

Learning Latent Forests for Medical Relation Extraction

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Task

- Input:** A sentence from medical literature together with given entities.
Crystallographic analysis of the human phenylalanine hydroxylase catalytic domain with bound catechol inhibitors at 2.0 Å resolution.
- Output:** The relation between given entities: **Down regulator**

Motivation

Dependency structures are often used for relation extraction as they are able to capture long-range relations that are only implicit in the surface form alone. However, dependency parsing accuracy is **relatively low** in the medical domain, which may downstream the relation extraction pipeline.

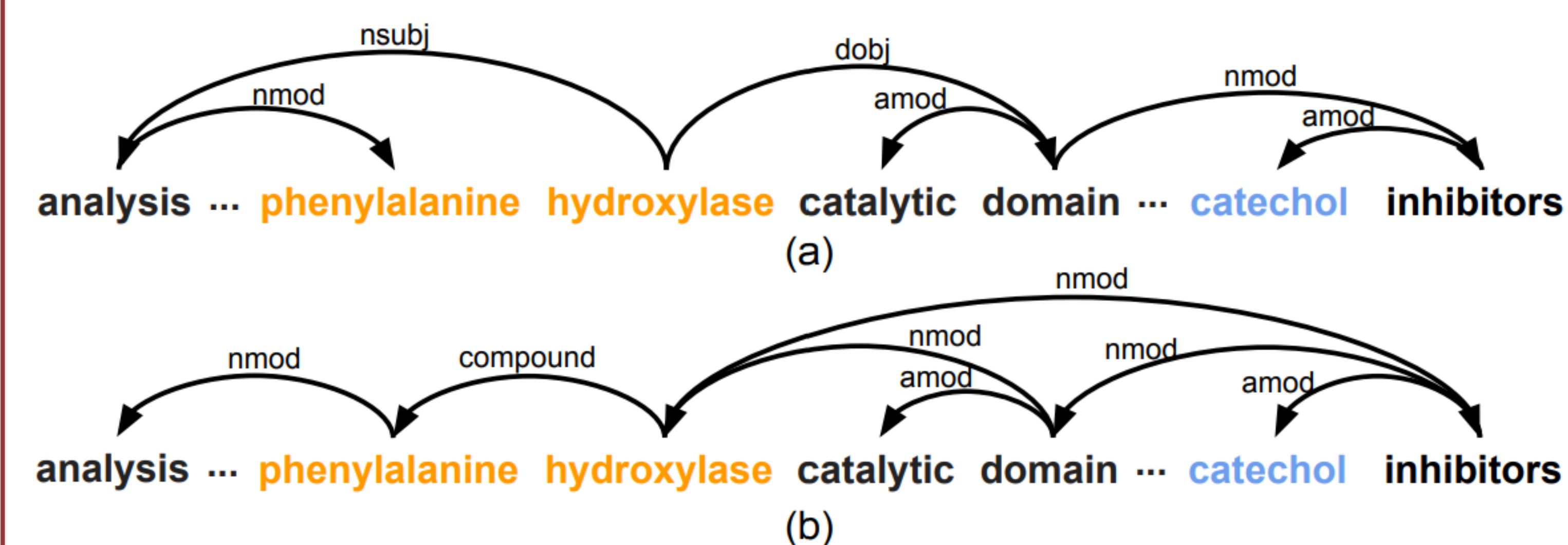
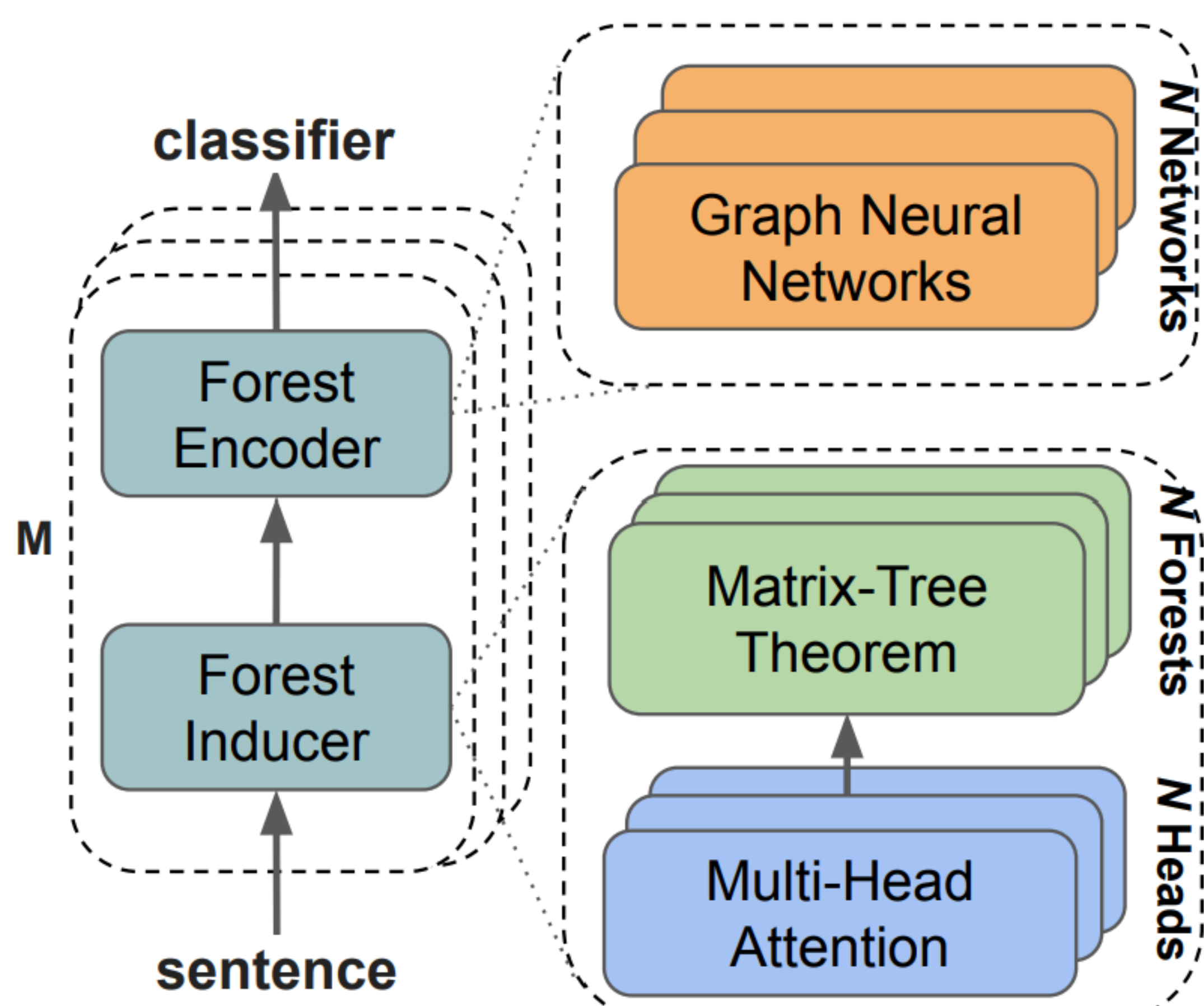


Figure (a) shows the 1-best dependency tree, (b) is the manually labeled tree.

- The 1-best dependency tree obtained by an off-the-shelf parser contains an error. Specifically, the entity phrase *phenylalanine hydroxylase* is broken since *hydroxylase* is mistakenly considered as the main verb.
- Unlike previous research efforts that rely on **dependency parsers trained on newswire text**, our model is able to generate **task-specific dependency structures** for capturing non-local interactions.

Model

- Our model consists of two modules: **Forest Inducer** and **Forest Encoder**. The forest inducer has two sub-modules, where the first one computes N attention matrices and the second sub-module takes the N attention matrices as inputs to obtain N **dependency forests**. Then the forest encoder uses graph neural networks to encode the induced forests.

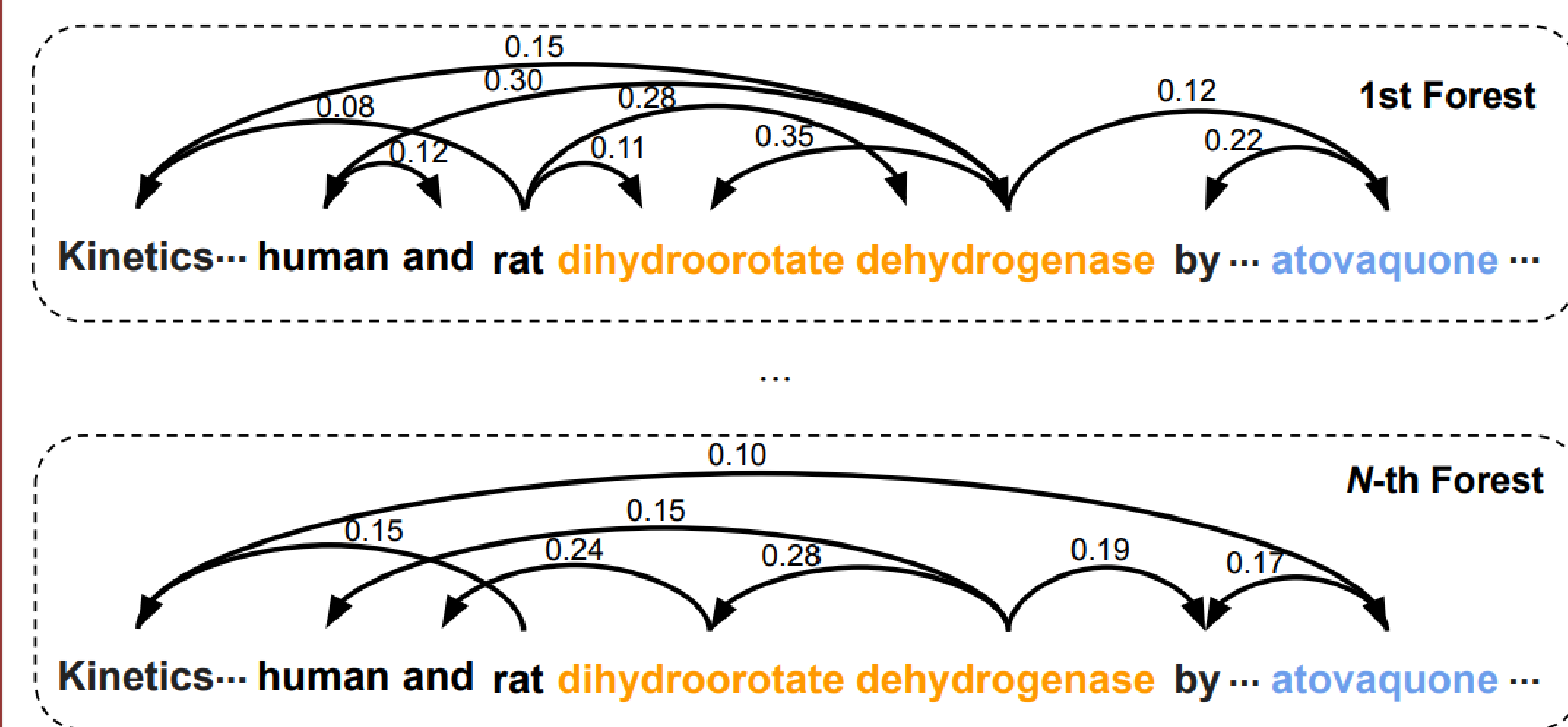


Latent Forests

Inspired by Liu and Lapata (2018), we use a **variant of Kirchhoff's Matrix-Tree Theorem** (MTT; Koo et al., 2007; Smith and Smith, 2007) to induce the latent structure of an input sentence. For the graph G , MTT takes the edge scores and root scores as inputs then generates a latent forest by computing the **marginal probabilities for each edge**:

$$P(z_{ij}^k = 1) = \sum_{y \in \mathbf{T}^k: (i,j) \in y} P(y|\mathbf{h}; \theta^k)$$

Induced latent structure as shown below can be viewed as **multiple full dependency forests**, which efficiently represent all possible dependency trees within a compact and dense structure.



Experiments

- We evaluate our proposed model with four medical relation extraction datasets on two tasks, including cross-sentence n -ary relation extraction (Peng et al., 2017) and sentence-level relation extraction, CPR (Krallinger et al., 2017) and PGR (Sousa et al., 2019).

Syntax Type	Model	Binary-class				Multi-class	
		Single	Cross	Single	Cross	Ternary	Binary
Full Tree	DAG LSTM [Peng et al., 2017]	77.9	80.7	74.3	76.5	-	-
	GRN [Song et al., 2018]	80.3	83.2	83.5	83.6	71.7	71.7
	GCN [Zhang et al., 2018]	84.3	84.8	84.2	83.6	77.5	74.3
Pruned Tree	GCN [Zhang et al., 2018]	85.8	85.8	83.8	83.7	78.1	73.6
Forest	AGGCN [Guo et al., 2019a]	87.1	87.0	85.2	85.6	79.7	77.4
	LF-GCN (Ours)	88.0	88.4	86.7	87.1	81.5	79.3

Syntax Type	Model	F1	Syntax Type	Model	F1
None	BioBERT [Lee et al., 2019]	67.2	None	Random-DDCNN [Lifeng et al., 2020]	45.4
Tree	BO-LSTM [Lamurias et al., 2019]	52.3	None	Att-GRU [Liu et al., 2017]	49.5
	GCN [Zhang et al., 2018]	81.3*	Tree	Bran [Verga et al., 2018]	50.8
	Tree-GRN [Lifeng et al., 2020]	78.9	Tree	GCN [Zhang et al., 2018]	52.2*
Forest	Edgewise-GRN [Song et al., 2019]	83.6	Tree	Tree-DDCNN [Lifeng et al., 2020]	50.3
	KBest-GRN [Song et al., 2019]	85.7	Tree	Tree-GRN [Lifeng et al., 2020]	51.4
	AGGCN [Guo et al., 2019a]	89.3*	Forest	Edgewise-GRN [Song et al., 2019]	53.4
	ForestFT-DDCNN [Lifeng et al., 2020]	89.3	Forest	KBest-GRN [Song et al., 2019]	52.4
	LF-GCN (Ours)	91.9	Forest	AGGCN [Guo et al., 2019a]	56.7*
				ForestFT-DDCNN [Lifeng et al., 2020]	55.7
			LF-GCN (Ours)	58.9	

References

- Liu and Lapata. Learning structured text representations. TACL18.
- Koo et al., Structured prediction models via the matrix-tree theorem. EMNLP07.
- Smith et al., Probabilistic models of nonprojective dependency trees. EMNLP07.