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Can Electronic Clinical Notes Identify Travelers with Zika?

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BACKGROUND / OBJECTIVES

Travel history can help differentiate a public health emergency from a travel-related infection by providing information on exposure. However such information is often available only in unstructured clinical documents.¹ We are not aware of existing work has reported on feasibility of automated extraction of travel history, likely due to a need for annotated data and a process for selecting data.

We aimed to assess feasibility of extracting past travel history mentions from the electronic health record in an automated fashion by first annotating a dataset and then developing a machine learning model to extract travel history locations from clinical documents.

METHODS

In collaboration with the National Biosurveillance Integration Center (NBIC), clinical notes were extracted from patient records for encounters with Zika, dengue and chikungunya virus testing in the Department of Veterans Affairs.

- Extracted 4,584 snippets from a set of >250k using a semi-automated bootstrapping process to identify documents containing potentially relevant information using locations and phrases (see right).²
- Manually annotated snippets for travel affirmation and locations visited including negation states (i.e. "pt denies travel to Puerto Rico")
- However, time period of the travel (if stated) was not annotated.
- Trained a Conditional Random Field (CRF) model to extract affirmed travel locations outside of the continental US.



Locations

- "recently returned from X..."
- "traveled on a cruise to X..."
- "10 day vacation to
- "Symptoms after a 12 hour flight from X"







Annotation agreement results Bootstrapping selection of annotation corpus Agree Disagre Type 769 24 Text Span **Annotation results** Phrases **Travel Evidence** Unique GeoNames.org (Any) **Snippets** Countries Goal : Acquire 4584 3006 (65.6%) Cities documents with Popular location names Most frequent annotated locations likely history Additional locations (e.g. "the big island", **Positive Locations** "caribbean islands", etc) Iraq Mexico Dominican Republic Costa Rica Vietnam **MODELING RESULTS** Model performance on 356 held out test documents **Positive Predictive Val** from Uganda months ago 85.6 Most frequent model extractions from over 250k notes **Model contextual features Extracted Location**

	Examples (i – 2, i – 1, i)
	Verb, Preposition, Noun,
	"returned", "from", "Uganda",
	"ret", "ned", "fro", "rom", "uga", "nda",
tion, digit, punctuation)	"Capital=False", "Capital=False", "Capital=True",
	"return", "from", "uganda",
ers ³	"213", "10", "320",
npound Clusters" ³	"213_10", "10_320",
ames.org lexicon)	False, False, True,

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2	Missing	% Agree	Карра
	75	89%	0.651

Evidence	Evidence
(Positive)	(Negated)
2659 (58%)	347 (7.6%)

Negated Locations
Liberia
Guinea
Sierra Leone
Democratic Republic of Congo
West Africa

е	Sensitivity	F1
	76.7	80.9

Count
571
569
392
301
259
254
160
134

Vietnam

Iraq

Mexico

Costa Rica

Dominican Republic

Afghanistan

Jamaica

Honduras



- Targeted travel history extraction is feasible in a large medical system with acceptable accuracy
- Approach capable of extracting novel locations that would not necessarily be found in a curated list (e.g. Mexican Riviera, Baja Cruise)
- Further research could incorporate automated extractions into models improving the early detection of autochthonous transmission
- Approach may also be beneficial to other biosurveillance (e.g. screening for CRE)
- Automated extraction now deployed in operations for continued validation

REFERENCES

- Chapman WW, Gundlapalli AV, South BR, Dowling JN. Natural language processing for biosurveillance. InInfectious Disease Informatics and Biosurveillance 2011 (pp. 279-310). Springer, Boston, MA
- Alba PR, Patterson OV, Viernes B, Denhalter DW, Bailey N, Wilson A, Kamauu AW, DuVall SL. The Super Annotator: A Method of Semi-Automated Rare Event Identification for Large Clinical Data Sets. In AMIA 2016.
- 3. Guo J, Che W, Wang H, Liu T. Revisiting embedding features for simple semi-supervised learning. In Proceedings of the 2014 Conference on Empirical Methods in Natural Language Processing (EMNLP) 2014 (pp. 110-120).

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