

Automated Patient Screening for Clinical Trials

Overview of the literature and challenges

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March, 12th 2019

Introduction : matching patients to clinical trials

Ontology + rule based feature extraction

Deep (representation) learning methods ?

Conclusion

Introduction : matching patients to clinical trials

- Procedure to assess new drug safety and efficiency
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- Need to select (screen) cohort of patients satisfying *eligibility criteria*
- Screening usually done **manually**, very **time consuming** (bottleneck in the CT process)
- Generalization of **electronic health records (EHRs)** can alleviate such tasks

Typical Clinical Trial

- Title, Summary, Condition name, Interventions
- List of **inclusion** and **exclusion** criteria (free text)
- <https://clinicaltrials.gov>

Electronic Health Record (EHR)

EHRs of hospital patients typically contains

- **Structured** data (age, demographic data, treatments, physical characteristics : BMI, blood pressure, *etc.*)
- **Unstructured** (free text) data (clinical narratives, progress notes, imaging reports, discharge summaries)

- Clinical trials descriptions : all on <https://clinicaltrials.gov>
- EHRs from patients : 50000 deidentified EHRs (for research, English) (without matching data)

Formalization of the matching problem

$x \in \mathcal{X}$ represents a patient's EHR

$y \in \mathcal{Y}$ represents a trial (list of criteria)

Goal :

find $f : \mathcal{X} \times \mathcal{Y} \rightarrow \{0, 1\}$

such that $f(x, y) = 1$ iff $x \in \mathbf{Elig}(y)$ (x is eligible for y).

Metrics ?

Given x_1, \dots, x_p patient records, y_1, \dots, y_T trials, and $M \in \{0, 1\}^{p \times T}$ assignment matrix such that $M_{i,j} = 1$ if patient i participated in trial j and 0 otherwise,

$$P = \sum_{\text{trial } j} \frac{\sum_{\text{patient } i} f(x_i, y_j) M_{i,j}}{\sum_{\text{patient } i} f(x_i, y_j)}$$

$$R = \sum_{\text{trial } j} \frac{\sum_{\text{patient } i} f(x_i, y_j) M_{i,j}}{\sum_{\text{patient } i} M_{i,j}}$$

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- Leverage common criteria across different trials ?

Formalization of the matching problem (ctd.)

Each trial = combination of inclusion / exclusion criteria.

$z \in \mathcal{Z}$ represents a criterion

$y_j = (z_j^{(1)}, \dots, z_j^{(n_j)})$ Goal :

find $\phi : \mathcal{X} \times \mathcal{Z} \rightarrow \{0, 1\}$

such that $\phi(x, z) = 1$ iff $x \in \mathbf{Elig}(z)$ (x satisfies z).

And $\tilde{M}_{i,k} = M_{i,j}$ for $k = 1, \dots, n_j$, for all trial j .

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Challenges

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- Still $\tilde{M}_{i,k} \neq \mathbb{1}[x_i \in \mathbf{Elig}(z_k)]$
- **No matching data** yet. Can we still make progress using proxys ?

Intermission : ICD10 classification

International Classification of Diseases (codes with descriptive sentence to tag patients' diseases. Essentially used for billing)

▼ ICD-10 Version:2016

▶ I Certain infectious and parasitic diseases

▼ II Neoplasms

▼ C00-C97 Malignant neoplasms

▶ C00-C75 Malignant neoplasms, stated or presumed to be primary, of specified sites, except of lymphoid, haematopoietic and related tissue

▼ C76-C80 Malignant neoplasms of ill-defined, secondary and unspecified sites

▶ C76 Malignant neoplasm of other and ill-defined sites

▼ C77 Secondary and unspecified malignant neoplasm of lymph nodes

C77.0 Secondary and unspecified malignant neoplasm: Lymph nodes of head, face and neck

C77.1 Secondary and unspecified malignant neoplasm: Intrathoracic lymph nodes

C77.2 Secondary and unspecified malignant neoplasm: Intra-abdominal lymph nodes

C77.3 Secondary and unspecified malignant neoplasm: Axillary and upper limb lymph nodes

C77.4 Secondary and unspecified malignant neoplasm: Inguinal and lower limb lymph nodes

C77.5 Secondary and unspecified malignant neoplasm: Intrapelvic lymph nodes

C77.8 Secondary and unspecified malignant neoplasm: Lymph nodes of multiple regions

C77.9 Secondary and unspecified malignant neoplasm: Lymph node, unspecified

▶ C78 Secondary malignant neoplasm of respiratory



C77.1 Intrathoracic lymph nodes

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Pectoral lymph nodes

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C77.5 Intrapelvic lymph nodes

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C77.9 Lymph node, unspecified

C78 Secondary malignant neoplasm of respiratory and digestive organs

C78.0 Secondary malignant neoplasm of lung

C78.1 Secondary malignant neoplasm of mediastinum

C78.2 Secondary malignant neoplasm of pleura

Malignant pleural effusion NOS

C78.3 Secondary malignant neoplasm of other and unspecified respiratory organs

C78.4 Secondary malignant neoplasm of small intestine

C78.5 Secondary malignant neoplasm of large intestine and rectum

C78.6 Secondary malignant neoplasm of retroperitoneum and peritoneum

Malignant ascites NOS

C78.7 Secondary malignant neoplasm of liver and intrahepatic bile duct

C78.8 Secondary malignant neoplasm of other and unspecified digestive organs

C79 Secondary malignant neoplasm of other and unspecified sites

C79.0 Secondary malignant neoplasm of kidney and renal pelvis

Intermission : ICD10 classification

International Classification of Diseases (codes with descriptive sentence to tag patients' diseases. Essentially used for billing)

- Well-posed classification (multilabel or multiclass) problem : input EHRs, output : ICD code (class)
- CNN works well with input text EHRs (Mullenbach et al. 2018)

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- Here, x and z is text. Represent x and z in same space (translation-like problem ?)
- Old-fashioned NLP : use ontology + NER to extract features. Broadly used for clinical text.

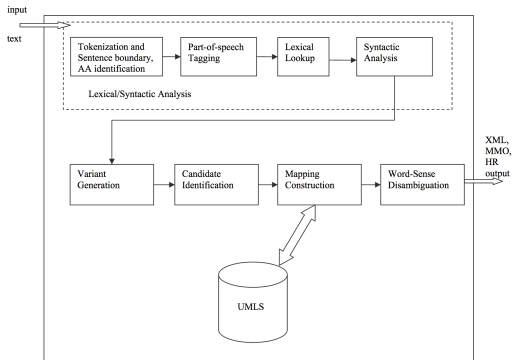
Ontology + rule based feature extraction

- ICD10 : disease codes with descriptive sentences
- MeSH (Medical Subject Headings) : thesaurus of controlled vocabulary used for PubMed indexing. Each term has short description and relations to other terms
- SNOMED CT : hierarchical+relational structure between classes of concepts
- UMLS : “Meta-thesaurus”. Millions of concept codes associated with descriptives and relations between them

Mapping text to clinical concepts

Tools using NER and/or UMLS (parse text and map to concepts)

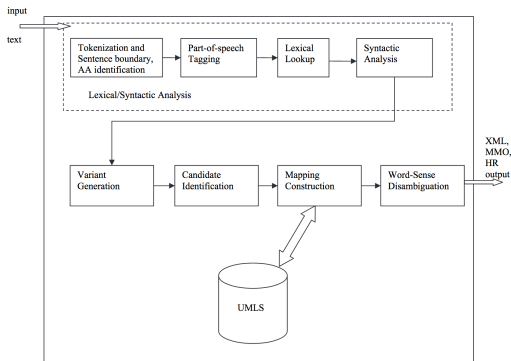
- MetaMap (https://ii.nlm.nih.gov/Interactive/UTS_Required/metamap.shtml) (Figure from Aronson & Lang (2010)), cTAKES, DNorm



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- ConText, NegEx : regex-based tools to find negative or context (family) in medical documents



Garcelon et al. (2016)

- context of rare diseases : text search may be sufficient
- family history important (e.g. father has Crohn disease)
- Text search + negation and context (family) yields good performance

Finding patients for clinical trials : use mapping to ontology to find similar patients

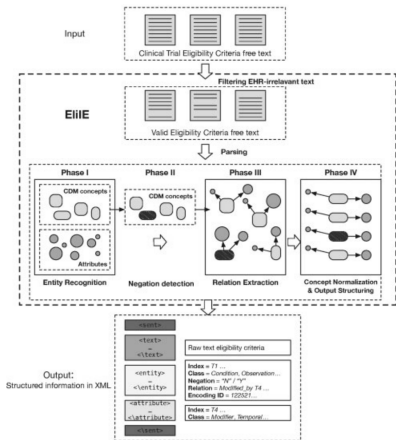
Garcelon et al. (2017)

- context of rare diseases : sparse set of relevant clinical concepts
- Method : map EHR to UMLS concepts to find representation vector of patients
- (Incorporate context and negation disambiguation)
- Given patient with rare disease, identify potentially similar patients based on their EHR

Use ontology-based mapping to extract information from clinical trials description

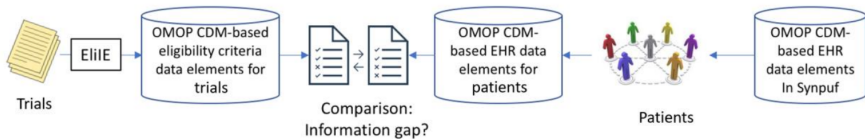
Kang et al. (2017)

- Goal : structure concepts in EC with terminology common to EHRs concepts (“normalization”)
- Specific entity recognition for eligibility criteria (relation between criteria, *etc.*)
- Fine-tuned on Alzheimer’s disease eligibility criteria



Join the dots between CT and EHRs : “the data gap”

Butler et al. (2018)



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- Involves manual unification of the clinical terms in EC before concept extraction

Table 1. Manual revision of clinical entities.

Types of Revision	Example	Times
Formatting; Typo	delerium -> delirium	207
Formatting; Plural	cancers -> cancer	253
Formatting; removal of non-informative words	heart rate measurement -> heart rate	364
Formatting; removal of abbreviations	absolute neutrophil count (ANC) -> absolute neutrophil count	1768
Simplification	asthmatic conditions -> asthma	573
Breaking down long phrases to logically-connected single phrases	basal or squamous cell carcinoma -> basal cell carcinoma or squamous cell carcinoma	445
Total		3610

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Join the dots between CT and EHRs : “the data gap”

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- Goal : Assess intersection of concepts extracted from EC and EHRs
- Involves manual unification of the clinical terms in EC before concept extraction
- Also on Alzheimer's disease data
- Intersection not so broad

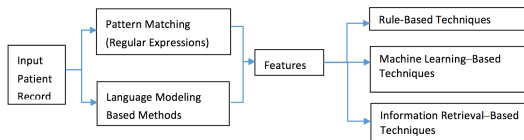
Table 4. The top 20 common SNOMED CT terms in AD trials and their prevalence in EHR dataset.

SNOMED CT Term	SNOMED-CT ID	Trial Count	Prevalence in Trials	Count of uses in EHR data for AD patients
<i>Alzheimer's disease</i>	26929004	972	64.29%	30,262
<i>Mini-mental state examination</i>	273617000	705	46.63%	0
<i>Presenile dementia</i>	12348006	599	39.62%	7,089
<i>Disease</i>	64572001	555	36.71%	12,029,900
<i>Current chronological age</i>	424144002	515	34.06%	0
<i>Mental disorder</i>	74732009	499	33.00%	505,870
<i>Magnetic resonance imaging</i>	113091000	482	31.88%	63,171
<i>Cerebrovascular accident</i>	230690007	371	24.54%	4
<i>Global assessment of functioning - 1993 Diagnostic and Statistical Manual of Mental Disorders- ver.4th</i>	284061009	361	23.88%	0
<i>Systemic disease</i>	56019007	353	23.35%	0
<i>Disorder of nervous system</i>	118940003	335	22.16%	780,478
<i>Substance abuse</i>	66214007	279	18.45%	9,466
<i>Parkinson's disease</i>	49049000	275	18.19%	0
<i>Impaired cognition</i>	386806002	260	17.20%	13,375
<i>Seizure disorder</i>	128613002	240	15.87%	28,586
<i>Hypersensitivity reaction</i>	421961002	218	14.42%	4,686
<i>Schizophrenic disorders</i>	191526005	216	14.29%	40,777
<i>History of clinical finding in subject</i>	417662000	207	13.69%	189,543
<i>Risk identification: childbearing family</i>	386414004	205	13.56%	0
<i>Clinical dementia rating scale</i>	273367002	204	13.49%	0

Extract information from EHRs: domain specific rules

Adupa et al. (2016)

- EHR information extraction method for a given clinical trial (PARAGON)



Extract information from EHRs: domain specific rules

Adupa et al. (2016)

- EHR information extraction method for a given clinical trial (PARAGON)
- Domain specific rules (Heart Failure)

Table 4. Regular expressions for extracting LVEF-containing sentences and values.

S/N	Regular Expression
1	(left ventricular ejection fraction lv lv ejection fraction left ventricle ejection fraction ejection fraction ef ejection fraction)[^_%\\,]*?((\\d-\\.\\,)+)\\s*;%
2	(left ventricular systolic function left ventricular function systolic function of the left ventricle lv systolic function left ventricular ejection fraction ejection fraction left ventricle)(normal normal global low normal well preserved severely reduced moderately decreased moderately depressed severely decreased severe markedly decreased markedly reduced severely globally reduced mildly decreased mildly depressed severely depressed)
3	(normal normal global low normal well preserved severely reduced moderately decreased moderately depressed severely decreased severe markedly decreased markedly reduced severely globally reduced mildly decreased mildly depressed severely depressed)
4	.*(moderate marked severe) (lv systolic dysfunction left ventricular dysfunction left ventricular systolic dysfunction).*
5	((\\d+\\s*(\\ - to \\s*\\d+)/(\\d*\\.\\d* \\s*(\\ - to \\s*\\d*\\.\\d* \\d* \\d+)/(\\d+)/(?=\\s*(\\%)))
6	\\d+{\\.\\d+)?

Extract information from EHRs: domain specific rules

Adupa et al. (2016)

- EHR information extraction method for a given clinical trial (PARAGON)
- Domain specific rules (Heart Failure)
- Goal : save time for prescreening with high recall

		Prescreening Gold Standard (Manual)	
		Patients Included	Patients Excluded
Classification outcome (algorithmic)	Patients included	38	6
	Patients excluded	2	152

Deep (representation) learning methods ?

- Think of Computer Vision
- Now transfer learning works with text too (BERT, ELMO, etc.)
- Unsupervised methods ? (Word2Vec)
- Yet, not always satisfying in domain-specific tasks (even in CV)

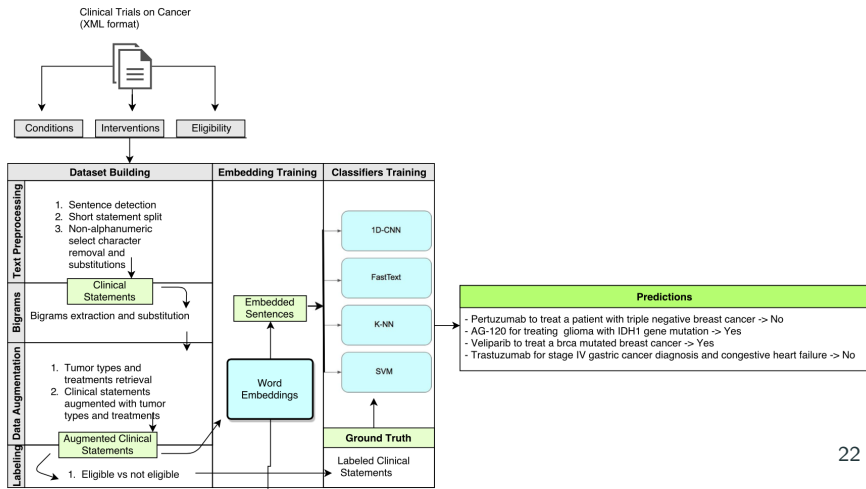
Training deep representation of clinical trials with a random classification task

Bustos & Pertusa (2018)

- Goal : train deep neural network (CNN) to obtain accurate embedding of clinical text (words)
- Task : classify statements as True or False (Eligible / Not eligible)
- Data : uses data from `clinicaltrials.gov` only) to generate data (labeling given by inclusion/exclusion, data augmentation through simple sentences)
- Belief in the magic of word embeddings

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Training deep representation of clinical trials with a random classification task

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Condition	Intervention
Rectal Neoplasms	Procedure: Irreversible electroporation (IRE)
Criteria Inclusion Criteria: <ul style="list-style-type: none">Rectal Neoplasms diagnosed by positive biopsy or non-invasive criteria,Not suitable for surgical resection,Eastern Cooperative Oncology Group (ECOG) score of 0-1,A prothrombin time ratio > 50%,Platelet count > 80x10⁹/L,Ability of patient to stop anticoagulant and anti-platelet therapy for seven days prior to and seven days post NanoKnife procedure,Able to comprehend and willing to sign the written informed consent form (ICF),Have a life expectancy of at least 3 months. Exclusion Criteria: <ul style="list-style-type: none">Cardiac insufficiency, ongoing coronary artery disease or arrhythmia,Any active implanted device (eg Pacemaker),Women who are pregnant or women of child-bearing potential who are not using an acceptable method of contraception,Have received treatment with an investigational agent/ procedure within 30 days prior to treatment with the NanoKnife™ LEDC System,Are in the opinion of the Investigator unable to comply with the visit schedule and protocol evaluations.	

A. Original Source: <https://clinicaltrials.gov/ct2/show/NCT02425059>

Label	Clinical Statement
Eligible	study intervention is irreversible electroporation . rectal neoplasm and not suitable for surgical resection
Not eligible	study intervention is irreversible electroporation . rectal neoplasm and cardiac insufficiency ongoing coronary artery disease or arrhythmia

B. Extracted features after preprocessing

Conclusion

Summary, TODOs, challenges and open questions

- Matching unstructured text data (EHRs) to unstructured text (Clinical Trials)
- Goal : prescreen patients with high recall, and provide reasonable number of patients for manual screening
- Domain restriction allows information retrieval with specifically designed rules (e.g., Alzheimer's or Heart Failure)
- Degree of precision for matching also depends on domain restriction (e.g., just output patients with "Heart Failure" in their EHR ?)
- Evaluate baselines (text-search and concept mapping tools)
- Make progress without matching data (other, simpler task (e.g., classification of diseases))
- Annotate data ?
- Reliably augment the matching data (e.g. with patient similarity, or leveraging external corpus or ontology)

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