Event Extraction as Dependency Parsing (in BioNLP 2011)

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Our approach in two slides...





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Full details in [McClosky, Surdeanu, and Manning, ACL 2011]



Outline





3 Experiments





Approach

... tax, acts as a costimulatory signal for GM-CSF and IL-2 gene transcription ...

Preprocessing: Segmentation, tokenization



Approach



... tax, acts as a costimulatory signal for GM-CSF and IL-2 gene transcription ...

Preprocessing: Segmentation, tokenization, syntactic parsing

Self-trained biomedical parser: [McClosky, 2010]



Approach



Anchor classification: Token classification for event anchors

(similar to [Björne et al., BioNLP 2009])





Overview

Approach



Event parsing: Parse anchors and proteins using reranking parser



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- Handles non-projective trees naturally
- Easy to extend feature extractor
- Support for *n*-best parsing



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 - Distributional similarity features in anchor detection



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 - Update event type information (EPI, ID)
 - Combine ID training data with GENIA (ID)
 - Removing nested entities (ID)



Results on Genia development

Decoder(s)	Parser	Reranker
1P	49.0	49.4
2P	49.5	50.5
1N	49.9	50.2
2N	46.5	47.9
All		50.7



Results on Epigenetics development

Decoder(s)	Parser	Reranker
1P	62.3	63.3
2P	62.2	63.3
1N	62.9	64.6
2N	60.8	63.8
All		64.1

(note: issues with our internal evaluator implementation)



Domain adaptation for Infectious Diseases

Model	Precision	Recall	f-score
ID	59.3	38.0	46.3
ID (×1) + GE	52.0	40.2	45.3
ID ($ imes$ 2) + GE	52.4	41.7	46.4
ID ($ imes$ 3) + GE	54.8	45.0	49.4
ID ($ imes$ 4) + GE	55.2	43.8	48.9
ID ($ imes$ 5) + GE	55.1	44.7	49.4

(parser only with 2N decoder)



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2P	47.8	49.8
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All		50.2



Summary

- New approach to event extraction
 - Parsing can be used for event extraction
 - Reranker further improves performance
- Minimal changes to adapt to new BioNLP domains
- Component in the FAUST system (stay tuned!)
- Code coming soon!

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Questions?

