EACL 2014

14th Conference of the European Chapter of the Association for Computational Linguistics



Proceedings of the 5th International Workshop on Health Text Mining and Information Analysis (Louhi)

> April 27, 2014 Gothenburg, Sweden



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ISBN 978-1-937284-90-9

Introduction

Welcome to Louhi 2014: The Fifth International Workshop on Health Text Mining and Information Analysis, in Gothenburg, Sweden. The Louhi workshop series is an international, scientific, forum for researchers and practitioners in the multidisciplinary area of health text mining and information analysis. This research area has progressed and grown since the First Louhi workshop in Turku, 2008, addressing challenging research issues in the health and biomedical domain, leading to more openly available annotated corpora, tools, terminologies, etc. Moreover, work on other languages than the previously dominating English is both increasing and maturing.

The importance of accurate and specific information extraction and classification from diverse health and biomedical documents such as Electronic Health Records (EHRs), scientific literature and online health forums is evident for several differing purposes, e.g. detecting drugs and medications, adverse events, building timelines, understanding information needs. Ontologies and terminologies for such tasks are also crucial. The papers presented in this workshop all address these issues from different perspectives, and on different languages such as Basque, Spanish, French, Danish, Swedish, German and English. The Fifth Louhi workshop will provide a platform for important and useful discussions in this vivid research area, and hopefully lead to many more fruitful endeavours.

We received in total 21 submissions from 11 countries and three continents, and after a rigorous doubleblind peer-review process we could accept 17 of these submissions (nine long papers and eight short papers) to be published in the 2014 Louhi proceedings. The acceptance rate for long papers was 60%, while the overall acceptance rate for the workshop was 81%.

A short description of each paper follows in order of appearance.

Long papers:

Medical queries in a Swedish health portal are studied from the perspective of supporting information needs by using semantic- and graph-based methods (*Moradi et al.*).

Collier et al. experiment with five strategies for mitigating the impact of near domain transference for biomedical named entity recognition. Distributional dissimilarities of domains need to be adequately compensated during learning, or else lower performance and higher annotation costs are expected.

Zhao and Tou Ng also address domain adaptation, but in the area of coreference resolution, using active learning methods and target domain instance weighting.

Discourse parsing is addressed in the paper by *Stepanov and Riccardi*, where cross-domain evaluation of a discourse relation parser trained on one domain generalises well across domains with feature-level domain adaptation.

Perez-de-Viñaspre and Oronoz present initial work on semi-automatically translating SNOMED CT into Basque, using the English version of SNOMED CT as source and then adapting the terms to Basque utilizing various rules.

A method for building FrameNet-like corpora using ontologies is described in the paper by *Tan*. The system includes algorithms for selecting and describing appropriate concepts to be translated into semantic frames.

Quan and Ren describe work on gene-disease association extraction by combining information filtering, grammar parsing and network analysis. With breast cancer as testing disease, they achieve 83.9% accuracy for the testing genes and diseases and 74.2% accuracy for the testing genes.

Segura-Bedmar et al. describe work on detecting drugs and adverse events from Spanish health social media posts. A gold standard is created for evaluation, and a multilingual text analysis engine, Textalytics, was applied for automatic detection, achieving 80%/56% precision, and 87%/85% recall for drugs/adverse events.

The paper by *Moen et al.* presents several methods for information retrieval, focusing on care episode retrieval based on textual similarity using distributional semantics and ICD-10 codes of diagnoses, to retrieve the most similar care episodes among the records.

Short papers:

Engel Thomas et al. present work on text mining of Danish health records, with a focus on handling spelling and ending variations, gaps and shuffling of terms, as well as negation identification and scope. Spelling variation was found to be the most important functionality.

A new annotated corpus for identifying phenotype information for congestive heart failure is presented by *Alnazzawi et al.* This corpus is unique in that it integrates information both from electronic health records and literature articles.

Medication extraction, as defined in the 2009 i2b2 challenge, is addressed by using agile text mining methods in the paper by *Shivade et al.* They report results of 92% precision and 71.5% recall.

Roller and Stevenson describe work using the Unified Medical Language System (UMLS) for distantly supervised relation extraction.

Casillas et al. describe work on extracting cause-effect relations between drugs and diseases, applied on Spanish health records.

Semantic relations are integrated in a vector space model to tackle the problem of context-unawareness and applied on an electronic health record corpus (*Périnet and Hamon*).

Kreuzthaler and Schulz present work on disambiguating period characters in German clinical discharge summaries. An accuracy of 93% is reported for abbreviation detection and sentence delimitation.

The Heideltime system is used for identifying time expressions in English and French in the paper by *Hamon and Grabar*. Results of 0.94 (French) and 0.85 (English) F-measure are reported on their adapted version of the system.

Dear reader, most welcome to study these proceedings, which we hope will raise interest, open new perspectives and generate new exciting research questions in health text mining and information analysis.

Stockholm and San Diego, March 2014

Sumithra Velupillai, Martin Duneld, Maria Kvist and Hercules Dalianis

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Invited Speaker:

Sophia Ananiadou, University of Manchester, U.K.

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	Keynote Talk
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	Session I
10:05	A Graph-Based Analysis of Medical Queries of a Swedish Health Care Portal Farnaz Moradi, Ann-Marie Eklund, Dimitrios Kokkinakis, Tomas Olovsson and Philippas Tsigas
10:30	Coffee
	Session II
11:00	<i>The impact of near domain transfer on biomedical named entity recognition</i> Nigel Collier, Mai-vu Tran and Ferdinand Paster
11:25	<i>Domain Adaptation with Active Learning for Coreference Resolution</i> Shanheng Zhao and Hwee Tou Ng
11:55	<i>Towards Cross-Domain PDTB-Style Discourse Parsing</i> Evgeny Stepanov and Giuseppe Riccardi
12:20	Lunch Break

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Session III

- 13:45Translating SNOMED CT Terminology into a Minor LanguageOlatz Perez-de-Viñaspre and Maite Oronoz
- 14:10A System for Building FrameNet-like Corpus for the Biomedical Domain
He Tan
- 14:35 *Gene-disease association extraction by text mining and network analysis* Changqin Quan and Fuji Ren

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Negation scope and spelling variation for text-mining of Danish electronic patient records Cecilia Engel Thomas, Peter Bjødstrup Jensen, Thomas Werge and Søren Brunak

Building a semantically annotated corpus for congestive heart and renal failure from clinical records and the literature Noha Alnazzawi, Paul Thompson and Sophia Ananiadou

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Session IV

16:00	Detecting drugs and adverse events from Spanish social media streams Isabel Segura-Bedmar, Ricardo Revert and Paloma Martínez
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16:50	Closing Remarks by Sumithra Velupillai, Department of Computer and Systems Sciences (DSV), Stockholm University