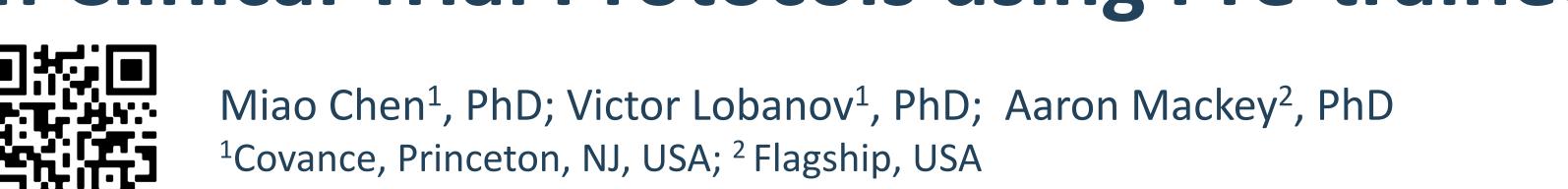


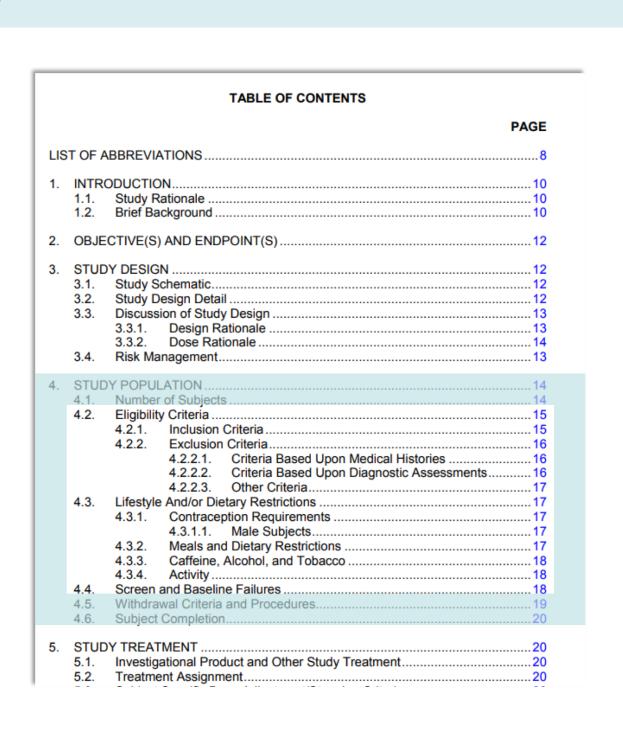
# Text Classification for Identifying Eligibility Criteria in Clinical Trial Protocols using Pre-trained Deep Learning Models





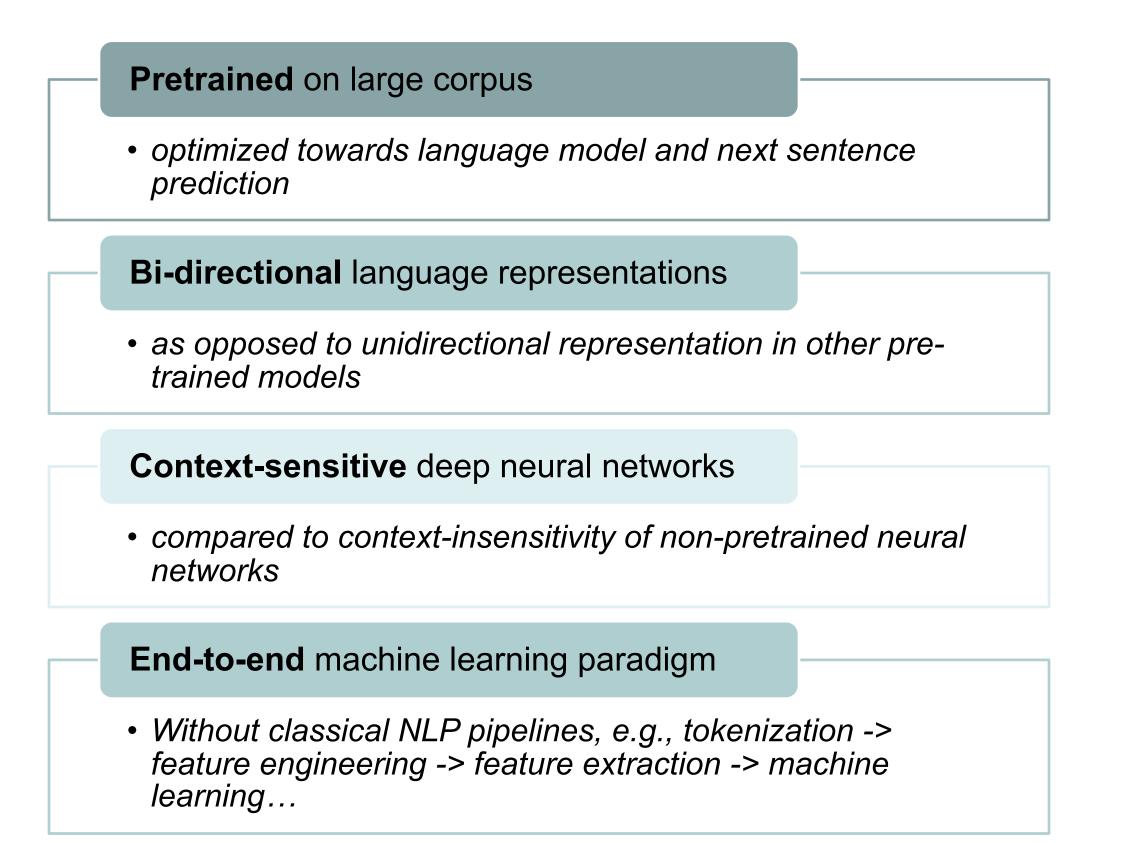
#### Background

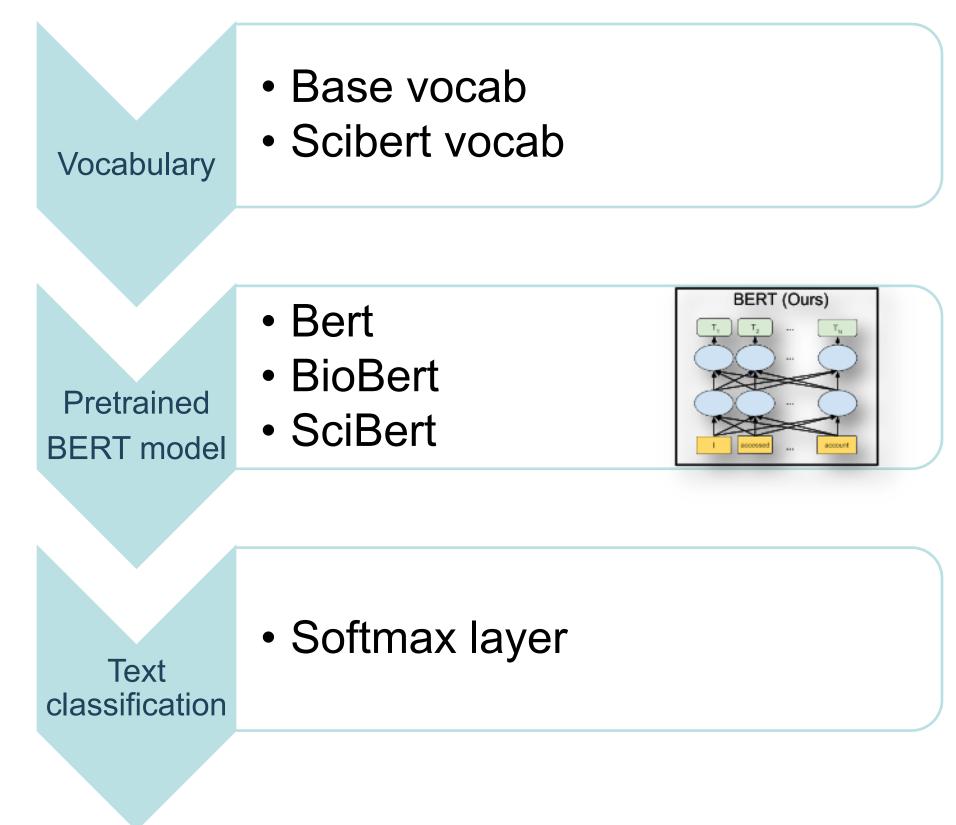
### Information extraction from clinical protocols



Importance of protocols and downstream tasks of protocol analytics	Challenges
For understanding operational requirements	Clinical protocols = semi-structured or unstructured document
For understanding systemic challenges	Lack of training and labeled
Implying probability of success	
cost implications and business planning	

#### Methods





#### Data & Methods

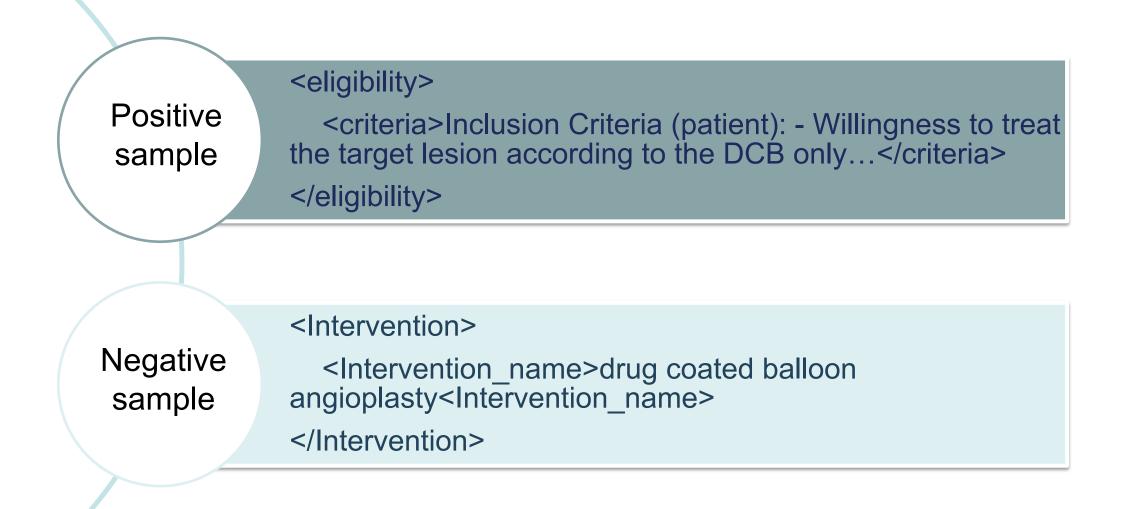
# Eligibility criteria data from clinicaltrials.gov (ct.gov)

- A small subset of ct.gov XML documents
- A XML doc contains multiple fields about a protocol,
   e.g. eligibility criteria, intervention, condition, etc.
- Used the eligibility criteria (EC) section of the XML docs as positive samples, and other sections (non-EC) as negative samples

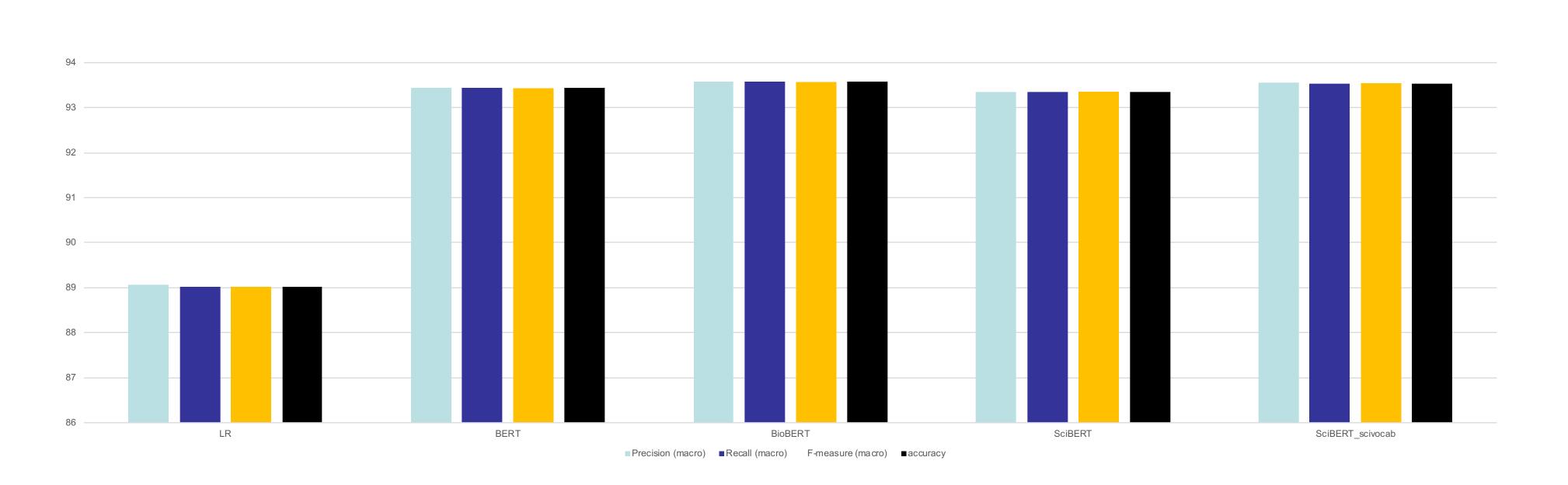
Eligibility criteria (EC)	Non-eligibility (non- EC)
Train – 14,000	Train - 14000
Test – 6,000	Test - 6,000

## Binary text classification task using pretrained deep learning model

- Using Bidirectional Encoder Representations from Transformers (BERT) as the pretrained language model
- Fine tuning the binary text classification task based on the BERT pretrained model
- Experimented with variants of BERT pretrained models:
   Bert/BioBert/Scibert



#### Results & Conclusions



Overall the BERT family models outperform the logistic regression baseline due to the deeper language representation and fine tuning

The BERT family models perform close to each other on our data set, likely due to the fact it was a small classification data set

Ongoing NLP efforts in Covance – entity recognition, syntactic relations, multi-task learning

contact: contact@ohdsi.org/2019-us-symposium-showcase-32/
BERT image from https://ai.googleblog.com/2018/11/open-sourcing-bert-state-of-art-pre.html



## Title here

Authors and affiliations

Abstract

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Methods

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Results

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Background

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Results

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Conclusions

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