



Extracting Drug Information with Apache cTAKES and ClearTK



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Background

Adverse drug events cause harm to patients and are not well-understood, especially in pediatric populations. We are interested in designing machine learning algorithms for extracting adverse drug events from texts in the electronic health record.

This dataset largely annotates *physician-inferred adverse drug events*. These represent cases where ADEs were believed by clinicians to have occurred, and then documented.

Data

Example ADEs (ADEs in Red, Drugs in Blue)
Elevated LFTs/coagulopathy: Secondary to shock liver versus tylenol toxicity.

During the admission the pt was thought to have had a **ceftriaxone** allergic reaction (**morbilliform drug rash**) and ceftriaxone was added to allergy list.

She did not experience significant nausea during hospitalization, but continued to have **diarrhea** related to her **chemotherapy** which was treated with Lomotil.

Your **rash** appeared to improve after discontinuation of the **antibiotics piperacillin** and **tazobactam**, but the exact cause was unknown.

Hypotension - The patient was initially hypotensive in the [**Hospital Unit Name 153**]. This was after receiving **Propofol** and then other **sedating medications**.

Methods

We participated in all 3 tracks:

- Entity extraction
- Relation identifiaction
- End-to-end relation extraction

Our focus was on adapting general-purpose clinical NLP tools to this new task. We used:

- ClearTK [1] entity extraction API to train a BIO tagger for entity extraction (above right)
- Features based on characters, word, and part-of-speech of focus word and context words (right)

increase to 50mg po BID for seven days then

Duration	0	0	0	0	0	В	1	1	0
Frequency	0	0	0	0	В	0	0	0	0
Strength	0	0	В	0	0	0	0	0	0
Route	0	0	0	В	0	0	0	0	0
Drug	0	0	0	0	0	0	0	0	0

increase to 50mg po BID for seven days then

ID	increase	to	50mg	ро	BID	for	seven	days	then
POS	VB	TO	Ν	ADJ	PN	Р	CD	NS	Р
Char	L*	L*	N*L*	L*	U*	L*	L*	L*	L*

Apache cTAKES [2] Relation Extractor module for relation classification [3]:

- She did not experience significant **nausea** during hospitalization, but continued to have **diarrhea** related to her **chemotherapy** which was treated with **Lomotil**
- Specify allowed types for each argument:
 - Arg1 : Medication Mention
 - Arg2 : Sign/Symptom/Disease/Disorder
- Compare all pairs within covering type (paragraph)
 - (Chemotherapy, nausea) -> False
 - (Chemotherapy, diarrhea) -> True

(Lomotil, diarrhea) -> False

• (Lomotil, nausea) -> False

		Relations	F1		End to end	F1
Drug	0.9375				Relations	
ADE	0.3092	ADE	0.7635		ADE	0.2758
Dosage	0.9082	Dosage	0.9503		Dosage	0.8780
Duration	0.7106	Duration	0.8223		Duration	0.5992
Form	0.9461	Form	0.9660		Form	0.9219
Frequency	0.9520	Frequency	0.9136		Frequency	0.8784
Reason	0.5077	Reason	0.7497		Reason	0.4195
Route	0.9369	Route	0.9416		Route	0.8894
Strength	0.9655	Strength	0.9428		Strength	0.9139
Average	0.9052	Average	0.9067		Average	0.8249
			0.0004			0.0007

0.810

Task

0.9052

0.8467

Median

Average

<u>Results</u>

0.7485

Task

Average

The classifier for all tasks is a linear SVM using the Liblinear library [4].

- With minimal domain-specific modeling, we were able to obtain results > 0.9 in tasks 1 and 2, and 0.82 for the end-to-end task.
- Novel challenges of the task ADE and Reason
 were some of the worst performing
- Scoring metric incentivized work on most prevalent categories

Blue indicates final scores, **bold** indicates trouble areas

Error Analysis

Reason/ADE

- Using Sign/Symptom/Disease/Disorder as Argument type is still too fine-grained (Recall errors)
- Drug categories are sometimes annotated as well as drugs, not always in ontologies (Recall errors)
- Reason and ADE relations are typically much farther apart than other relations, harder to memorize context between arguments (Recall errors)

Duration

• Confusion with time expressions (Precision errors)

References

- [1] S. Bethard, P. Ogren, L. Becker, ClearTK 2.0: Design Patterns for Machine Learning in UIMA., LREC (2014).
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- [3] D. Dligach, S. Bethard, L. Becker, T. Miller, G. K. Savova, Discovering body site and severity modifiers in clinical texts., Journal of the American Medical Informatics Association: JAMIA 21 (????) 448–54.
- [4] R.-E. Fan, K.-W. Chang, C.-J. Hsieh, X.-R. Wang, C.-J. Lin, LIBLINEAR: A Library for Large Linear Classification, Journal of Machine Learning Research 9 (2008) 1871–1874.

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