Deciphering *Clinical Narratives* – key to automated Decision Making in Health Care

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TCS Research

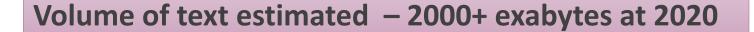
Outline of the talk

- Clinical Texts brief introduction
- Clinical Text Processing challenges
- Clinical NLP pipeline
- Clinical Text Processing Use-cases
 - Handling Clinical Trials
 - Predicting ICU stay for patients
 - Detecting Adverse effect of drugs

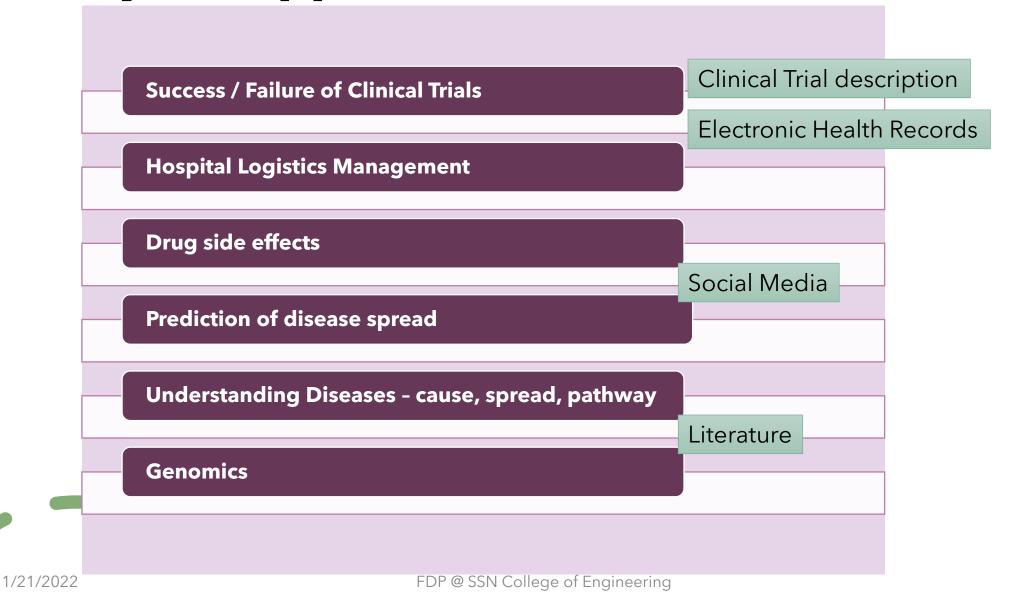
Clinical narratives - Main form of communication within health care

Clinical Data

- Electronic Medical Records (EMR)/Electronic Health Records (EHR) (classic)
- Physician and Care-given notes Account of patient history and assessments offering rich insights about clinical decision making
- Clinical trials management trial description for recruiting patients, monitor trial progress
- Social Media (tweets, Facebook comments, message boards, etc.)
 - personal accounts of patients signals for mental health adverse effects of drugs
 - Health care system feedback
- Medical Literature
 - News feeds, Medical journals
- Insurance Providers (claims from private and government payers)
 - Underwriter notes



Analytics applications



Analytics Life-Cycle

Ingest

Measure

Impact

Process

Natural Language Processing Knowledge - based Reasoning

Act

 Predict, Pre-empt, Reduce Risk Analyze



Clinical NLP - ingest and process

- Natural language processing (NLP) plays an important role in unlocking insights embedded in clinical narratives
- Machine learning is the back-bone that aids development of NLP tools by leveraging large amounts of text data
- ✓ Information Extraction
- ✓ Text Classification
- ✓ Paraphrase detection
- ✓ Content Matching
- ✓ Predictive Modeling
- ✓ Summarization



Challenge - Data is totally unstructured

 Bulk of this data does not exist in discretely-labeled fields - but rather available as completely unstructured free text clinical notes

 Traditional healthcare analytics depends predominantly on discrete data fields



Challenge - Unavailability of annotated data

- Adverse-drug effect?
- Bankruptcy?
- Humor in Uniform?



"You have a choice. An ultra-expensive medication that may cure you but, has the side-effect of bankruptcy, OR a low-priced medication with a side-effect of a near-death experience."

Privacy and Security concerns

It's more difficult to mask text data



"Excuse me doctor, could you spell that medical term? I'm updating all my social media friends about this lady's strange condition."

Clinical text is very different

- Domain specific features abound
- Shared vocabularies are needed
 - When we say breathless the doctor hears "dyspnea"
- Contextual interpretation
 - History
 - Ruling out
 - Grammar is not the mainstay
- Paraphrases
 - Obese with hyperinsulimenia
 - Type II Diabetes



"The Doctor will see you now. Here's your medical jargon dictionary."

Clinical NLP - pillars

Entities

Named Entity Detection and Classification Relations

Extraction Contextualization

Concept mapping

Paraphrase detection

Knowledge Graph

Reasoning



Named Entity Recognition

- A subtask of Information Extraction that aims to locate and classify Named Elements within unstructured text
- Named Entity Recognition and Classification into medical concepts
 - Diseases
 - Symptoms
 - Drugs
 - Genes
 - Chemicals

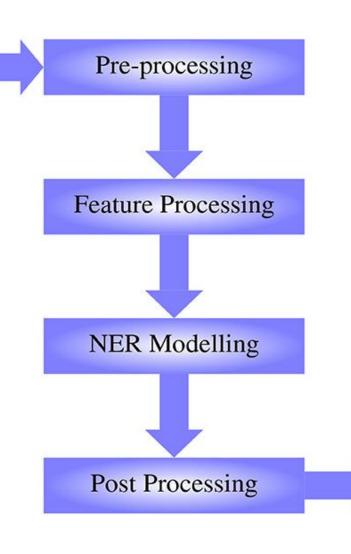
BioNER Challenges

- Boundary detection
 - 2,4,4,6-Tetramethylcyclohexa-2,5-dien-1-one," "epidemic transient diaphragmatic spasm
- Phrases
 - Alice in wonderland syndrome Neuropsychological condition
- Synonyms
 - Lymphocytic Leukemia" and "Lymphoblastic Leukemia"
- Sharing common Head noun in an article
 - "91 and 84 kDa proteins"
- Polysemy
 - *GLP1R* may refer to either the gene or protein needs resolution from context
- Non-standard abbreviations
 - CLD Cholesterol-lowering Drug," "Chronic Liver Disease," "Congenital Lung Disease," or "Chronic Lung Disease

Machine Learning Pipeline – detecting BioNER

E1A gene expression induces susceptibility to killing by NK cells following immortalization but not adenovirus of human cells. Adenovirus infection and E1A transfection were used to model changes in susceptibility to NK cell killing caused by transient vs stable E1A expression in human cells.

Unstructured Text



Structured Text

E1A gene (DNARegion)
expression induces susceptibility
to killing by NK cells (CellType)
following immortalization but not
adenovirus (Virus) of human
cells (CellType).

Adenovirus (Virus) infection and
E1A (ProteinMolecule)
transfection were used to model
changes in susceptibility to NK
cells (CellType) killing caused
by transient vs stable
E1A (ProteinMolecule)
expression in human
cells (CellType).

BioNER Detection and Classification

Rule Based

- Syntactic features
- Statistically significant features

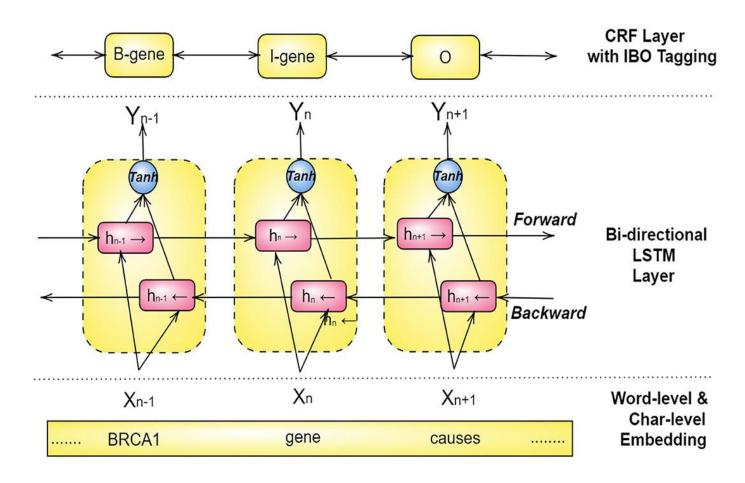
Support Vector Machines, Decision Trees, Conditional Random Fields

 Semantic features - Words, POS tags, dependencies

Deep Neural Networks

- LSTM with Word Embeddings sequence probabilities
- Transformers –
 Contextual Text
 Embedding

BIO tagging for NER



Sequence Labeling Architecture

A few words on building models

- Training is expensive
- Annotated resources are not available
 - Spent resources for building annotated data sets
 - Expert availability is a problem
- Transfer Learning
 - Trained on areas for which we have experts
 - Fine-tuned with small sets
 - Distant supervision

Annotated Entity Dataset

Dataset	Entity type	Number of annotations
NCBI Disease (Doğan et al., 2014)	Disease	6881
2010 i2b2/VA (Uzuner et al., 2011)	Disease	19 665
BC5CDR (Li et al., 2016)	Disease	12 694
BC5CDR (Li et al., 2016)	Drug/Chem.	15 411
BC4CHEMD (Krallinger et al., 2015)	Drug/Chem.	79 842
BC2GM (Smith et al., 2008)	Gene/Protein	20 703
JNLPBA (Kim et al., 2004)	Gene/Protein	35 460
LINNAEUS (Gerner et al., 2010)	Species	4077
Species-800 (Pafilis et al., 2013)	Species	3708

BioBERT – a pre-trained model for BioNER

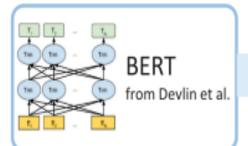
Pre-training of BioBERT

Pre-training Corpora

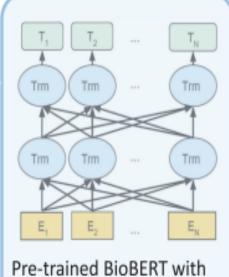




Weight Initialization



BioBERT Pre-training



biomedical domain corpora

Fine-tuning of BioBERT

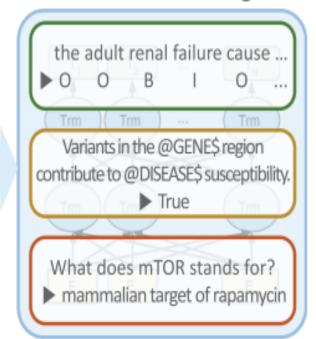
Task-Specific Datasets

Named Entity Recognition NCBI disease, BC2GM, ...

Relation Extraction EU-ADR, ChemProt, ...

Question Answering BioASQ 5b, BioASQ 6b, ...

BioBERT Fine-tuning



Bio-medical Named Entity Recognizers

Tools	Source	Description
CLiNER	https://github.com/text- machine-lab/CliNER	Clinical Named Entity Recognition system (CliNER) is an open-source natural language processing system for named entity recognition in clinical text of electronic health records.
C-NER	Clinical Named Entity Recognition (NER)	
SCiSpacy	https://allenai.github.io/s cispacy/	scispaCy is a Python package containing spaCy models for processing biomedical, scientific or clinical text.
PICO_Parser	https://github.com/Tian3 12/PICO_Parser	Recognize PICO Elements from Clinical Trial Literature - (Patient, Intervention, Condition, Outcome)
BIO_POS_DEP	_	Biomedical POS tagging and dependency parsing models

Example 2

Sample Text

The patient was given some hydrocodone for control of her pain. The patient suffers from bulimia and eating disorder, bipolar disorder, and severe hypokalemia.

Label	Id	Туре	Context
Hydrocodone	lxg-079584b7217f	Drug	Medication (Ingredients), Medication (Generic)
Control brand of phenylpropanolamine	lxg-b3f1a567f46b	Drug	Medication (Brand), Medication (Ingredients)
Pain	lxg-28ab8b90a066	Disease	Current issue
Bulimia	lxg-793bc328e6b8	Disease	<mark>C</mark>
Eating Disorders	lxg-aa5be6fbee41	MentalHealth	Chronic ailments
Bipolar Disorder	lxg-ca57a7a5d0c9	MentalHealth	ments
Hypokalemia	lxg-335c798f8bcc	Disease	Modifier

Analytics Tasks

Document Classification

No fever or chills. No nausea or vomiting. He had a mild fever and fainted. He does not recall falling. There is no apparent fracture.

Diagnosis

Deriving more context

Negation

The patient was given some hydrocodone for control of her pain. The patient suffers from bulimia and eating disorder, bipolar disorder, and severe hypokalemia.

Prescription

Deriving more context

Medication

Drug Dose

Different types of Contexts

- ✓ Negation and Hypothetical
- ✓ Vitals and Lab Values
- ✓ Medication dosage
- ✓ Anatomy
- √ History past / present
- ✓ Allergies
- ✓ Modifiers

Entities are not enough

Entities by themselves are not enough - relationship between entities or concepts around entities are more important

- The AIDS pandemic was caused by the spread of HIV infection.
- Infectious diseases or communicable diseases are caused by bacteria, viruses, and parasites.
- Serious adverse effect was observed in patients with heart disease due to high dosage of Spironolactone

Cause - effect relations

Example 1

Sample Text

No fever or chills. No nausea or vomiting. He had a mild fever and fainted. He does not recall falling. There is no apparent fracture.

Label	Id	Type Reference text	
Fever	lxg-45c0d6e146f4	Disease No [fever] or chills	
Chills	lxg-fc8fdc162ff2	Disease fever or [chills] . No	
Nausea	lxg-9bdf71250b2d	Disease No [nausea] or vomiting	
Vomiting	lxg-5f5bf2c446ed	Disease nausea or [vomiting] . He	
Fever	lxg-45c0d6e146f4	Disease a mild [fever] and fainted	
Syncope	lxg-7bf138779152	Disease fever and [fainted] . He	
Accidental Falls	lxg-91f1adb6e169	Other not recall [falling] . There .	
Fractures, Bone	lxg-2d8902924530	Disease to apparent [fracture]	
	SSN College of Engineering		

Presenting Contexts

No fever or chills. No nausea or vomiting. He had a mild fever and fainted. He does not recall falling. There is no apparent fracture.

```
"contexts": [
       "type": "negative",
       "subtype": "definite",
       "explanation": {
            "begin": 0,
            "end": 14,
            "matchedTokens": [
                    "token": "no",
                    "position": 0
                },
                    "token": "evidence",
                    "position": 2
                    "token": "of",
                    "position": 4
            "fuzzyValues": [],
            "triggerId": "N0000057",
            "triggerLabel": "no evidence"
```

1/21/2022 FDP @ SSN Collec

Relation extraction – an NLP task

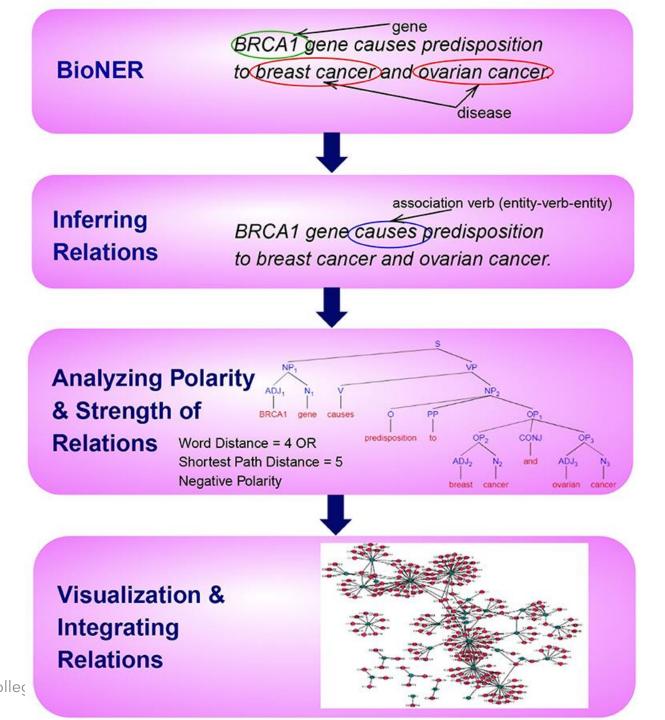
- Relations are defined between entities
 - Defines the context for entities
- Drugs *cure* disease
- Diseases are caused by organisms
- Auxins influence plant growth
- Genes are expressed through the process of protein synthesis.

Relation extraction – *Find relations embedded in text*

- Information about possible new drugs from Literature
- Side-effects from Social Media

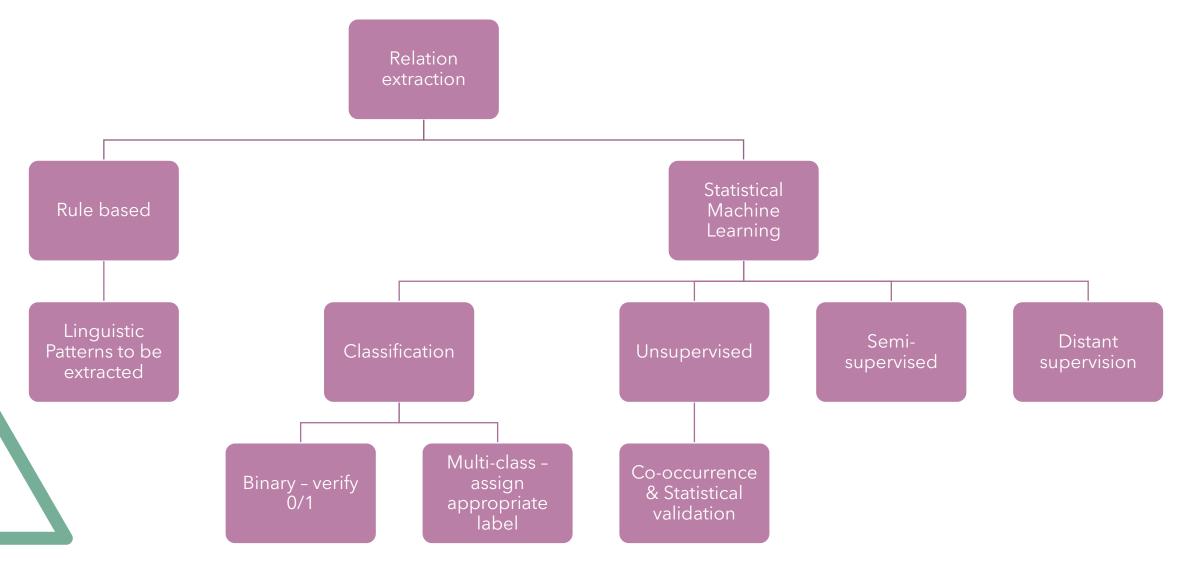


Inferring Relationships through Big Data Analytics



1/21/2022 FDP @ SSN Colleg

Bio-medical Relation Extraction



Distant Supervision for Relation Learning

- Applicable when a large pool of relevant but unlabeled data exists
 - Label a small sample using rules / heuristic / noisy operator / existing imperfect classifier
- Small pool of labeled data may also exist but not necessary
- Create a model utilizing the original labeled training data if it existed and the new noisily labeled data to create final output

Obtaining Annotations for Distant Supervision

- Weak Labels: Non-expert labels from crowdsourcing / heuristic rules
- Constraints: Specified as constraints entity-based constraints
- Distributions: Probability distribution
- Invariances: extend the coverage of the labeled distribution to all transformations e.g. if it is applicable for one entity, it is applicable to conjunction of similar entities

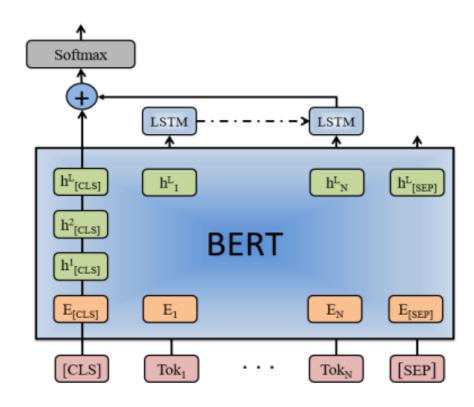
https://www.snorkel.org/blog/weak-supervision



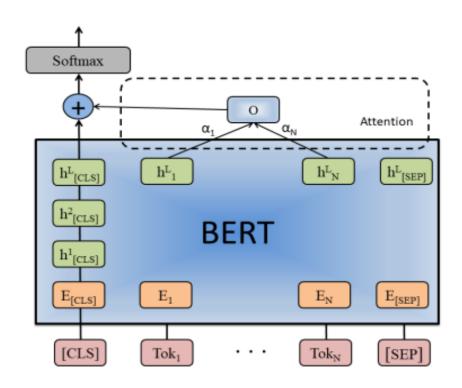
Relation extraction using BERT based classification

- Assume a set of relations
 - Example label set L = {ADVICE, EFFECT, INT, MECHANISM}
- Relation extraction as a classification problem
 - Does this text contain a Drug-Drug Interaction relation?
- Example -
 - S = Tok1 Tok2 . . . Tokn
 - Sentence has to contain drug occurrences, otherwise it will be seen as negative
 - A few tokens should be labeled by two entities Et1, Et2
 - Model needs to predict the probability of each label P(L|Tok1, Tok2, . . . , Tokn, Et1, Et2) based on the sentence text in the sentence

BERT based Architectures



(a) LSTM on the last layer



(b) Attention mechanism on the last layer

Focus shift needed

Dataset	Entity type	Number of relations
GAD (Bravo et al., 2015)	Gene-disease	5330
EU-ADR (Van Mulligen et al., 2012)	Gene-disease	355
CHEMPROT (Krallinger et al., 2017)	Protein-chemical	10 031

Relation between diseases and other risk factors

- nutritional habits
- life style
- side effects of combinations of drugs rather than single drugs alone
- analyzing the experience of patients on mass diseases such as asthma or diabetes

KnowLife: A knowledge graph for health and life sciences

KnowLife Portal Input Sources Knowledge Base hypertension UMLS Entities creates risk 4 has symptom Seed sickle cell smoking **Facts** anemia Relational creates risk erythromelalgia aggravates Patterns Raynaud's Health Portals: disease RxList Wikipedia Health Portal observed in causes Mayo Clinic lupus fingers Medline Plus Scientific Publications: diabetes Pubmed Medline mutation Pubmed Central External Online Communities: Professional Sources: Images: clinicaltrials.gov drugstalk.com pillbox pubmed.org user supplied text bodyparts3d

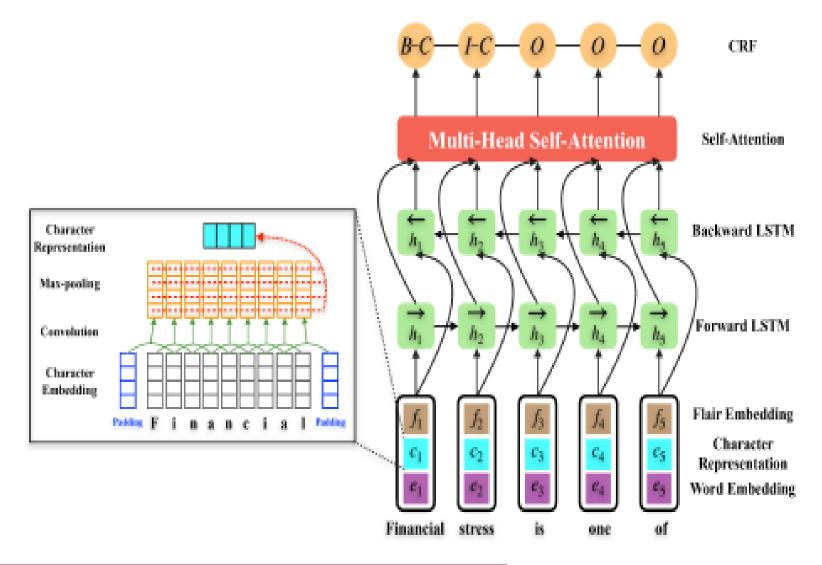
•Using advanced IE methods for constructing a large KB on a wide range of health-centric relations with entity linking to the Unified Medical Language System (UMLS, <u>uts.nlm.nih.gov</u>): a total of 214k canonical entities and 78k facts for 14 relations •Tapping into health-related online forums for evidence for relational facts and populate KG Automatically annotating new documents from scientific literature or from social media with relevant entities and relationships

Use Case 1 – Extract Causal relations from text

Identify drug side-effects

Deep Learning Mechanisms for Relation Extraction

- ✓ Use Deep recurrent Networks
- ✓ Train relation classifiers
- ✓ Attention mechanism to learn stronger correlations between words, motifs observed for a particular kind of relation

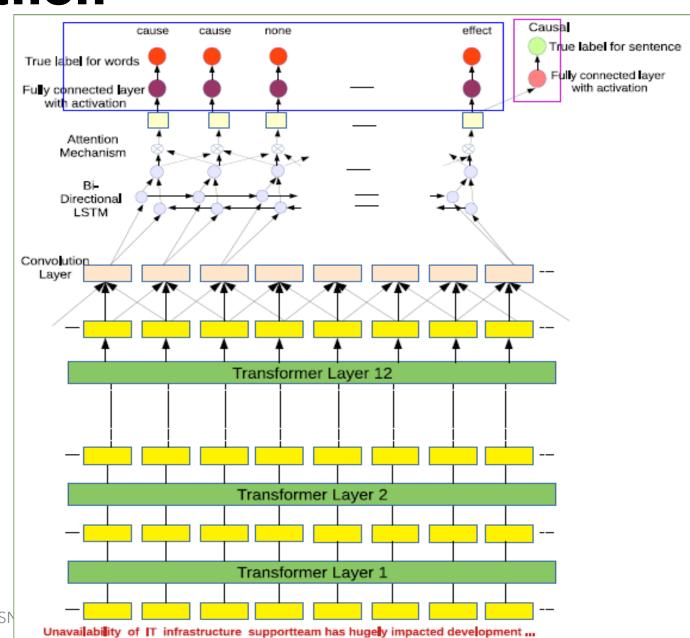




Self-Attentive BiLSTM-CRF with Transferred Embeddings (Li et al., 2020)

Causal Relation Extraction

- Proposed Joint Model for Causal Sentence Classification and Relation Extraction
- Uses a multi-layer bidirectional Transformer encoder architecture based on the transformer model
- 12 layers of transformer blocks, 768 hidden units, and 12 self-attention heads.
- Pretrained with two strategies on large-scale unlabeled text - masked language model and next sentence prediction.
- The pre-trained BERT model provides a powerful context-dependent sentence representation and can be used for various target tasks through the fine-tuning procedure.



Learning The Joint Model for Sentence Classification and Sequence Labeling Tasks

Loss function

$$L_1(\theta) = -\sum_{t=1}^{M} \sum_{k=1}^{K} \bar{y}_t^k log(y_t)$$

$$L_2(\theta) = -\sum_{t=1}^{N} \sum_{j=1}^{J} \bar{q}_t^{i,j} log(q_t^i)$$

$$L_{joint}(\theta) = \lambda * L_1(\theta) + (1 - \lambda) * I_{[y_{sentence} = 1]} * L_2(\theta)$$

Resource Creation - Causal fragments

Source	Sentence count	Average sentence length
Analyst Report (AR)	4500	23.7
SEMEVAL (SEM)	1331	18.7
BBC News (BBC)	503	22.5
ADE	3000	20.5
Recall News (RN)	1052	23.1

Results

Features	Dataset	Precision	Recall	F1-Score
	Analyst Reports(AR)	0.71 ± 0.11	0.87 ± 0.09	0.78 ± 0.11
BiLSTM	BBC News (BBC)	$0.71 \pm\ 0.14$	$0.82 \pm\ 0.03$	0.75 ± 0.08
	SEMEVAL(SEM)	$0.79 \pm\ 0.11$	$0.87 {\pm}~0.14$	$0.81 \pm\ 0.01$
	Adverse Drug(ADE)	$0.73 \!\pm 0.07$	0.86 ± 0.03	$0.75 \pm\ 0.06$
	Recall News (R)	$0.87 \!\pm 0.02$	0.93 ± 0.14	$0.87 \!\pm 0.10$
CNN+BiLSTM	Analyst Reports(AR)	0.84 ± 0.11	0.87 ± 0.04	0.82 ± 0.03
	BBC News (BBC)	$0.81 \!\pm 0.01$	$0.82 \pm\ 0.03$	$0.79 \pm\ 0.04$
	SEMEVAL(SEM)	$0.89 \!\pm 0.03$	0.87 ± 0.01	$0.86 \pm\ 0.05$
	Adverse Drug(ADE)	$0.83 \!\pm 0.07$	0.96 ± 0.02	$0.90 \pm\ 0.05$
	Recall News (R)	$0.87 \!\pm 0.08$	0.93 ± 0.05	$0.89 \!\pm 0.06$
$BERT_{base}$	Analyst Report	$0.87 {\pm}~0.14$	0.92 ± 0.09	0.85 ± 0.07
	BBC News (BBC)	$0.81 \!\pm 0.01$	0.92 ± 0.05	$0.84 \pm\ 0.06$
	SEMEVAL(SEM)	$0.91 {\pm}~0.05$	0.97 ± 0.03	$0.95 \pm\ 0.02$
	Adverse Drug(ADE)	$0.91 \!\pm 0.03$	0.96 ± 0.06	$0.93 \pm\ 0.05$
	Recall News (R)	$0.87 \!\pm 0.04$	0.93 ± 0.05	0.90 ± 0.07
	Analyst Report	$0.91 {\pm}~0.05$	0.97 ± 0.07	0.90 ± 0.02
	BBC	$0.94 \!\pm 0.13$	$0.97 \pm\ 0.16$	$0.94 \!\pm 0.09$
BERT+CNNBiLSTM	SEMEVAL	$0.91 {\pm}~0.15$	0.97 ± 0.07	$0.94 \!\pm 0.05$
	Adverse Drug	$0.89 {\pm}~0.05$	0.97 ± 0.01	$0.95 {\pm}~0.05$
	Recall News (R)	$0.87 \!\pm 0.02$	0.93 ± 0.01	$0.90 {\pm}~0.02$

Use Case 2 - Enabling Automatic Patient Recruitment For Clinical Trials

Clinical Trials

Research studies that are aimed at evaluating a medical, surgical, or behavioral intervention.

Whether a new treatment, like a new drug or diet or medical device is more effective than the existing treatments for a particular ailment.

Clinical Trial description - a sample

TITLE: Randomized Trial of Acetazolamide for Uveitis-Associated Cystoid Macular Edema

CONDITION: Macular Edema, Cystoid

INTERVENTION: Acetazolamide

SUMMARY: To test the efficacy of acetazolamide for the treatment of uveitis-associated cystoid

macular edema.

DETAILED DESCRIPTION: Uveitis, an intraocular inflammatory disease, is the cause of about 10 percent of visual impairment in the United States. Uveitis may lead to many sight-threatening conditions including cataract, vitreal opacities, glaucoma, and, most commonly, cystoid macular edema. Reduction of swelling or edema within the retina depends on the movement of fluid from the retina through the choroid. A number of studies indicate that this process requires active transport of fluid ions by the retinal pigment epithelium and may involve the carbonicm anhydrase system.

ELIGIBILITY Gender: All Age: 8 Years to N/A

Males and females 8 years of age or older and weighing at least 35 kg (77 lb) were eligible for the study. Patients had to have a best corrected visual acuity of 20/40 or worse in at least one eye with cystoid macular edema demonstrable on fluorescein angiography. Patients were allowed to receive systemic therapy for their uveitis. Exclusion criteria included current use of acetazolamide as part of a therapeutic regimen; a history of hypersensitivity reactions to acetazolamide, sulfonamides, or angiography dye; unclear ocular media that would obscure fluorescein angiography. Also, patients with macular subretinal for medical reasons may be excluded.

What is the analytics use-case here?

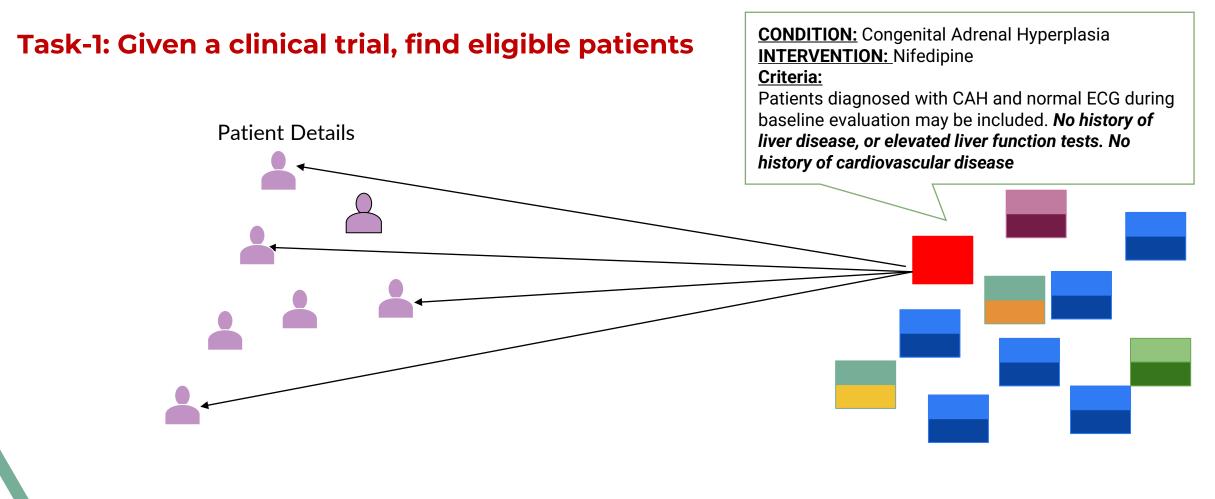
- ☐ A successful completion of a trial is dependent on achieving a significant sample size of patients enrolled into the trial within a limited time period.
- □ Identification of participants for a given trial is not trivial.
 - □ Involves repeated readings of the patient's electronic health record (EHR) for matching against the inclusion/exclusion conditions.

The Inclusion/Exclusion do not follow standard specification format – makes the problem more complex



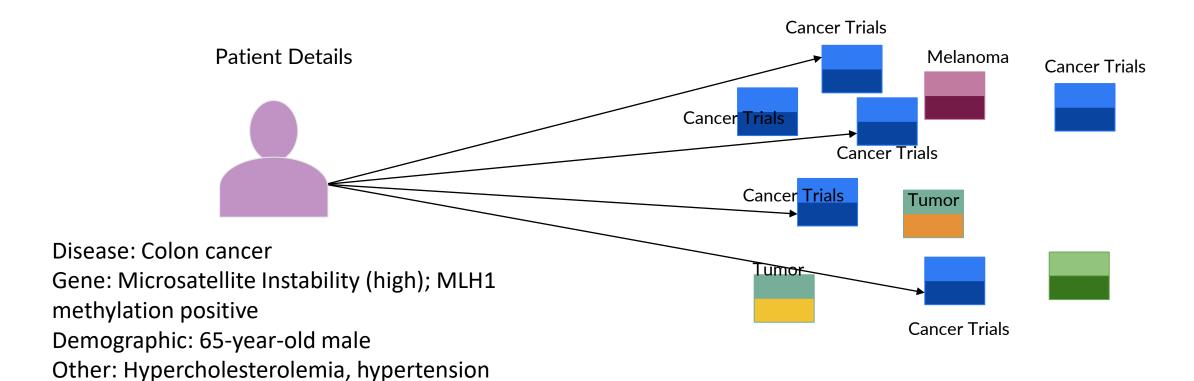
- Limits the number of patients that can be evaluated.
- May overlook adverse drug reactions

Text REtrieval Conference (TREC) task1

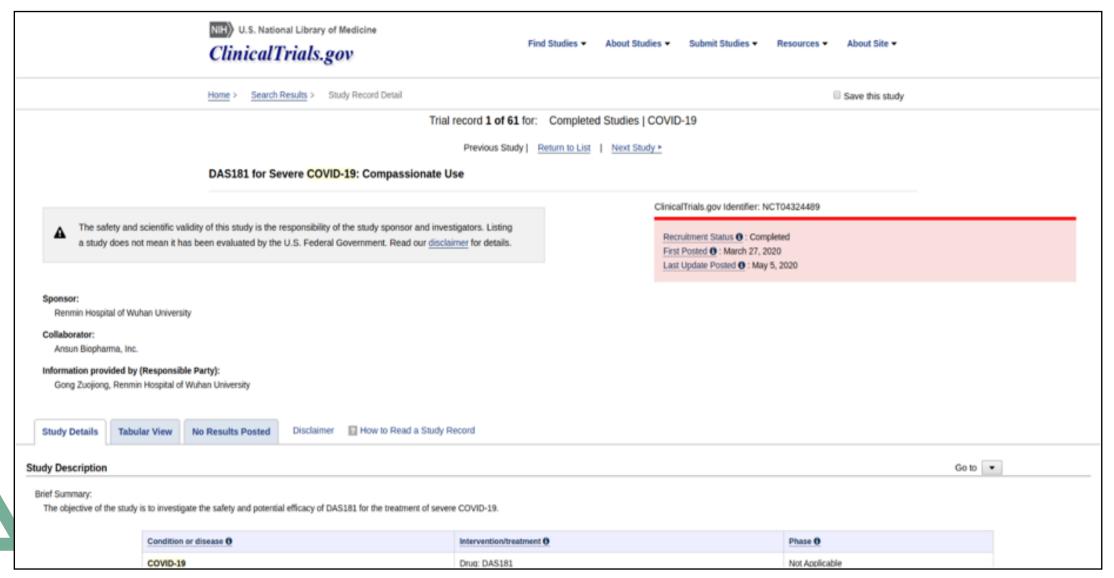


Text REtrieval Conference (TREC) task2

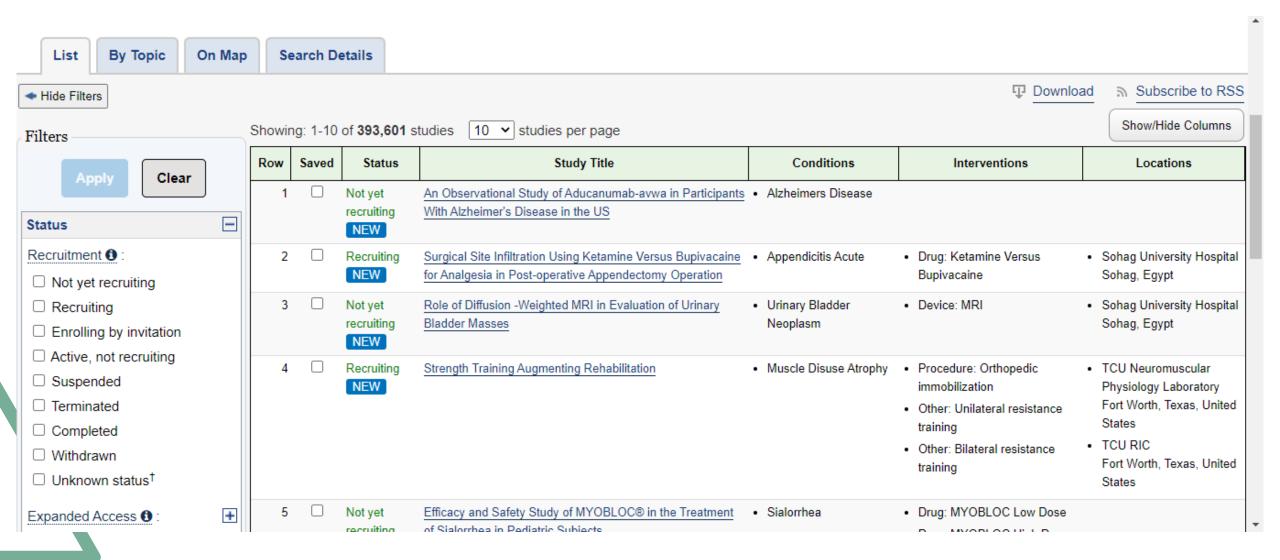
Task-2: Given a patient detail, find its relevant trials



Clinical Trial Search



Existing Search Engines/Filters



Problem Statement 1

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media that would obscure fluorescein angiography. Also, patients with macular subretinal for medical reasons

may be excluded.

Inclusion

Exclusion

Engineering

Matching Pipeline

Find Inclusion
- Exclusion
segments
within a
description

Within each segment - label text sequences as clinical aspects

Within each clinical aspect
- identify
Named
Entities with their labels

Detect
paraphrases different ways
of saying the
same thing

Derive match score using Named Entities

Inclusion - Patients had to have a best corrected visual acuity of 20/40 or worse in at least one eye with cystoid macular edema

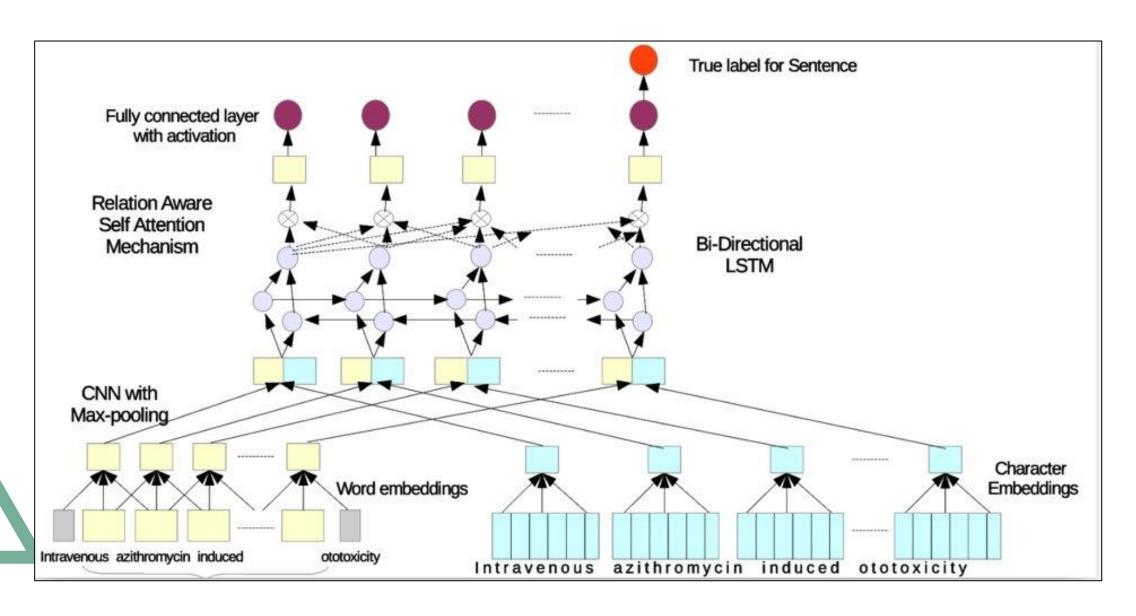
Exclusion - a history of hypersensitivity reactions to acetazolamide, **sulfonamides**, or angiography dye

Look up appropriate content in EHR

Automatic Segregation and Classification of Inclusion and Exclusion Criteria – Text Sequence Labeling task

Sentences	Prediction
Patients with active infections, including HIV, will be excluded, due to unknown effects of the vaccine on lymphoid precursors.	exclusion
Patients with HIV infection (but not AIDS) are eligible for this trial. Therefore, no HIV testing will be required.	Inclusion
HIV-1 infection as documented by any licensed ELISA (enzyme-linked immunosorbent assay) test kit and confirmed by Western blot at any time prior to study entry; HIV-1 culture, HIV-1 antigen, plasma HIV-1 ribonucleic acid (RNA), or a second antibody test by a method other than ELISA is acceptable as an alternative confirmatory test.	Inclusion
Patients who are HIV seropositive can have decreased immune competence and thus be less responsive to the experiment and more susceptible to its toxicities).	exclusion

Sequence Labeling Task



Dataset Available for Research

Source: TREC 2018 Precision Medicine Task, Clinical trials

Dataset-1	Inclusion instances	Exclusion instances	Total instances
Training Data	10,000	12,280	22,280
Test Data	8,000	5,700	13,700

https://www.kaggle.com/auriml/eligibilityforcancerclinical trials

Dataset-1	Inclusion instances	Exclusion instances	Total instances
Training Data	29,855	32,280	61865
Test Data	11,000	12,000	23,000

Results

(AAAI (W) on Health Intelligence, 2020)

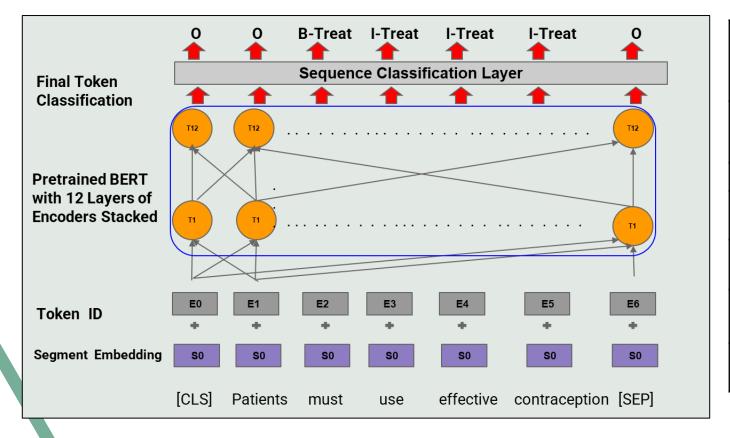
	Dataset-I (TREC-2019)					Dataset-II (Kaggle)						
Word Embedding	Inclusion			Exclusion			Inclusion			Exclusion		
	W	FT	E	W	FT	\mathbf{E}	W	FT	\mathbf{E}	W	FT	Е
BiLSTM	0.70	0.74	0.78	0.72	0.72	0.76	0.70	0.73	0.80	0.71	0.75	0.76
BiLSTM*	0.73	0.76	0.80	0.74	0.75	0.78	0.72	0.77	0.84	0.73	0.76	0.78
CNN	0.68	0.67	0.68	0.65	0.65	0.70	0.71	0.73	0.62	0.61	0.65	0.70
CNN*	0.69	0.71	0.73	0.67	0.66	0.74	0.79	0.77	0.67	0.63	0.68	0.71
C-BiLSTM	0.70	0.72	0.76	0.71	0.71	0.74	0.73	0.74	0.81	0.70	0.72	0.79
C-BiLSTM*	0.77	0.75	0.81	0.73	0.73	0.78	0.75	0.77	0.86	0.72	0.75	0.80
$C - BiLSTM - S_{att}$	0.75	0.78	0.82	0.72	0.73	0.80	0.82	0.83	0.89	0.78	0.85	0.88
$C - BiLSTM - S_{att}*$	0.80	0.82	0.83	0.78	0.76	0.81	0.84	0.86	0.90	0.82	0.88	0.90
$C - BiLSTM - Re_{att}$	0.81	0.79	0.84	0.79	0.80	0.81	0.85	0.87	0.91	0.86	0.85	0.91
$C - BiLSTM - Re_{att}*$	0.84	0.84	0.89	0.82	0.81	0.85	0.87	0.87	0.95	0.88	0.90	0.93
BERT-base	Dataset-I				Dataset-II							
	Inclusion Exclusion				on	In	clusi	on	Ex	clusi	ion	
BERT-base		0.86			0.80			0.91			0.92	

Problem-2: Matching EHRs against trial – more complex problem

- Medical Literature proposed five different clinical aspects
 - Lab Test Results
- Health Status
- Treatment status
- Demography
- Lifestyle

Total bilirubin less than or equal to 1.5 mg/dl, except in patients with history of anaemia. Have had their ileostomy or colostomy for at least 3 months. Subjects must be between the ages of 18 to 65 years old and must not intake alcohol. Pregnant women of stage 3 and age 18 years and older attending delivery room. Life expectancy of at least 6 months and willing to provide informed consent. Live vaccine within 4 weeks prior to therapy or potential need for a live vaccine. Serum ALAT or serum ASAT > 5 x upper limit of normal (ULN) at screening. Current alcohol abuse or drug addiction that in the opinion of the investigator Lab Test Results Health status Treatment status Demography status Life style (Lou et al., 2011)

Transformer based Architecture for Clinical Aspect Extraction – multi-class classification problem



	Baselin	e (LST	M)	Our	Model				
	Precision	Recall	F1	Precision	Recall	F1			
Health	0.67	0.68	0.68	0.78	0.75	0.77			
Demography	0.92	0.97	0.94	0.94	0.96	0.95			
Treatment	0.69	0.71	0.70	0.71	0.79	0.74			
Lab Test	0.79	0.83	0.81	0.82	0.83	0.82			
Lifestyle	0.62	0.76	0.69	0.72	0.83	0.77			

(Clinical Natural Language Processing, 2020)

Problem-3: Different manifestations of the same criteria- Paraphrase detection

history of traumatic brain or head injury with loss of consciousness

diagnosis of traumatic brain injury

history of tbi

history of known brain metastases

subjects with a history of stroke or traumatic brain injury

volunteers must have history of at least one mild traumatic brain injury

- The ability to provide informed consent before any trial-related activities.
- written informed consent to participate in the study Ability and willingness to give written informed consent
- The child and parent or legal guardian is able to provide assent and/or consent.
- Willingness and ability to sign informed consent document
- Provide assent and have a legal guardian that will participate and provide parental permission.
- Able to comprehend and willing to provide written consent

Obese with Hyperinsulinemia can be included in the study

Patients with Type-II Diabetes can be included

Detecting different manifestations of the same criteria classification - similar / dissimilar (ICDMAI, 2020)

	P	R	F			
BiLSTM Siamese	0.51	0.48	0.49			Fully connected
CNN	0.45	0.47	0.45			Layer with sigmoid activation
C-BiLSTM Siamese	0.60	0.68	0.63			
BERT	0.75	0.78	0.76			
BERT+KB	0.81	0.79	0.79			
		T_[d:	Ī	T_N T_[se	P. T_1 T_M	Similarity matrix
						Knowledge base UMLS
		E_[cl		E_N E_[se	1	Sentence1 Sentence2
				Sentence1	Sentence2	

Matching Patients to Trials – work in progress

Disease: Colon cancer

Gene: Microsatellite Instability (high);

Demographic: 25-year-old female

Other: Early pregnancy

Disease: Colon cancer

Gene: MLH1 methylation positive

Demographic: 65-year-old male

Other: Hypercholesterolemia,

SGOT=73.0, history of hypertension

Involves aspect resolution followed by matching.

Total bilirubin less than or equal to 1.5 mg/dl, except in patients with history of anaemia. Have had their ileostomy or colostomy for at least 3 months. Subjects must be between the ages of 18 to 65 years old, able to provide informed consent and must not intake alcohol. Life expectancy of at least 6 months and willing to provide informed consent. Live vaccine within 4 weeks prior to therapy or potential need for a live vaccine. SGOT and SGPT < 3x the normal, Serum ALAT or serum ASAT $> 5 \times 10^{-2} \text{ m}$ (ULN) at screening. Current alcohol abuse or drug addiction that in the opinion of the investigator, Hypertensive patients Pregnant women of stage 3 and age 18 years and older must be excluded.

Use Case 3 – Predicting ICU Length of stay

Using Clinical Notes for ICU stay prediction

- Predicting length of stay in ICU helps in better logistics planning ensures better resource usage for critically ill patients
- MIMIC Dataset a publicly available dataset was developed by the Laboratory for Computational Physiology
 - Comprises deidentified health data associated with thousands of intensive care unit admissions
 - The dataset is widely used by investigators and engineers around the world to drive research in clinical informatics, epidemiology, and machine learning
- During hospitalization a nursing notes contain information about patient's condition, nursing assessments, care provided to a patient.
- Objective
 - Predict length of stay of a patient in Intensive care unit at the time of hospital admission from their nursing notes of first 24 hours.



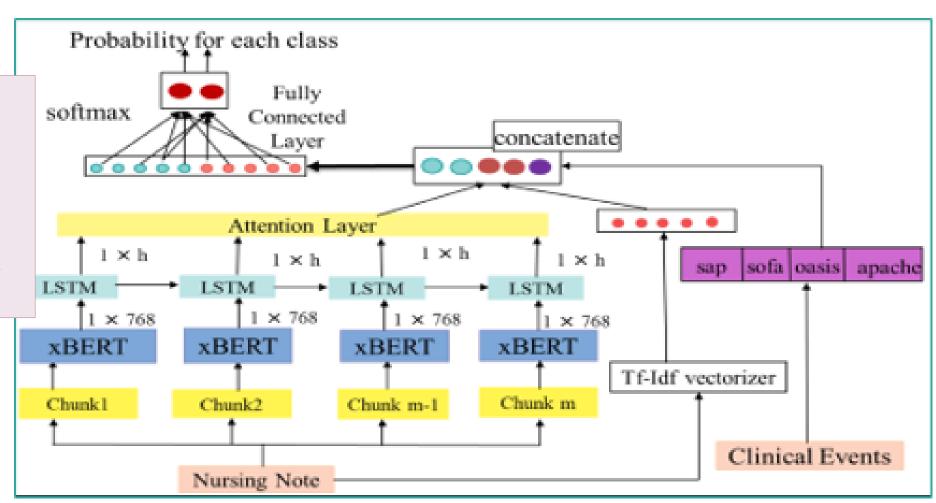
	Dataset	Feature used	Method	Best Result
Alghatani et al., 2021	44,000 ICU stays from MIMIC	patient's vital signs like, heart rate, BP, temp., resp. etc	Random Forest	65% accuracy
Su et al., 2021	2224 Sepsis patients PICMISD	Age, P(v-a)CO ₂ /C(a-v)O, SO,wbc etc.	XG-Boost model	F1: 0.69, AUC- ROC:0.76
Rocheteau,Liò, et al., 2020	elCUcritical care dataset	medical features, Gender, Age, Ethnicity, etc.	Temporal convolution	Kappa score = 0.58
Harutyunyan et al., 2019	42276 ICU stays of 33798 uniquepatients from mimic database	17 clinical variables like, Capillary refill rate, Diastolicblood pressure etc. from first 24 hours of admission.	LSTM	AUC-ROC: 0.84
van Aken et al., 2021	38013 admission notes from MIMIC III	Created admission notes from discharge summaries	Pretrained CORe +Bi oBERT	AUC-ROC : 0.72%
Improved upon SOTA	22789 Nursing Notes from MIMIC III	nursing notes + TF-IDF Vector+ SOI scores	BlueBERT+LSTM+TF -IDF+SOI	Acc: 79%; AUC- ROC:0.87; Kapp a: 0.59

Prediction Architecture

Explainability

- Attempts to understand the model - by perturbing the input of data samples and assess how the predictions change
- Weighted average from multihead attention models

Long notes



Indicative Phrases

- •Phrases like "HR dropping", "requiring mask ventilation for resp. failure", "couldn't breathe"
 i indicative of high risk patients needing longer ICU stays
- •"good effect from Ativan", "comfortable breathing", "hemodynamically stable"

 short ICU stays.

- Many more problems
 - •Trigger alerts
 - More exact predictions
 - •Correlations between vital parameters and clinical notes
 - Contra-indications

Summary

- Clinical texts contain a wealth of information about state of patients, healthcare
 - Long Covid
 - Precision Medicine
- Automation possibilities to reduce risks
 - Physician's Aid Expert insights
- Explainable, Secure models are needed
- Resource-light models are needed

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"Doctor and physician are outdated terms. I'm your biological tech support specialist."

Thank You

References

• Perera et al. Named Entity Recognition and Relation Detection for Biomedical Information Extraction - https://internal-journal.frontiersin.org/articles/10.3389/fcell.2020.00673/full

