

Biomedical Event Annotation with CRFs and Precision Grammars

Andrew MacKinlay, David Martinez, Timothy Baldwin

Genomic Event Extraction

*TRADD was the only protein that **interacted** with wild-type TES2 and not with isoleucine-mutated TES2.*

- ▶ Proteins marked in input: *TRADD*, *TES2*, *TES2*
- ▶ Events:
 - ▶ ID: Ev1, Type: Binding, Trigger: **interacted**, Theme1: *TRADD*, Theme2: *TES2*
 - ▶ ID: Ev2, Type: Binding, Trigger: **interacted**, Theme1: *TRADD*, Theme2: *TES2*
 - ▶ ID: Modif1, Type: Negation, Theme: Ev2

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BioNLP task

- ▶ 3 tasks:
 1. Identify events and theme(s)
 2. Identify additional arguments, e.g. localization
 3. Detect event modifiers (negations and speculations)
- ▶ Event types:
 - ▶ single-theme events: e.g. *gene expression*
 - ▶ multiple-theme events: *binding* of proteins or events
 - ▶ regulation events: regulate other events. E.g. X **inhibits** Y
- ▶ 9 different entities: subset of Genia event ontology

Motivation and Architecture

- ▶ Our motivation: deep linguistic processing for detection of speculation and negation
- ▶ Architecture:
 - ▶ Task 1:
 - ▶ Trigger word detection: CRF and Lookup systems
 - ▶ Event-theme construction (hand-crafted rules)
 - ▶ Task 3:
 - ▶ Deep parsing for semantic representation
 - ▶ Classification of events using maximum entropy learners

Trigger Word Detection with CRFs

- ▶ Conditional probability distribution over label sequences given a particular observation sequence
- ▶ CRF++ toolkit
- ▶ Tested features: word-form, lemma, POS, chunking marks, protein NER, grammatical dependencies (from Bikel parser and GDep)
- ▶ JULIE-Lab sentence splitter and Genia Tagger for pre-processing
- ▶ Window sizes: ± 3 and ± 4

Trigger Word Detection with CRFs

- ▶ Best results (training data): Precision $\sim 66\%$, Recall $\sim 30\%$
- ▶ All features help except for grammatical dependencies
- ▶ ± 3 window size

Trigger Word Detection with Dictionary Look-up

- ▶ Decision list for each trigger string found in training data
 - ▶ Simply assign highest frequency class
- ▶ Frequency cut-off
- ▶ We can reach high recall ($\sim 77\%$) but at the cost of precision ($\sim 13\%$)
- ▶ Best f-score $\sim 36\%$ ($\sim 50\%$ recall)

Trigger Word Detection: Combination

- ▶ Add all trigger words identified by CRF and look-up
- ▶ Two approaches:
 - ▶ Optimise per class (Optim)
 - ▶ Always preference to CRF (All)

Event-theme Construction

- ▶ Approach: assign closest events/proteins as themes (without crossing sentence boundaries)
- ▶ Basic events:
 - ▶ Single closest protein
- ▶ Binding events:
 - ▶ Closest proteins
 - ▶ Parameters: maximum distance and number of themes
- ▶ Regulation events
 - ▶ Single closest protein or event (give precedence to events)
 - ▶ Parameters: maximum distance and detect/ignore CAUSE

Task 1 Results

System	Rec.	Prec.	F-Score
Combined (Optim.)	17.44	39.99	24.29
Combined (All)	24.36	30.87	27.23
CRF	12.23	62.24	20.44
CRF (+ synt feats)	12.01	61.91	20.11
Look-Up	22.88	29.67	25.84
Look-Up (freq \geq 20)	23.26	26.74	24.88
Look-Up (freq \geq 30)	21.37	30.50	25.13

Table: Task 1 results with approximate span matching, recursive evaluation (our final submission is in bold)

Negation/Speculation Detection

- ▶ English Resource Grammar (ERG): high-precision grammar in the HPSG framework
- ▶ GENIA tagger to deal with named entities
- ▶ 72% of training sentences parsed

Feature Extraction

- ▶ Semantic formalism: Robust Minimal Recursion Semantics
- ▶ Elementary Predicates (EP): Predicates with their arguments
- ▶ Relationships between trigger EP and lexical cues
 - ▶ Outscoping and shared-argument

Features for Negation Identification

- ▶ Pre-identify word lists:
 - ▶ Conjunctions: *_not_c*, *_but+not_c*, *_nor_c*
 - ▶ Other markers: *_only_a*, *_never_a*, *_not+as+yet_a*, *_not+as+yet_a*, *_unable_a*, *neg_rel*
- ▶ Negative-outscope feature: when negative EP outscopes trigger-EP
 - ▶ E.g. “...product was **not (NEG-EP)** able to **bind (TRIG-EP)** DNA and...”
 - ▶ NegOutscope *neg_rel* = 1
 - ▶ NegOutscope *not* = 1

Features for Negation Identification

- ▶ ...product was **not** able to **bind** DNA and was recovered in cytoplasmic cellular extracts...
- ▶ ERG analysis
 - ▶ **I8**: neg_rel⟨692 : 695⟩(e9, ARG1: **h10**)
 - ▶ **I11**: _able_a_1⟨696 : 700⟩(e12, ARG1: x6, ARG2: **h13**)
 - ▶ **I14**: _bind_v_to⟨704 : 708⟩(e17, ARG1: x6, ARG2: x16, ARG3: u15)
 - ▶ h10 qeq **I11**, h13 qeq **I14**
- ▶ Thus I8 immediately outscopes I11, and I11 immediately outscopes I14

Features for Negation Identification

- ▶ Negative conjunction: when trigger-EP is the argument (ARG0) of a negative conjunction EP
 - ▶ E.g. “...**but not (NEG-EP) binding (TRIG-EP) DNA...**”
- ▶ When trigger-EP is the argument (ARG0) of a negatively-outscoped EP
 - ▶ E.g. “...the **product (TRIG-EP) was never (NEG-EP) considered...**”

Features for Speculation Identification

- ▶ Pre-identify word lists:
 - ▶ Speculation verb short list: *_investigate*, *_study*, *_examine*, *_test*, *_evaluate*, *_observe*}
 - ▶ Extended list: adding WordNet sisters
- ▶ SpecVOBJ: when verb part of “speculative-verbs” set, and object is a trigger word
 - ▶ E.g. “IkappaBalpha phosphorylation and **degradation (TRIG-EP)** was **analyzed (SPEC-EP)**”
 - ▶ SpecVObj2+WN-seed:examine = 1
 - ▶ SpecVObj2+wn-sister:_analyze_v_1(examine) = 1
 - ▶ SpecVObj2+wn-gen = 1

More Features

- ▶ Speculation:
 - ▶ Modal verb outscopes trigger
 - ▶ ARG0 of trigger-EP occurs as argument of the word 'analysis'
- ▶ General features:
 - ▶ E.g. (Modifier adjective) "...Fas **upregulation (TRIG-EP)** is **central (ADJ-EP)** to the preservation..."
 - ▶ 'ModAdj:_central_a_1' = 1
 - ▶ Trigger name, trigger POS, etc.

Negation/Speculation Classifiers

- ▶ Maximum Entropy classifier (Maxent Toolkit)
- ▶ Different feature combinations
- ▶ Baseline: bag of words
- ▶ Development phase:
 - ▶ Goldstandard events
 - ▶ 10-fold cross-validation
- ▶ Test phase:
 - ▶ Trained over goldstandard event extraction
 - ▶ Output of task-1 classifier as source of trigger words

Development Results: Speculation

Feats.	Rec.	Prec.	F-Score
BOW	22.1	47.7	30.2
Spec. + BOW	23.2	57.9	33.1

- ▶ Very low performance over automatic classification, due to pipelining nature of evaluation
- ▶ Linguistic features improve (slightly) over BOW
- ▶ Combination of features works best

Development Results: Negation

Feats.	Rec.	Prec.	FSc.
BOW	15.0	30.2	20.0
Neg. + BOW	24.3	68.4	35.9

- ▶ Bigger improvement over BOW

Official Results for Task 3

TEAM	gold (match)	answer (match)	Rec.	Prec.	F-Score
ConcordU	3617 (1182)	1943 (1182)	32.68	60.83	42.52
VIBGhent	3617 (1105)	2227 (1104)	30.55	49.57	37.80
ASU+HU+BU	3617 (710)	1185 (710)	19.63	59.92	29.57
NICTA	3617 (577)	1450 (575)	15.95	39.66	22.75
USzeged	3617 (722)	3113 (722)	19.96	23.19	21.46
CCP-BTMG	3617 (446)	777 (446)	12.33	57.40	20.30

Lessons Learned

- ▶ Keyword detection suffers from data sparseness
- ▶ Rules for event construction are too naive
- ▶ Deep parsing better than lexical baseline, but there are coverage problems
- ▶ Combined approach (detect triggers and themes together) to be explored for Task 1

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