

Data and text mining

DeepEventMine: End-to-end Neural Nested Event Extraction from Biomedical Texts

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Associate Editor: XXXXXXX

Received on XXXXX; revised on XXXXX; accepted on XXXXX

Abstract

Motivation: Recent neural approaches on event extraction from text mainly focus on flat events in general domain, while there are less attempts to detect nested and overlapping events. These existing systems are built on given entities and they depend on external syntactic tools.

Results: We propose an end-to-end neural nested event extraction model named DeepEventMine that extracts multiple overlapping directed acyclic graph structures from a raw sentence. On the top of the Bidirectional Encoder Representations from Transformers (BERT) model, our model detects nested entities and triggers, roles, nested events and their modifications in an end-to-end manner without any syntactic tools. Our DeepEventMine model achieves the new state-of-the-art performance on seven biomedical nested event extraction tasks. Even when gold entities are unavailable, our model can detect events from raw text with promising performance.

Availability and implementation: Our codes and models to reproduce the results are available at: <https://github.com/aistairc/DeepEventMine>

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Supplementary information: Supplementary data are available at *Bioinformatics* online.

1 Introduction

Biomedical information is scattered in the substantial body of literature, so natural language processing (NLP) techniques are required to extract such underlying information without manual efforts. Many studies have been carried out to develop biomedical information extraction (IE) techniques, and, in these studies, extracting events among entities from text is considered as one of the most important tasks (Kim *et al.*, 2011b; Pyysalo *et al.*, 2011; Kim *et al.*, 2013). Event structures are key to reveal the underlying knowledge and to capture biological processes described in the unstructured text. Events capture dynamic *n*-ary relations among entities (e.g., proteins and genes) and their attributes (e.g., locations and sites). They can facilitate the development of practical applications such as information retrieval (Tsuruoka *et al.*, 2008; Van Landeghem *et al.*, 2013), knowledge base enrichment (Hakala *et al.*, 2012) and pathway curation (Miwa *et al.*, 2013a).

An event consists of a trigger and zero or more arguments.¹ A trigger is a textual mention that denotes the presence of an event in text. Each argument is either an entity or an event, and it plays a role that characterises its contribution to the event. An event is called nested event (Pyysalo *et al.*, 2011; Miwa and Ananiadou, 2013) when it has other events in its arguments, while an event is called flat event when there are only entities in its arguments. Figure 1 illustrates an example of the flat and nested events, as well as the nested triggers and entities. There is a flat event *Cell transformation* (“transformed”) in this example. The flat event has a *Theme* argument from *Cell* entity (“erythroid cells”). The *Positive regulation* (“transformed”) is a nested event at level 1² with the *Cell transformation* event as its *Theme* and the *Organism* entity (“*erbA/myb IRES*”) as its

¹ In this work, all our target events have triggers.

² We define the level *k* of a nested event. *k* = 1 means the event has flat events in its arguments and *k* = 2 means the event has at least one nested event at level 1 as its argument.

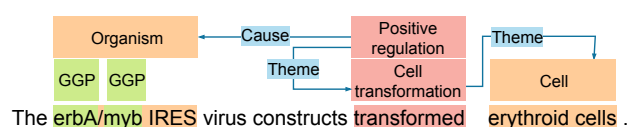


Fig. 1. Example flat and nested events from the BioNLP-ST 2013 Cancer Genetics task. Green and orange denote entities, while red are event triggers, e.g., *erbA*, erythroid cells are two entities and *transformed* is a trigger. Blue connection denotes the role of an argument to a trigger, where an argument can be an entity or a trigger, e.g., erythroid cells is a Theme of the trigger *transformed*. Event structures are constructed by an event trigger and its arguments. The flat event, *transformed* erythroid cells, has only one entity as its argument. The nested event, *erbA/myb IRES* transformed erythroid cells, has the entity *erbA/myb IRES* as Cause and the flat event *transformed* erythroid cells as its argument. The nested event is a tree whose root is the red rectangle (Positive regulation) with blue connections to its entity argument (Organism) and its flat event argument (*transformed* erythroid cells).

Cause. In this example, there are two triggers that share the same textual span “*transformed*”. We call these triggers nested triggers, which allow us to deal with multiple events indicated by the same phrase. Similarly, this example also involves nested or overlapping entities; the *Organism* entity contains two *GGP* (*Gene or gene product*) entities (“*erbA*” and “*myb*”). Nested events have this complex schema that allows multiple, overlapping, directed acyclic graph (DAG) structures, because they are essential to deeply understand texts written in condensed styles that often appear in scientific and technical domains such as bio-medicine.

Many neural models have been proposed for event extraction and have shown to produce better results than traditional feature-based models, e.g., (Miwa *et al.*, 2012a; Li *et al.*, 2013; Yan and Wong, 2019); however, most efforts have been dedicated to extracting flat events on flat entities in general domain, e.g., (Yang and Mitchell, 2016; Nguyen *et al.*, 2016; Liu *et al.*, 2018; Sha *et al.*, 2018; Nguyen and Nguyen, 2019), rather than nested entities (Ju *et al.*, 2018; Katiyar and Cardie, 2018; Sohrab and Miwa, 2018). There are few neural-based models (Björne and Salakoski, 2018; Li *et al.*, 2019) for nested events, even though they are important to capture relationships between events such as causality, which is essential to represent biological processes. Furthermore, most event extraction models assume the entities are given, so we need to prepare an external named entity recognizer before extracting events from raw texts. This leads to potential error propagation in the entity-event pipeline, which requires additional tuning. Especially when the argument entities are nested, we also need to consider a nested named entity recognizer (Finkel and Manning, 2009; Katiyar and Cardie, 2018; Ju *et al.*, 2018; Sohrab and Miwa, 2018; Wang and Lu, 2018). Although there are a few end-to-end models (Yang and Mitchell, 2016; Nguyen and Nguyen, 2019) to extract flat events on flat entities, none of these models can treat nested events on nested entities that may further overlap with event triggers.

Deep models pre-trained on large-scale corpora have achieved impressive results on several NLP tasks. Such models include Embeddings from Language Models (ELMo) (Peters *et al.*, 2018), OpenAI Generative Pre-Training (GPT) (Radford *et al.*, 2018), and Bidirectional Encoder Representations from Transformers (BERT) (Devlin *et al.*, 2019). High performance NLP systems have been built by stacking simple task-specific architecture on these models. Such models can also be helpful in modeling complex structures like events.

We propose a novel neural model named DeepEventMine that extracts nested events from a sentence in an end-to-end manner. We build a relatively simple but representative model that captures nested event structures, relying on the highly expressive power of BERT. Taking BERT as the base representation, the model performs a flow of entity and trigger detection, role detection, and event and their modification (e.g., negation and speculation) detection in an end-to-end manner. The state-of-the-art event extraction models depend on external syntactic tools (Björne and

Salakoski, 2018; Nguyen and Nguyen, 2019), which leads to domain dependent models. Unlike these existing models, our model does not rely on any external syntactic tools and this allows the application of our model to several domains. We fine-tune the SciBERT model (Lee *et al.*, 2019) that is pre-trained with a large scale biomedical and computer science texts while simultaneously training the event extraction model. We evaluated our model on seven biomedical nested event extraction data sets: Cancer Genetics 2013, Epigenetics and Post-translational Modifications 2011, GENIA 2011, GENIA 2013, Infectious Diseases 2011, Pathway Curation 2013, and Multi-Level Event Extraction. The experimental results show that our model can detect triggers, roles, events, and their modifications with better performance than existing models by a large margin. The end-to-end training makes the model focus more on event detection than the pipeline training. As a result, our model achieved the new state-of-the-art (SOTA) on the test sets of all seven tasks when given gold (manually annotated) entities. We also show that our model can detect the events from raw sentences with a slight performance loss.

2 Related Work

Deep neural networks including recurrent and convolutional neural networks have boosted event extraction performance, e.g., (Björne and Salakoski, 2018; Nguyen and Nguyen, 2019). These models show better performance than traditional hand-crafted feature-based approaches (Björne and Salakoski, 2013; Miwa and Ananiadou, 2013; Yang and Mitchell, 2016). While pre-trained models for fine-tuning contextual representations are recently shown to perform well on several NLP tasks (Peters *et al.*, 2018; Radford *et al.*, 2018; Devlin *et al.*, 2019), they have not been taken into consideration to build an event extraction model.

Previous neural models on flat event extraction have been mainly focused on event trigger and argument detection (Chen *et al.*, 2015; Nguyen *et al.*, 2016; Liu *et al.*, 2018; Sha *et al.*, 2018). They assume that entities are given, therefore, a separate entity recognition module is required to extract events from plain text. Due to the independence between the event model and the entity detection, information is not shared among them. Nguyen and Nguyen (2019) introduced an end-to-end neural model utilizing a bi-directional recurrent neural network and they addressed the independent issue of entity detection. However, they only extract flat events with flat entity arguments and rely on several syntactic features, so the application is limited to general domain and it cannot be applied to biomedical event extraction tasks. Meanwhile, our model can deal with both flat and nested entities and it is independent of external syntactic tools.

Contrary to flat event extraction, fewer attempts have been made to investigate nested event detection. Early work mainly relied on hand-crafted features as well as gold entities (Björne and Salakoski, 2013; Miwa and Ananiadou, 2013). Björne and Salakoski (2018) presented a pipeline-based neural system that has been developed for nested event extraction. The system performs event extraction by tackling four consecutive sub-tasks: detecting entities/triggers, relations, events, and event modifiers. All modules share a unified sentence representation, which is based on a convolutional neural network. Their approach relies on syntactic information, e.g., part-of-speech tags and dependencies, to encode a sentence. In addition, they ignore the dependencies among entities, relations and events due to independent detection. In contrast, our model enables rich semantic representations shared among sub-tasks and errors can be addressed through the entire task sequence.

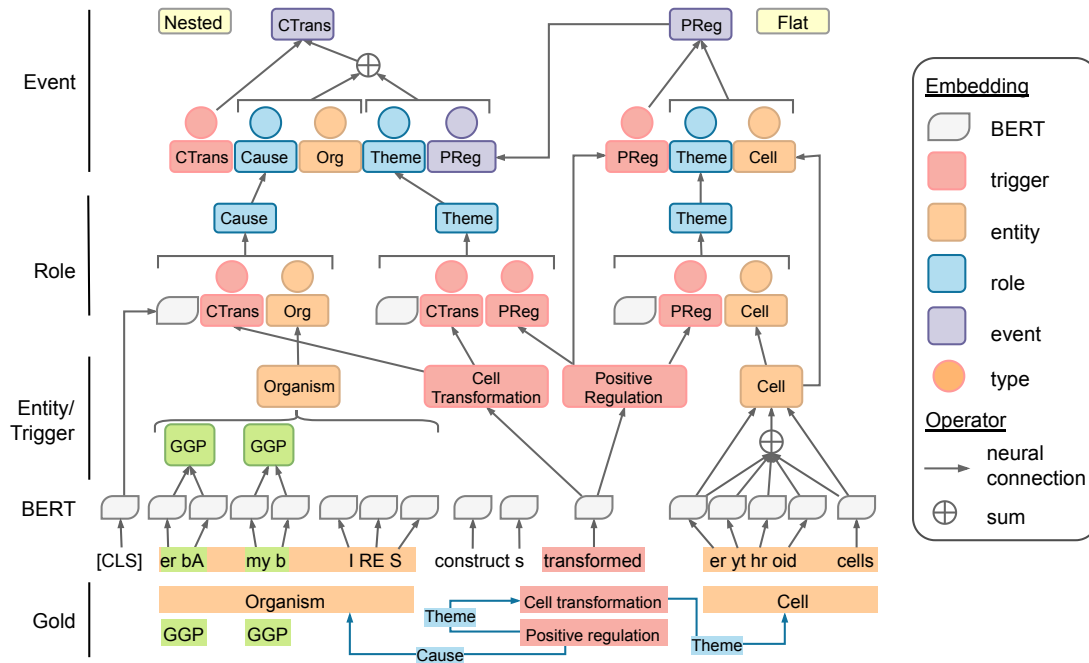


Fig. 2. Overview of our deep end-to-end event extraction model. The hidden layers are omitted for brevity. We use the example in Figure 1 to present our method. Due to the space limitation, we omit some unimportant tokens (e.g., the, virus, and .). We use abbreviations to represent triggers and entities, i.e., CTrans, PReg stand for Cell Transformation and Positive Regulation triggers, respectively, whilst Org and GGP are the short forms of Organism and Gene or gene product entities, respectively. We sum the representations of the tokens and concatenate with the first and last tokens to form the representation of a span. We show the combination details of an entity erythroid cells, while those of the others are omitted for simplicity.

3 DeepEventMine

Our DeepEventMine model consists of four layers: BERT, entity/trigger, role, and event layers. We illustrate the overview of our model in Figure 2. As shown in this figure, the representations of textual spans built on the outputs of the BERT layer are shared and updated by the subsequent layers.

3.1 BERT Layer

The BERT layer receives sub-word sequences and assigns contextual representations to the sub-words via BERT (Devlin *et al.*, 2019). We assume each sentence S has n words and the i -th word is split into s_i sub-words. This layer assigns a vector $v_{i,j}$ to the j -th sub-word of the i -th word. It also produces the representation v_S for the sentence, which corresponds to the [CLS] embedding in Devlin *et al.* (2019).

3.2 Entity/Trigger Layer

The entity/trigger layer assigns entity or trigger types to overlapping text spans, or word sequences, in a sentence. We employ the modeling of Sohrab and Miwa (2018) since this enables the extraction of nested triggers and entities at once. This model has the advantage that it can utilize all the sub-words' information inside the target candidate unlike the NER model of Devlin *et al.* (2019) that only uses the first sub-words of words. Particularly, this layer exhaustively considers all possible spans of a sentence with the sizes less than or equal to the maximum span size L_{ent} for entities and L_{trig} for triggers.³

The representation $\mathbf{m}_{k,l} \in \mathbb{R}^{d_m}$ for the span from the k -th word to the l -th word in a sentence is calculated as follows:

$$\mathbf{m}_{k,l} = \left[\mathbf{v}_{k,1}; \frac{\sum_{i=k}^l \sum_{j=1}^{s_i} \mathbf{v}_{i,j}}{\sum_{i=k}^l s_i}; \mathbf{v}_{l,s_l} \right], \quad (1)$$

³ L_{ent} and L_{trig} are tuned as hyper-parameters and presented in Appendix B.

where $[\cdot; \cdot]$ denotes concatenation, while $\mathbf{v}_{i,j}$ denotes the j -th sub-word representation in the i -th word. More specifically, $\mathbf{v}_{k,1}$ is the first sub-word representation of the k -th word, while \mathbf{v}_{l,s_l} is the last sub-word representation of the l -th word. Since each span may have multiple types, we solve this multi-label problem by using multiple binary classifiers with sigmoid functions. In particular, the representation $\mathbf{m}_{k,l}$ is passed to multiple binary classifiers corresponding to all possible types. For each classifier, the span is selected as the corresponding entity/trigger type if the output score is greater than 0.5.

3.3 Role Layer

The role layer enumerates all trigger-argument pairs (trigger-trigger and trigger-entity pairs) given triggers and entities detected by the entity/trigger layer and assigns a role type or no role to each pair.

Since each role is constructed by a trigger and an argument, we firstly compute representations of all triggers and arguments detected by the entity/trigger layer. The representations of a trigger and an argument are calculated in the same way. A trigger t ranging from the starting t_s -th word to the ending t_e -th word is represented with the concatenation of its span representation $\mathbf{m}_t (= \mathbf{m}_{t_s,t_e}$ from Eq. (1)) and an n_t -dimensional type embedding \mathbf{s}_t , as follows:

$$\mathbf{v}_t = [\mathbf{m}_t; \mathbf{s}_t]. \quad (2)$$

Similarly, the representation of an argument a can be calculated as:

$$\mathbf{v}_a = [\mathbf{m}_a; \mathbf{s}_a]. \quad (3)$$

Meanwhile, the context representation \mathbf{c} is obtained from the sentence representation of the BERT layer, i.e., $\mathbf{c} = \mathbf{v}_S$. We empirically investigated the utility of the context and found that the context contributed to the performance.⁴

⁴ The detailed scores are presented in Appendix E.

The representation $\mathbf{r}_i \in \mathbb{R}^{d_r}$ for a pair i is then calculated from its trigger representation \mathbf{v}_t , argument representation \mathbf{v}_a , and context representation \mathbf{c} :

$$\mathbf{r}_i = \text{GELU}(\mathbf{W}_r [\mathbf{v}_t; \mathbf{v}_a; \mathbf{c}] + \mathbf{b}_r), \quad (4)$$

where \mathbf{W}_r and \mathbf{b}_r are learnable weights and biases respectively and GELU is the Gaussian Error Linear Unit (GELU) activation function (Hendrycks and Gimpel, 2016). After getting the pair representation \mathbf{r}_i , we pass it to a fully connected layer with a softmax function to predict the corresponding role type.

3.4 Event Layer

Given the triggers and entities detected by the entity/trigger layer and the role pairs detected by the role layer, the event layer enumerates all legal combinations of role pairs⁵ to construct event candidates for each trigger. These event candidates include events with no arguments. Each candidate is then classified as an event or non-event. Additionally, this layer detects event modification such as speculation or negation.

To allow the construction of nested events, we build events in a bottom-up manner. Concretely, we first classify event candidates with non-trigger arguments (containing no argument or a set of entity arguments only) to obtain a set of detected (flat) events. We then repeat the following process of constructing and classifying nested event candidates, which contain trigger arguments. In the construction step, we construct nested event candidates by replacing trigger arguments by their corresponding detected events. In the case there are multiple detected events linked to a trigger argument, we create (or duplicate) an event candidate corresponding to each of them. The nested event candidates are classified and the detected events may serve as arguments for the higher nested-level nested event candidates. The process is repeated until all the event candidates are classified or no event candidate with fixed arguments remains.

The representation $\mathbf{e}_i \in \mathbb{R}^{d_e}$ of an event candidate i is calculated from the representations of the trigger t by Eq. (2), the role r by Eq. (4), the n_t -dimensional embedding of the role type u , and all the arguments $a_* \in e_a$ by Eq. (3) as follows:

$$\mathbf{e}_i = \left[\mathbf{v}_t; \sum_{j \in e_a} \text{GELU}(\mathbf{W}_p [\mathbf{r}_j; \mathbf{u}_j; \mathbf{v}_{a_j}] + \mathbf{b}_p) + \sum_{j \notin e_a} \text{GELU}(\mathbf{W}_n [\mathbf{r}_j; \mathbf{u}_j; \mathbf{v}_{a_j}] + \mathbf{b}_n) \right], \quad (5)$$

where \mathbf{W}_p and \mathbf{b}_p are learnable weights and biases respectively for roles that are in the event structure, meanwhile, \mathbf{W}_n and \mathbf{b}_n are learnable weights and biases for roles that are detected by the role layer and share the trigger but are not in the event structure. We include not only roles in the structure but also those outside of the structure, aiming at choosing the appropriate set of roles from all possible role sets. When the event candidate has a trigger with a fixed event as its argument, we replace \mathbf{v}_{a_j} by the event candidate representation \mathbf{e}_{a_j} in Eq. (5) after adjusting the dimensions via a hidden layer with GELU. The representation \mathbf{e}_i is passed to a hidden layer with GELU to produce \mathbf{e}'_i , and then \mathbf{e}'_i is fed to a binary classifier to classify as an event or not. When the candidate event is classified as an event, \mathbf{e}'_i is additionally passed to a softmax classifier to attach event modification types to the detected event.

4 Experimental Settings

4.1 Data and Task Settings

We evaluated our model on the six event extraction shared task data sets in biomedical domain: Cancer Genetics 2013, Epigenetics and

Post-translational Modifications 2011 (Ohta et al., 2011) (EPI), GENIA 2011 (Kim et al., 2011b) (GE11), GENIA 2013 (Kim et al., 2013) (GE13), Infectious Diseases 2011 (Pyysalo et al., 2011) (ID), and Pathway Curation 2013 (Pyysalo et al., 2015) (PC). We also evaluated it on Multi-Level Event Extraction (Pyysalo et al., 2012) (MLEE), a major event extraction data set. All the data sets include nested events and the data statistics are presented in Appendix A.

For all of the data sets, we used the data partition provided by the shared tasks or by the data provider for MLEE. We followed the official task evaluation settings for the tasks, which are originally from the setting in the BioNLP 09 shared task (Kim et al., 2011a). In particular, an extracted trigger span can be differed from the gold span by a single word (approximate span matching) and a sub-event are treated as a valid argument if the *theme* arguments of the sub-event are correctly detected (approximate recursive matching).

For the six event extraction tasks, we evaluated using the official online evaluation provided by the BioNLP11 and BioNLP13 shared tasks for both development and test sets except for CG and PC development sets, where the online evaluation are not provided. We use the same evaluation criteria as in Björne and Salakoski (2018). Specifically, for CG, EPI, ID, and PC tasks, we apply the primary evaluation criteria for full task results. For GE11 and GE13 tasks, we apply the approximate span/approximate recursive for the whole data (i.e., Task 1: full papers + abstract).

For MLEE task, we follow the evaluation setting of the MLEE provider (Pyysalo et al., 2012), which adapts the evaluation protocol and tools from the BioNLP'09 shared task (Kim et al., 2011a). Specifically, we apply the primary matching criteria, in which event structures are identical but include the approximate span and approximate recursive relaxations.

4.2 Training Settings

We train the model in an end-to-end manner based on the pre-trained BERT model. We employed the pre-trained SciBERT model (Beltagy et al., 2019), which was pre-trained on large-scale biomedical text. For pre-trained models on biomedical text, there are several existing models such as the BioBERT (Lee et al., 2019) and SciBERT models. We have empirically compared the BioBERT and SciBERT models on the CG development set and found that we obtained better performance when using the SciBERT model.⁶ Due to the large size of the pre-trained BERT, we employed multiple GPUs in the AI Bridging Cloud Infrastructure (ABCI)⁷ for speeding up.

Our model was implemented with PyTorch (Paszke et al., 2017). We employed the BERT model from the PyTorch Pretrained BERT repository⁸ as our BERT layer and implemented the end-to-end event extraction layers on the top of it. For all the data sets, we split the sentences into words by spaces and we further split each word into sub-words, which is a prerequisite for the input of BERT, following the setting of the BERT tokenizer.

Our training consists of two steps. In the first step, we warm up each layer separately by training each layer using gold annotations in order to avoid using unreliable predictions in the subsequent layers and to fully make use of gold annotations. At the second step, we train the entire model including entity/trigger, role and event layers simultaneously. The training loss is the summation of the losses from all the layers. We use corresponding binary or categorical cross entropy loss functions as the learning objectives for the binary or softmax classifiers in each layer.

⁶ The detailed scores of the comparison with using BioBERT are presented in Appendix D.

⁷ <https://abci.ai/>

⁸ <https://github.com/huggingface/pytorch-pretrained-BERT/tree/34cf67fd6c>

⁵ We build templates from the event structure definition.

We train the model with the Adam optimizer (Kingma and Ba, 2015). We include gradient clipping, dropout, and L2 regularization. The model is trained for 100 epochs, with the training mini-batch size as 16. Dropout is employed to the input of each layer: entity/trigger, role and event layers. All the above hyper-parameters were tuned with the Optuna framework.⁹ The model hyper-parameters used in our experiments are listed in Appendix B.

4.3 Models in Comparison

We compared our model with the SOTA models on the test sets. We briefly describe the models here.

- **TEES-CNN** (Björne and Salakoski, 2018) (SOTA on CG and GE11 tasks) – a pipeline system that detects events by sequentially performing entity, argument and event extraction. Each module shares a convolutional neural network (CNN)-based sentence encoder architecture.
- **BioMLN** (Venugopal *et al.*, 2014) (SOTA on GE13 task) – this is a markov logic network-based joint model that is built on multiple feature-based event extraction modules.
- **EventMine** (Miwa *et al.*, 2013b) (SOTA on ID task) – a pipeline system using hand-crafted lexical and syntactic features that employs a domain adaptation method and trained on seven event corpora.
- **PMCNN** (Li *et al.*, 2018) (SOTA on MLEE task) – a parallel multi-pooling CNN model to represent compositional semantic features of a sentence. We also compared with two recent neural-based models: **MultiRep-CNN** (Wang *et al.*, 2017) and **BLSTM** (He *et al.*, 2019).

For our models, we implemented the two following settings:

- **DeepEventMine** – Our end-to-end model that is able to predict events in one pass from raw text. The model is simultaneously trained for entity/trigger, role, and event detection.
- **Pipeline** – Pipeline model where each layer is trained separately. This is prepared to show the effectiveness of our end-to-end approach.

We also show performance gains when ensembling five versions (same as Björne and Salakoski (2018)) of our DeepEventMine with different parameter initialization. We employed a simple majority voting method, in which an event is predicted only when it appears in the outputs of at least three of the five models.

5 Results and Discussion

We present the performance of our DeepEventMine model on the seven corpora and compare it with the SOTA models on the test sets given gold entities. We also conduct analyses to compare the end-to-end and pipeline models, and evaluate the extraction from raw text and text with gold entities. We finally deeply analyze the results using the CG data set. Since the gold data of the test sets are not provided by the tasks, and also to avoid overfitting on the test sets, the analyses are conducted on the development sets.

5.1 Comparison with the state-of-the-art

Table 1 compares our model with the SOTA models on the seven tasks. For the shared tasks, the test evaluation can only be performed through the shared task website and all submissions are required to use gold entities. We use the same setting for MLEE, which uses gold entities. Overall, our DeepEventMine model consistently outperforms the SOTA models in all

Table 1. Results of event extraction on the test sets given gold entities.

Task	Model	P	R	F (%)
CG	TEES-CNN (Björne and Salakoski, 2018)	66.55	50.77	57.60
	DeepEventMine (single)	69.54	54.24	60.94
	DeepEventMine (ensemble)	72.23	53.92	61.74
EPI	EventMine (Miwa <i>et al.</i> , 2013b)	54.42	54.28	54.35
	TEES-CNN (Björne and Salakoski, 2018)	64.93	50.00	56.50
	DeepEventMine (single)	73.73	55.95	63.62
	DeepEventMine (ensemble)	78.34	56.39	65.57
GE11	EventMine (Miwa <i>et al.</i> , 2012b)	63.48	53.35	57.98
	BioMLN (Venugopal <i>et al.</i> , 2014)	63.61	53.42	58.07
	TEES-CNN (Björne and Salakoski, 2018)	69.45	49.94	58.10
	DeepEventMine (single)	71.71	56.20	63.02
	DeepEventMine (ensemble)	76.28	55.06	63.96
GE13	TEES-CNN (Björne and Salakoski, 2018)	65.78	44.38	53.00
	BioMLN (Venugopal <i>et al.</i> , 2014)	59.24	48.95	53.61
	DeepEventMine (single)	60.98	49.80	54.83
	DeepEventMine (ensemble)	67.08	49.14	56.72
ID	TEES-CNN (Björne and Salakoski, 2018)	66.48	50.66	57.50
	EventMine (Miwa <i>et al.</i> , 2013b)	61.33	58.96	60.12
	DeepEventMine (single)	63.56	57.30	60.27
	DeepEventMine (ensemble)	68.51	55.99	61.62
PC	EventMine (Miwa and Ananiadou, 2013)	53.48	52.23	52.84
	TEES-CNN (Björne and Salakoski, 2018)	62.16	50.34	55.62
	DeepEventMine (single)	64.12	49.19	55.67
	DeepEventMine (ensemble)	68.13	50.07	57.72
MLEE	MultiRep-CNN (Wang <i>et al.</i> , 2017)	60.56	56.23	58.31
	PMCNN (Li <i>et al.</i> , 2018)	67.23	53.61	59.65
	BLSTM (He <i>et al.</i> , 2019)	–	–	59.61
	DeepEventMine (single)	67.39	56.35	61.38
	DeepEventMine (ensemble)	69.91	55.49	61.87

the seven tasks. Specifically, our DeepEventMine model achieved 1.5 to 5.2 percent point (pp) improvements in comparison with the SOTA models. Event modification performance is presented in Appendix C.

We also presented the comparison on nested event categories in Table 2. In these tasks, nested events belong to one of the following categories: *Regulation*, *Positive_regulation*, *Negative_regulation*, and *Planned_process* (for CG and MLEE tasks). As the result shows, our DeepEventMine model obtains the better performance on almost all of the nested event categories (except few cases in ID and MLEE tasks).

5.2 End-to-end Extraction Performance

We show the performance of our end-to-end extraction system and compare it with the pipeline system in Table 3. For reference, we also show the performance of these models given gold entities.¹⁰

As shown in Table 3, our end-to-end model consistently performed better than the pipeline model in event extraction in all of the tasks. This implies that our end-to-end model can overcome the drawback of the pipeline training systems, where errors can be cascaded through different pipeline modules. With our end-to-end model, we represent and discover relationships among triggers, roles, and events via shared vector representation and pass the training loss from event detection back to the role detection and trigger detection, which alleviate the problem of error propagation in pipeline training.

¹⁰ We also tried randomly-initialized BERT model in the setting of Table 3 given gold entities, but as expected, the model performed poorly (with the F1-score of 32.8% in event detection of the CG task) since the parameters in BERT is too many compared to the size of the training data.

⁹ <https://github.com/pfnet/optuna>

Table 2. Results of nested event categories on the test sets. (ens.: ensemble, TOTAL_REGULATION: micro-averaged scores on the three regulation types. % nested events: the proportion of the events in total nested categories among the events in a corpus.)

Nested category	SOTA			DeepEventMine (ens.)		
	P	R	F	P	R	F (%)
CG (vs. TEES-CNN)						
Regulation	50.52	30.93	38.36	61.13	34.07	43.75
Positive_regulation	61.28	44.24	51.38	66.26	45.72	54.11
Negative_regulation	56.60	45.97	50.73	62.23	44.29	51.75
TOTAL REGULATION	57.33	41.14	47.90	63.88	42.19	50.82
Planned_process	48.53	34.10	40.05	57.45	42.33	48.75
% nested events	50.76%					
EPI (vs. TEES-CNN)						
Catalysis	46.67	6.14	10.85	81.25	23.68	36.68
% nested events	8.27%					
GE11 (vs. TEES-CNN)						
Regulation	53.47	34.03	41.59	67.69	34.29	45.52
Positive_regulation	63.63	38.67	48.10	69.42	47.68	56.53
Negative_regulation	54.89	43.26	48.38	64.27	46.94	54.25
TOTAL REGULATION	59.54	39.02	47.14	67.87	45.35	54.37
% nested events	53.83%					
GE13 (vs. BioMLN)						
Regulation	–	–	–	64.04	25.35	36.32
Positive_regulation	–	–	–	62.84	38.76	47.95
Negative_regulation	–	–	–	64.31	44.87	52.86
TOTAL REGULATION	50.86	36.47	42.48	63.41	38.43	47.85
% nested events	58.89%					
ID (vs. EventMine)						
Regulation	44.00	22.80	30.03	59.65	17.62	27.20
Positive_regulation	63.95	49.23	55.63	64.77	59.49	62.02
Negative_regulation	74.63	55.25	63.49	74.02	51.93	61.04
TOTAL REGULATION	62.47	42.18	50.36	67.22	42.88	52.36
% nested events	41.50%					
PC (vs. TEES-CNN)						
Regulation	50.48	35.81	41.90	60.33	33.11	42.75
Positive_regulation	56.05	36.86	44.48	66.24	35.88	46.55
Negative_regulation	52.38	43.67	47.63	59.90	44.05	50.76
TOTAL REGULATION	53.69	38.43	44.80	62.94	37.43	46.95
Activation	75.48	79.76	77.56	81.28	77.33	79.25
Inactivation	51.67	47.69	49.60	52.17	36.92	43.24
% nested events	57.57%					
MLEE (vs. PMCNN)						
Regulation	42.96	27.16	33.28	50.00	25.00	33.33
Positive_regulation	45.92	39.65	42.56	61.33	49.16	54.57
Negative_regulation	51.01	38.49	43.87	57.21	41.83	48.32
TOTAL REGULATION	-	-	-	57.86	40.53	47.67
Planned_process	44.03	37.36	40.42	50.00	32.14	39.13
% nested events	53.17%					

Although successfully achieving our goal which is event detection, our end-to-end model performed worse than the pipeline model on entity, trigger and role detection (except for two cases of trigger detection in ID and MLEE). To analyze the reason behind this performance, we additionally calculated the ratio of predicted entities generated correct events over the total predicted entities. For instance, in the CG task given gold entities, the ratios are 64.79% (2,501/3,860) and 62.13% (2,459/3,958) for the end-to-end and pipeline models, respectively. The scores evidently indicate that our end-to-end model pays more attention on predicting event-related elements, i.e., entities, rather than the pipeline model. This is acceptable for event extraction, which is our main aim in this work, but in practice, we may need to choose the model settings by applications. We will leave this for future work.

When we compare the performance of our model (DeepEventMine) with gold entities and one without gold entities, the performance difference is in the range of 3 to 10% points (pp). This performance gap is relatively small considering the entity detection performances, i.e., 81–86%. This reveals that our end-to-end model, which includes its own entity recognizer and does not depend on any external entity recognizer, is capable to detect events from raw texts with a promising result. The gap in GE13 task is higher with 10% when the entity detection is only 81%.

5.3 Detailed Analysis on Each Layer and Nested Events

In order to investigate the contribution of each layer to the performance improvement, we compare our model with the TEES-CNN model on the CG development set on each layer. The results presented in Table 4 reveal that our model achieves better performance than the TEES-CNN model in all trigger, role, and event scores.

We also conducted an analysis of our model in extracting nested events on the CG development set. Nested events cover up to 34% of the total number of events. Please refer to Appendix A for the detailed statistics of the nested events. Table 5 demonstrates the performance of TEES-CNN and our model on different nested event levels. Level 0 corresponds to flat event extraction, while level 1 – 3 show the level of nested event. For flat events, our model obtained 68.98% in F1-score, which is 5 pp higher than the TEES-CNN. Although the performance on nested event extraction is lower than that on flat events, our model still detected more nested events than TEES-CNN even for the higher nested levels 2 and 3.

5.4 Error Analysis

We conducted error analysis on the event extraction results of our model and summarized them in Table 6. In particular, we analyze here six types of errors that can cause incorrect or missing events, including:

- **Missing role:** when the model cannot extract an event due to errors that occur before the event layer.
- **Missing trigger:** when the model cannot extract an event because it cannot detect the trigger.
- **Incorrect event class:** when the textual span of a trigger is correctly detected but classified to a wrong class.
- **Missing entity:** similar to “missing trigger”, we count the number of missing events due to undetected entity.
- **Missing argument:** when an event trigger is correctly detected, but its arguments are missing.
- **Missing nested argument:** when the event argument of a nested event has incorrect core arguments.¹¹

As presented in Table 6, missing events from our prediction were mainly caused by missing roles between triggers and arguments, accounted for a total of 89.0%. In detail, around 38.4% (497/1,294) and 50.2% (650/1,294) of missing role were due to undetected entities and triggers in the Entity/Trigger layer respectively, although the entity/trigger layer performed remarkably well on entity/trigger extraction. These numbers are high considering the performance in Table 3. To overcome this pitfall, we need to consider how to put more focus on entities and triggers that are related to multiple events. Furthermore, 3.9% (25/650) of missing triggers were due to incorrect classification to different event classes, which means we need to improve the disambiguation performance rather than span detection performance. When we focus on the cases where the arguments are incorrect or missing while their triggers are correct, 78.7% (107/136) of these cases were caused by incorrect or missing child events. This result

¹¹ We refer the readers to Pyysalo *et al.* (2015) for the details of core arguments.

Table 3. Comparison of our model with the pipeline training setting when extracting events from raw text and given gold entities on the development sets. (gold: given gold entities)

Task	Model	Entity			Trigger			Role			Event		
		P	R	F	P	R	F	P	R	F	P	R	F (%)
CG	Pipeline (gold)	-	-	-	78.85	82.89	80.82	67.15	66.38	66.76	59.64	53.66	56.49
	DeepEventMine (gold)	-	-	-	79.17	82.93	81.01	63.18	66.65	64.87	65.24	55.93	60.23
	Pipeline	85.47	84.12	84.79	79.18	81.90	80.52	61.99	60.84	61.41	53.25	47.49	50.20
	DeepEventMine	85.57	82.02	83.76	79.54	81.45	80.48	59.72	60.70	60.20	61.23	48.74	54.27
EPI	Pipeline (gold)	-	-	-	74.43	83.11	78.53	75.74	65.50	70.25	71.02	55.29	62.18
	DeepEventMine (gold)	-	-	-	76.99	82.52	79.66	76.83	67.25	71.72	75.90	56.18	64.56
	Pipeline	85.06	84.02	84.54	74.43	83.11	78.53	67.53	61.07	64.14	59.61	49.55	54.12
	DeepEventMine	83.84	82.04	82.93	73.52	81.94	77.50	66.46	62.35	64.34	59.69	52.40	55.81
GE11	Pipeline (gold)	-	-	-	71.11	69.97	70.54	69.96	59.63	64.38	67.99	56.37	61.64
	DeepEventMine (gold)	-	-	-	71.23	70.31	70.77	63.52	59.09	61.22	70.52	56.52	62.75
	Pipeline	88.69	84.64	86.62	73.32	68.72	70.95	66.25	55.52	60.41	60.63	50.25	54.95
	DeepEventMine	88.51	84.29	86.35	72.05	68.89	70.43	60.82	57.14	58.92	62.36	51.88	56.64
GE13	Pipeline (gold)	-	-	-	76.59	68.73	72.45	69.45	57.07	62.65	59.37	48.37	53.31
	DeepEventMine (gold)	-	-	-	74.29	71.25	72.74	62.50	54.64	58.31	64.50	49.25	55.85
	Pipeline	82.36	80.02	81.17	75.21	70.36	72.70	62.51	53.42	57.61	48.39	41.90	44.91
	DeepEventMine	81.11	80.74	80.93	74.96	69.42	72.08	58.09	52.04	54.90	49.49	42.88	45.95
ID	Pipeline (gold)	-	-	-	71.96	84.06	77.54	52.34	61.68	56.65	52.82	56.39	54.54
	DeepEventMine (gold)	-	-	-	74.56	80.24	77.30	52.17	58.88	55.32	59.91	53.94	56.77
	Pipeline	81.97	86.24	84.05	71.79	83.36	77.15	48.07	55.66	51.59	47.94	51.72	49.76
	DeepEventMine	81.93	85.46	83.66	73.15	82.15	77.39	44.36	48.95	46.54	58.14	44.02	50.10
PC	Pipeline (gold)	-	-	-	75.26	80.71	77.89	65.60	65.06	65.33	53.82	52.11	52.95
	DeepEventMine (gold)	-	-	-	76.97	77.74	77.35	63.54	62.96	63.25	65.94	49.52	56.57
	Pipeline	88.26	89.61	88.93	75.26	80.71	77.89	61.73	61.59	61.66	48.13	47.67	47.90
	DeepEventMine	87.80	89.25	88.52	75.73	78.73	77.20	57.77	60.44	59.08	56.90	45.45	50.53
MLEE	Pipeline (gold)	-	-	-	80.77	73.93	77.20	62.09	56.22	59.01	55.66	48.60	51.89
	DeepEventMine (gold)	-	-	-	79.73	76.39	78.03	56.64	55.92	56.28	63.52	48.26	54.85
	Pipeline	82.69	80.78	81.72	81.49	78.43	79.93	56.47	55.32	55.89	50.14	45.79	47.86
	DeepEventMine	81.87	81.41	81.64	79.37	78.86	79.12	51.93	56.52	54.13	56.33	47.83	51.73

Table 4. Comparison of our model with the TEES-CNN on the CG development set given gold entities.

Model	Trigger			Role			Event		
	P	R	F	P	R	F	P	R	F (%)
TEES-CNN	76.81	80.87	78.78	65.10	62.83	63.95	59.24	51.81	55.27
Pipeline	78.85	82.89	80.82	67.15	66.38	66.76	59.64	53.66	56.49
DeepEventMine	79.17	82.93	81.01	63.18	66.65	64.87	65.24	55.93	60.23

Table 5. Results on different nested event levels on the CG development set.

Nested level	TEES-CNN			DeepEventMine(gold)		
	P	R	F	P	R	F (%)
Flat + nested events	60.42	53.45	56.72	66.22	57.19	61.37
TOTAL flat	64.99	62.99	63.98	71.19	66.91	68.98
TOTAL nested	48.82	35.23	40.93	53.90	38.62	45.00
Level 1	51.81	38.62	44.25	54.49	42.02	47.45
Level 2	26.93	16.78	21.43	46.67	19.58	27.59
Level 3	0.00	0.00	0.00	100.00	16.67	28.57
% nested events	34.37%					

Table 6. Analysis of missing events on the CG development set.

Error Type	#events	%
Missing role	1,294	89.0
Missing trigger	650	44.7
Incorrect event class	25	1.7
Missing entity	497	34.2
Missing argument	136	9.4
Missing event argument	107	7.4
Total	1,454	100

aligns with the performance drop in nested event detection in Table 5 and there is still room for improvement. Some examples of events predicted by our model are presented in Appendix F.

6 Conclusion

We proposed a neural nested event extraction model DeepEventMine that can predict nested entities and nested events in an end-to-end manner. On top of the expressive pre-trained model BERT, our model can simultaneously detect entities/triggers, roles and events. Experimental results and analysis showed that given gold entities, the proposed DeepEventMine model consistently improves the performance in all the

subtasks and the end-to-end learning further improves the event detection performance with slightly degrading entity/trigger and role detection performance. As a result, our DeepEventMine model has achieved the SOTA performance on the test set portion of the seven biomedical domain data sets: Cancer Genetics 2013, Epigenetics and Post-translational Modifications 2011, GENIA 2011, GENIA 2013, Infectious Diseases 2011, Pathway Curation 2013, and MLEE. For future work, we aim at generalizing and applying the model to other NLP tasks to evaluate the ability of our model to represent and extract any hyper-graphs from text.

Funding

This paper is based on results obtained from a project commissioned by the New Energy and Industrial Technology Development Organization (NEDO). This work is also supported by PRISM (Public/Private R&D Investment Strategic Expansion PrograM). This research was partially supported with funding from Biotechnology and Biological Services Research Council, Japan Partnership Award (Grant ID: BB/P025684/1) (SA).

Acknowledgements

We would like to thank the reviewers for their insightful comments. We also would like to thank the Institute of Advanced Industrial Science and Technology/Artificial Intelligence Research Center for providing the computational resources for the realization of the experiments conducted in this work. In addition, TTT thanks the University of Manchester for the Research Impact Scholarship Award.

References

- Beltagy, I., Lo, K., and Cohan, A. (2019). SciBERT: A pretrained language model for scientific text. In *Proceedings of EMNLP-IJCNLP*, pages 3606–3611. ACL.
- Björne, J. and Salakoski, T. (2013). TEES 2.1: Automated annotation scheme learning in the BioNLP 2013 shared task. In *Proceedings of BioNLP*, pages 16–25. ACL.
- Björne, J. and Salakoski, T. (2018). Biomedical event extraction using convolutional neural networks and dependency parsing. In *Proceedings of BioNLP*, pages 98–108. ACL.
- Chen, Y., Xu, L., Liu, K., Zeng, D., and Zhao, J. (2015). Event extraction via dynamic multi-pooling convolutional neural networks. In *Proceedings of ACL-IJCNLP*, volume 1, pages 167–176.
- Devlin, J., Chang, M.-W., Lee, K., and Toutanova, K. (2019). BERT: Pre-training of deep bidirectional transformers for language understanding. In *Proceedings of NAACL*, pages 4171–4186.
- Finkel, J. R. and Manning, C. D. (2009). Nested named entity recognition. In *Proceedings of EMNLP*, pages 141–150. ACL.
- Hakala, K., Van Landeghem, S., Kaewphan, S., Salakoski, T., Van de Peer, Y., and Ginter, F. (2012). Cyevex: Literature-scale network integration and visualization through cytoscape. *Proceedings of SBBM*, 12, 91–96.
- He, X., Li, L., Song, X., Huang, D., and Ren, F. (2019). Multi-level attention based blstm neural network for biomedical event extraction. *IEICE TRANSACTIONS on Information and Systems*, 102(9), 1842–1850.
- Hendrycks, D. and Gimpel, K. (2016). Gaussian error linear units (gelus). *arXiv preprint arXiv:1606.08415*.
- Ju, M., Miwa, M., and Ananiadou, S. (2018). A neural layered model for nested named entity recognition. In *Proceedings of NAACL*, pages 1446–1459. ACL.
- Katiyar, A. and Cardie, C. (2018). Nested named entity recognition revisited. In *Proceedings of NAACL*, pages 861–871. ACL.
- Kim, J.-D., Ohta, T., Pyysalo, S., Kano, Y., and Tsujii, J. (2011a). Extracting biomolecular events from literature. In the bionlp shared task. *Computational Intelligence*, 27(4), 513–540.
- Kim, J.-D., Wang, Y., Takagi, T., and Yonezawa, A. (2011b). Overview of genia event task in bionlp shared task 2011. In J. Tsujii, J.-D. Kim, and S. Pyysalo, editors, *Proceedings of BioNLP*, pages 7–15. ACL.
- Kim, J.-D., Wang, Y., and Yamamoto, Y. (2013). The genia event extraction shared task, 2013 edition - overview. In C. Nédellec, R. Bossey, J.-D. Kim, J. jae Kim, T. Ohta, S. Pyysalo, and P. Zweigenbaum, editors, *Proceedings of BioNLP*, pages 8–15. ACL.
- Kingma, D. P. and Ba, J. L. (2015). Adam: A method for stochastic optimization. In *Proceedings of ICLR*.
- Lee, J., Yoon, W., Kim, S., Kim, D., Kim, S., So, C. H., and Kang, J. (2019). BioBERT: a pre-trained biomedical language representation model for biomedical text mining. *Bioinformatics*, btz682.
- Li, D., Huang, L., Ji, H., and Han, J. (2019). Biomedical event extraction based on knowledge-driven tree-lstm. In *Proceedings of NAACL*.
- Li, L., Liu, Y., and Qin, M. (2018). Extracting biomedical events with parallel multi-pooling convolutional neural networks. *IEEE/ACM transactions on computational biology and bioinformatics*.
- Li, Q., Ji, H., and Huang, L. (2013). Joint event extraction via structured prediction with global features. In *Proceedings of ACL*, pages 73–82. ACL.
- Liu, X., Luo, Z., and Huang, H. (2018). Jointly multiple events extraction via attention-based graph information aggregation. In *Proceedings of EMNLP*, pages 1247–1256. ACL.
- Miwa, M. and Ananiadou, S. (2013). NaCTeM EventMine for BioNLP 2013 CG and PC tasks. In *Proceedings of BioNLP*, pages 94–98. ACL.
- Miwa, M., Thompson, P., and Ananiadou, S. (2012a). Boosting automatic event extraction from the literature using domain adaptation and coreference resolution. *Bioinformatics*, 28(13), 1759–1765.
- Miwa, M., Thompson, P., and Ananiadou, S. (2012b). Boosting automatic event extraction from the literature using domain adaptation and coreference resolution. *Bioinformatics*, 28(13), 1759–1765.
- Miwa, M., Ohta, T., Rak, R., Rowley, A., Kell, D. B., Pyysalo, S., and Ananiadou, S. (2013a). A method for integrating and ranking the evidence for biochemical pathways by mining reactions from text. *Bioinformatics*, 29(13), i44–i52.
- Miwa, M., Pyysalo, S., Ohta, T., and Ananiadou, S. (2013b). Wide coverage biomedical event extraction using multiple partially overlapping corpora. *BMC Bioinformatics*, 14, 175.
- Nguyen, T. H., Cho, K., and Grishman, R. (2016). Joint event extraction via recurrent neural networks. In *Proceedings of NAACL*, pages 300–309.
- Nguyen, T. M. and Nguyen, T. H. (2019). One for all: Neural joint modeling of entities and events. In *Proceedings of AAAI*. Association for the Advancement of Artificial Intelligence.
- Ohta, T., Pyysalo, S., and Tsujii, J. (2011). Overview of the epigenetics and post-translational modifications (EPI) task of BioNLP shared task 2011. In *Proceedings of BioNLP*, pages 16–25, Portland, Oregon, USA. ACL.
- Paszke, A., Gross, S., Chintala, S., Chanan, G., Yang, E., DeVito, Z., Lin, Z., Desmaison, A., Antiga, L., and Lerer, A. (2017). Automatic differentiation in pytorch. In *NIPS-W*.
- Peters, M., Neumann, M., Iyyer, M., Gardner, M., Clark, C., Lee, K., and Zettlemoyer, L. (2018). Deep contextualized word representations. In *Proceedings of NAACL*, pages 2227–2237. ACL.
- Pyysalo, S., Ohta, T., Rak, R., Sullivan, D. E., Mao, C., Wang, C., Sobral, B. W. S., Tsujii, J., and Ananiadou, S. (2011). Overview of the infectious diseases (id) task of bionlp shared task 2011. In J. Tsujii, J.-D. Kim, and S. Pyysalo, editors, *Proceedings of BioNLP*, pages 26–35. ACL.
- Pyysalo, S., Ohta, T., Miwa, M., Cho, H.-C., Tsujii, J., and Ananiadou, S. (2012). Event extraction across multiple levels of biological organization. *Bioinformatics*, 28(18), i575–i581.
- Pyysalo, S., Ohta, T., Rak, R., Rowley, A., Chun, H.-W., Jung, S.-J., Choi, S.-P., Tsujii, J., and Ananiadou, S. (2015). Overview of the cancer genetics and pathway curation tasks of bionlp shared task 2013. *BMC bioinformatics*, 16(10), S2.
- Radford, A., Narasimhan, K., Salimans, T., and Sutskever, I. (2018). Improving language understanding by generative pre-training. Technical report, Open AI.
- Sha, L., Qian, F., Chang, B., and Sui, Z. (2018). Jointly extracting event triggers and arguments by dependency-bridge RNN and tensor-based argument interaction. In *Proceedings of AAAI*.
- Sohrab, M. G. and Miwa, M. (2018). Deep exhaustive model for nested named entity recognition. In *Proceedings of EMNLP*, pages 2843–2849. ACL.
- Tsuruoka, Y., Tsujii, J., and Ananiadou, S. (2008). Facta: a text search engine for finding associated biomedical concepts. *Bioinformatics*, 24(21), 2559–2560.
- Van Landeghem, S., Björne, J., Wei, C.-H., Hakala, K., Pyysalo, S., Ananiadou, S., Kao, H.-Y., Lu, Z., Salakoski, T., Van de Peer, Y., et al. (2013). Large-scale event extraction from literature with multi-level gene normalization. *PLoS one*, 8(4), e55814.
- Venugopal, D., Chen, C., Gogate, V., and Ng, V. (2014). Relieving the computational bottleneck: Joint inference for event extraction with high-dimensional features. In *Proceedings of EMNLP*, pages 831–843.
- Wang, A., Wang, J., Lin, H., Zhang, J., Yang, Z., and Xu, K. (2017). A multiple distributed representation method based on neural network for biomedical event extraction. *BMC medical informatics and decision making*, 17(3), 171.

Wang, B. and Lu, W. (2018). Neural segmental hypergraphs for overlapping mention recognition. In *Proceedings of EMNLP*, pages 204–214. ACL.

Yan, S. and Wong, K.-C. (2019). Context awareness and embedding for biomedical event extraction. *Bioinformatics*. btz607.

Yang, B. and Mitchell, T. M. (2016). Joint extraction of events and entities within a document context. In *Proceedings of NAACL*, pages 289–299. ACL.