes.

3.5 Node Features

In order to describe the nodes in the dependency graph, we have adopted two categories of features judged to be of great importance for relation extraction: entity features and dependency features described by:

3.5.1 Word Embedding:

We have used a GloVe model pre-trained on clinical text to generate a static embedding of size 100 for each word (Flamholz et al., 2022). This embedding takes into account the global co-occurrence between words which is a significant factor in determining the relations (Shahab, 2017; Perera et al., 2020; Mahendran et al., 2022). Since we have fed all the input features into a BiLSTM layer, we can capture contextual information rather than using a contextual embedding like BERT (Mahendran et al., 2022; Ed-drissiya El-allaly et al., 2022).

3.5.2 Part of Speech:

The PoS is essential information to the relation extraction since it is highly dependent on grammatical properties (Xu et al., 2018; Li et al., 2022). Thus, we have used a trainable embedding layer of size $N_{pos} = 10$ to represent 45 fine-grained PoS tags such as singular noun, past participle verb, and determiner.

3.5.3 Entity Type:

Since we have to detect the relations between entities, the entity types are essential information for this task

(Patel et al., 2021; Yuan et al., 2022). Thus, we have used the entity types as lexical word features represented by a trainable embedding layer of size 10.

3.5.4 Dependency Type:

Most methods ignore the dependency types and consider that the different dependency relations are identical (Zhou et al., 2020; Li et al., 2022; Yuan et al., 2022). However, we have used a trainable embedding layer of size 10 to represent the type of the incoming dependency relation for each word. Thus, we are able to distinguish 40 types such as nominal subject, object predicate, particle, etc.

3.5.5 Matching a Relation Verb:

Relevant side information is usually ignored while it enables to benefit from knowledge about the relations (Yuan et al., 2022). To get information beyond the sentence, we have automatically collected verbs for each relation type from the training set. Thus, the feature vector indicates the types which have a verb matching the word. We were inspired by the post-processing rules of Chikka and Karlapalem (2018) to select a verb that connects a pair of entities. For that, the verb should appear in the shortest dependency path that relies these entities, otherwise it should be in sequence between them. Each verb is normalized by a stemming process.

3.5.6 Distance to the Entities:

Many methods have included information about the entity pair position in their features (Chikka and Karlapalem, 2018; Patel et al., 2021; Yuan et al., 2022). Since it is essential to know which words are supposed to belong to the relation entities. To represent the relative position of the subject and the object, we have used a trainable embedding layer of size 10 for each one. We have used 21 classes to represent each position where the direction and belonging to an entity are taken into account.

3.6 Graph Convolutional Network

In our work, we have used the GCN to perform deep learning on graph-structured data. The GCN layer represents the first-order neighborhood dependency by propagating the node features to the direct neighbor nodes. Usually, a larger number of GCN layers is required to represent higher order which increases the model complexity (Mahendran et al., 2022; Yuan et al., 2022). By adding the shortest path edges (Zhou et al., 2020), one layer is able to perform long-distance propagation, especially with edge weights.

In addition, we have used the mask pooling method as an entity-attention mechanism to resume the GCN output into three parts: subject entity, sentence, and object entity (Zhou et al., 2020). By this mechanism, the model is able to focus deeply on the relation entities during classification. The next section analyzes experimentally the performance of our proposal.

4 EXPERIMENTATION

We have chosen the benchmark dataset of the i2b2/VA 2010 challenge (Uzuner et al., 2011). This dataset contains patient reports which are divided into 170 for training and 256 for testing. It embeds sentences annotated by 8 types of relations that hold between 3 types of entities: medical problem, treatment, and test. More information is shown in Table 1.

Table 1: Information about relations in i2b2/VA 2010 dataset.

Entity types	Relation type	Train set	Valid set
Treatment-Problem	Treatment improves problem (TrIP)	107	198
	Treatment worsens problem (TrWP)	56	143
	Treatment causes medical problem (TrCP)	296	444
	Treatment is administered for medical problem (TrAP)	1 422	2 487
	Treatment is not administered because of problem (TrNAP)	106	191
Test-Problem	Test reveals medical problem (TeRP)	1 733	3 033
	Test conducted to investigate medical problem (TeCP)	303	588
Problem-Problem	Medical problem indicates medical problem (PIP)	1 239	1 986
Total		5 262	9 070

In our research, we used a BiLSTM hidden layer of size 64 and a WGCN layer of size 128, enabling effective residual computation. Our model consists of 2 WGCN layers with a dropout rate of 0.5 and was trained over 150 epochs with batch sizes of 50. The computations were carried out on a system with 8 GB of RAM and an Intel(R) Core(TM) i5-9300H CPU @ 2.40GHz processor. We implemented our work using the Python programming language with the Spacy module for natural language preprocessing and PyTorch for deep learning models.

For the evaluation, we have adapted the standard

metrics: precision, recall, and F1-score. For that, we use 9 classes: a "Non-relation" class and a class for each relation type. Each sample represents a sentence with a selected entity pair. In order to analyze the importance of each category of features, we have implemented different versions of our proposal as follows:

- GCN+WE: The GCN model with our edge weighting method and essential features.
- +DEP: Adding the dependency type features.
- +TYRE: Adding the relation verb features.
- +DIST: Adding the entity pair position features (full set of features).

The overall and the relation type-level results are shown in the tables 2 and 3, respectively. In the table 2, all components contribute to reaching high results since the minimal result is $\approx 81\%$. The added features contribute to enhancing the F1-score results by +0.71%. The best recall improvement is achieved by adding the dependency type features by +0.67%. Hence, the model can cover well the relations by distinguishing the different dependency types. The best precision improvement (+0.30%) was made by the entity distance features. Thus, recognizing the words of the relation entities is useful to precisely classify the relation.

In table 2, all the features contribute to increasing the results for almost all the relation types with an average improvement of +4% for 7 different classes. However, the TrWP relations have the lowest result (6.30%) due to the lack of data since there are only 56 training samples for this type. After adding our features, the result for this type is improved by +4.48% and especially after adding the relation verbs features. Thus, our model benefits from the automatically constructed dictionary to cover well the different relation types and to deal with the lack of data.

Table 2: Overall evaluation of the GREED components.

Model	Precision	Recall	F1-score
GCN+WE	81.87%	81.83%	81.38%
+DEP	81.77%	82.50%	81.97%
+TYRE	81.99%	82.67%	81.97%
+DIST	82.29%	82.40%	82.09%

In order to assess the effectiveness of GREED, we have compared it with four other recent models: BiLSTM (Li et al., 2019), WGCN (Zhou et al., 2020), LSTM+R (Chikka and Karlapalem, 2018), and CNN (Patel et al., 2021). The overall and the type-level results of all models are shown in tables 4 and 5,

Table 3: Evaluation of the GREED components in type-level according to F1-score. The "No relation" class means that there is no defined relation between the entity pair.

Model	No relation	TrWP	TrAP	TrNAP	TrCP	TrIP	TeCP	TeRP	PIP
GCN+WE	88.90%	01.82%	73.63%	26.09%	53.37%	26.40%	52.48%	84.14%	43.26%
+DEP	89.27%	00.00%	74.53%	16.90%	57.31%	27.72%	55.87%	85.39%	44.19%
+TYRE	89.32%	06.61%	74.87%	32.94%	57.09%	29.17%	51.34%	85.19%	42.77%
+DIST	89.21%	06.30%	75.26%	36.00%	58.10%	29.17%	49.73%	83.43%	47.58%

Table 4: Overall comparison of GREED with state-of-the-art models.

Model	Method	Precision	Recall	F1-score
BiLSTM (Li et al., 2019)	ET	75.69%	73.03%	74.34%
WGCN (Zhou et al., 2020)		80.94%	82.44%	81.14%
GREED		82.29%	82.40%	82.09%
WGCN (Zhou et al., 2020)	T	86.20%	87.91%	86.60%
CNN (Patel et al., 2021)		75.00%	72.00%	74.00%
GREED		87.88%	89.08%	87.72%

Table 5: Comparison of GREED with state-of-the-art models according to type-level F1-score.

Model	Method	TrWP	TrAP	TrNAP	TrCP	TrIP	TeCP	TeRP	PIP
BiLSTM (Li et al., 2019)	ET	44.57%	79.74%	42.27%	62.13%	61.59%	61.17%	84.44%	63.33%
WGCN (Zhou et al., 2020)		00.00%	72.72%	30.28%	49.02%	34.07%	42.13%	81.73%	40.85%
GREED		06.30%	75.26%	36.00%	58.10%	29.17%	49.73%	83.43%	47.58%
LSTM+R (Chikka and Karlapalem, 2018)	T	58.00%	47.00%	42.00%	30.00%	17.00%	NA	NA	NA
WGCN (Zhou et al., 2020)		07.35%	88.55%	33.20%	65.73%	43.39%	55.81%	93.63%	100.0%
CNN (Patel et al., 2021)		16.66%	78.54%	17.10%	04.00%	28.57%	46.66%	83.95%	92.43%
GREED		11.11%	89.02%	38.16%	65.64%	37.40%	68.20%	95.36%	99.97%

respectively. The abbreviation "ET" in these tables means that the model should extract the relation before identifying its type; the abbreviation "T" means that we have only to identify the type of relation between a pair of related entities; while the abbreviation "NA" means Not Available. In table 4, our GREED model outperforms all the state-of-the-art models according to all metrics. WGCN (Zhou et al., 2020) provides the closest results to ours with F1-score difference of -0.95% and -1.12% according to "ET" and "T" methods, respectively. Note that this is the only model based on GCN to process weighted dependency graphs and it has not been applied in the medical field before (Zhou et al., 2020). However, by our features and edge weighting method, GREED is able to achieve higher results. These two GCN-based models have significantly outperformed the others with a difference of more than $+6.8$ according to the overall F1-score.

We can see in Table 5 that our proposal GREED is more stable than the rest of the models and provides the best results in most cases for all relation types. However, GREED is outperformed by the LSTM+R (Chikka and Karlapalem, 2018) model in the TrWP relation type with a difference of -46.89% , while we significantly outperform this model by an average of $+32.68\%$ on three other types: TrAP, TrCP, and TrIP. Note that LSTM-R used only 5 classes to classify "treatment-problem" relations. This reduces the confusion to focus more on the TrWP class, but the results are significantly decreased for other classes. We have outperformed the CNN (Patel et al., 2021) model with an average of $+20.35\%$ on 7 relation classes. Indeed, CNN provides higher results than GREED for the TrWP relations, with a difference of $+5.55\%$. In fact, this class has the lowest number of training samples (56), but CNN benefited from a big manually constructed meta-thesaurus called UMLS (Bo-

denreider, 2004) to make lexical features. Although we don't make any manual effort to construct our dictionary, we significantly outperform LSTM+R in the other classes. We have outperformed the two versions of WGCN on all the classes, especially with the "ET" method, with an average increase of +5.66%. WGCN provides a result of 0% for the TrWP class while we obtain a result of 6.30%. Thus, our added features and edge weighting, and pre-processing technique allow GREED to cover more relations. Although we outperformed BiLSTM (Li et al., 2019) by +7.75% in terms of overall F1-score, this model achieved the highest results on all relation type classes. Note that BiLSTM uses the "non-relation" class but its result is not available. Thus, this class must have a very low result and this means that many entity pairs without relations are misclassified. This model uses dependency relations and their types with CNN and BiLSTM layers. Thus, we can conclude that the use of GREED of this information with GCN leads to higher overall results without affecting the "non-relation" class.

5 CONCLUSION

In this paper, we have proposed a hybrid method named GREED (Graph learning based Relation Extraction with Entity and Dependency relations) which is able to extract clinical relations between entities in sentences by using GCN on dependency graphs. Our model processes the dependency relations efficiently by appropriately weighting and filtering the edges, taking into account the entity pair. GREED outperforms four state-of-the-art models especially which are not based on graphs, without the need for a big manual effort. Moreover, it can deal with the lack of data. However, the time complexity of extracting relations in one sentence may be unreasonable since every possible entity pair needs to be processed. For that, we need to find a way to extract all relations in a sentence with only one process.

REFERENCES

- Aronson, A. R. and Lang, F.-M. (2010). An overview of metamap: historical perspective and recent advances. *Journal of the American Medical Informatics Association*, 17(3):229–236.
- Ben Abdesslem Karaa, W., Alkhamash, E. H., and Bchir, A. (2021). Drug disease relation extraction from biomedical literature using nlp and machine learning. *Mobile Information Systems*, 2021.
- Bodenreider, O. (2004). The unified medical language system (umls): integrating biomedical terminology. *Nucleic acids research*, 32(suppl_1):D267–D270.
- Chikka, V. R. and Karlapalem, K. (2018). A hybrid deep learning approach for medical relation extraction. *arXiv preprint arXiv:1806.11189*.
- Ed-drissiya El-allaly, Sarrouti, M., En-Nahnahi, N., and Ouatik El Alaoui, S. (2022). An attentive joint model with transformer-based weighted graph convolutional network for extracting adverse drug event relation. *Journal of Biomedical Informatics*, 125:103968.
- Flamholz, Z. N., Crane-Droesch, A., Ungar, L. H., and Weissman, G. E. (2022). Word embeddings trained on published case reports are lightweight, effective for clinical tasks, and free of protected health information. *Journal of Biomedical Informatics*, 125:103971.
- Hearst, M. A., Dumais, S. T., Osuna, E., Platt, J., and Scholkopf, B. (1998). Support vector machines. *IEEE Intelligent Systems and their applications*, 13(4):18–28.
- Kim, Y., Heider, P. M., Lally, I. R., and Meystre, S. M. (2021). A hybrid model for family history information identification and relation extraction: Development and evaluation of an end-to-end information extraction system. *JMIR Medical Informatics*, 9(4):e22797.
- Kong, H.-J. (2019). Managing unstructured big data in healthcare system. *Healthcare informatics research*, 25(1):1–2.
- Landolsi, M. Y., Hlaoua, L., and Ben Romdhane, L. (2022a). Information extraction from electronic medical documents: state of the art and future research directions. *Knowledge and Information Systems*, pages 1–54.
- Landolsi, M. Y., Romdhane, L. B., and Hlaoua, L. (2022b). Medical named entity recognition using surrounding sequences matching. *Procedia Computer Science*, 207:674–683.
- Li, T., Ma, L., Qin, J., and Ren, W. (2022). Dtcgn: a method combining dependency tree and graph convolutional networks for chinese long-interval named entity relationship extraction. *Journal of Ambient Intelligence and Humanized Computing*, pages 1–13.
- Li, Z., Yang, Z., Shen, C., Xu, J., Zhang, Y., and Xu, H. (2019). Integrating shortest dependency path and sentence sequence into a deep learning framework for relation extraction in clinical text. *BMC medical informatics and decision making*, 19(1):1–8.
- Liu, H. (2008). Dependency distance as a metric of language comprehension difficulty. *Journal of Cognitive Science*, 9(2):159–191.
- Luo, Y., Uzuner, Ö., and Szolovits, P. (2017). Bridging semantics and syntax with graph algorithms—state-of-the-art of extracting biomedical relations. *Briefings in bioinformatics*, 18(1):160–178.

- Mahendran, D., Tang, C., and McInnes, B. (2022). Graph convolutional networks for chemical relation extraction. *Proceedings of the Semantics-enabled Biomedical Literature Analytics (SeBiLAN)*.
- Patel, R., Tanwani, S., and Patidar, C. (2021). Relation extraction between medical entities using deep learning approach. *Informatica*, 45(3).
- Perera, N., Dehmer, M., and Emmert-Streib, F. (2020). Named entity recognition and relation detection for biomedical information extraction. *Frontiers in cell and developmental biology*, page 673.
- Quevedo, X. and Chicaiza, J. (2023). Using nlp to enrich scientific knowledge graphs: A case study to find similar papers. In *ICAART (2)*, pages 190–198.
- Shahab, E. (2017). A short survey of biomedical relation extraction techniques. *arXiv preprint arXiv:1707.05850*.
- Uzuner, Ö., South, B. R., Shen, S., and DuVall, S. L. (2011). 2010 i2b2/va challenge on concepts, assertions, and relations in clinical text. *Journal of the American Medical Informatics Association*, 18(5):552–556.
- Xu, J., Gan, L., Cheng, M., and Wu, Q. (2018). Unsupervised medical entity recognition and linking in chinese online medical text. *Journal of healthcare engineering*, 2018.
- Yang, X., Yu, Z., Guo, Y., Bian, J., and Wu, Y. (2021). Clinical relation extraction using transformer-based models. *CoRR*, abs/2107.08957.
- Yuan, C., Huang, H., Feng, C., and Cao, Q. (2022). Piecewise graph convolutional network with edge-level attention for relation extraction. *Neural Computing and Applications*, pages 1–13.
- Zhou, L., Wang, T., Qu, H., Huang, L., and Liu, Y. (2020). A weighted gcn with logical adjacency matrix for relation extraction. In *ECAI 2020*, pages 2314–2321. IOS Press.