BioNLP 2018

# SIGBioMed Workshop on Biomedical Natural Language Processing

# **Proceedings of the 17th BioNLP Workshop**

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### Biomedical natural language processing in 2018: Spotlight on Deep Learning

Dina Demner-Fushman, Kevin Bretonnel Cohen, Sophia Ananiadou, and Jun-ichi Tsujii

The number of community challenges, corpora and publicly available tools in the domain continues to grow rapidly. The past year has seen several hackathons, a variety of shared tasks and growing numbers of workshops dedicated to specific biomedical and clinical sublanguages and tasks. The BioNLP meeting has now been ongoing for 17 years. BioNLP continues to stay the flagship and the generalist in biomedical language processing, accepting noteworthy work independently of the tasks and sublanguages studied. The quality of submissions continues to impress the program committee and the organizers. BioNLP 2018 received 28 submissions, of which 13 were accepted for oral presentation and 12 as poster presentations. This year, Deep Learning approaches are explored in the overwhelming majority of the papers, with focus on interesting new models and in-depth exploration of the state-ofthe-art publicly available tools. As for the past several years, the themes in this year's papers and posters continue to focus equally on clinical text and biological language processing, as well as reveal growing interest in consumer language processing. The morning session presents clinical text processing for extraction of causes of death, risk factors identification and named entity recognition, among others. The next session presents work on fundamental NLP problems, such as ontology alignment and key-phrase extraction, whereas the afternoon session presents exceptionally strong work on complex text mining tasks, such as event extraction and question answering.

The invited talk and the invited presentation reflect thus growing interest in automated support for systematic reviews of the literature. In the invited talk, professor Paul Glasziou discusses progress and challenges in automating systematic reviews. Paul Glasziou, FRACGP, PhD is Professor of Evidence-Based Medicine at Bond University and a part-time General Practitioner. He was the Director of the Centre for Evidence-Based Medicine in Oxford from 2003-2010. His key interests include identifying and removing the barriers to using high quality research in everyday clinical practice. He is the author of six books related to evidence based practice: Systematic Reviews in Health Care, Decision Making in Health Care and Medicine: integrating evidence and values, An Evidence-Based Medicine Workbook, Clinical Thinking: Evidence, Communication and Decision-making, Evidence-Based Medicine: How to Practice and Teach EBM, and Evidence-Based Medical Monitoring: Principles and Practice. He has authored over 160 peer-reviewed journal articles and his h-index is currently 43. He is the recipient of an NHRMC Australia Fellowship which he commenced at Bond University in July, 2010.

The invited presentation follows suit by bringing to our attention a new corpus of about 5,000 abstracts of randomized control trials annotated with granular information regarding the study populations, interventions, comparators and outcomes.

### **Organizers:**

Kevin Bretonnel Cohen, University of Colorado School of Medicine, USA Dina Demner-Fushman, US National Library of Medicine Sophia Ananiadou, National Centre for Text Mining and University of Manchester, UK Jun-ichi Tsujii, National Institute of Advanced Industrial Science and Technology, Japan

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### **Invited Speaker:**

Paul Glasziou, Bond University, Australia

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## **Conference Program**

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9:30–9:45	Multi-task learning for interpretable cause of death classification using key phrase prediction Serena Jeblee, Mireille Gomes and Graeme Hirst
9:45-10:00	<i>Identifying Risk Factors For Heart Disease in Electronic Medical Records: A Deep Learning Approach</i> Thanat Chokwijitkul, Anthony Nguyen, Hamed Hassanzadeh and Siegfried Perez
10:00-10:15	Keyphrases Extraction from User-Generated Contents in Healthcare Domain Using Long Short-Term Memory Networks Ilham Fathy Saputra, Rahmad Mahendra and Alfan Farizki Wicaksono
10:15–10:30	<i>Identifying Key Sentences for Precision Oncology Using Semi-Supervised Learning</i> Jurica Ševa, Martin Wackerbauer and Ulf Leser

10:30–11:00 Coffee Break

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### 11:00–12:30 Session 2: Foundations

- 11:00–11:15 *Ontology alignment in the biomedical domain using entity definitions and context* Lucy Wang, Chandra Bhagavatula, Mark Neumann, Kyle Lo, Chris Wilhelm and Waleed Ammar
- 11:15–11:30 Sub-word information in pre-trained biomedical word representations: evaluation and hyper-parameter optimization Dieter Galea, Ivan Laponogov and Kirill Veselkov
- 11:30–11:45 PICO Element Detection in Medical Text via Long Short-Term Memory Neural Networks
   Di Jin and Peter Szolovits
- 11:45–12:00 *Coding Structures and Actions with the COSTA Scheme in Medical Conversations* Nan Wang, Yan Song and Fei Xia
- 12:00–13:30 Lunch break
- 13:30–14:30 Invited Talk: "Automating systematic reviews: progress and challenges" Paul Glasziou

### 14:30–15:30 Session 3 Literature mining and retrieval; Question Answering

- 14:30–14:45 A Neural Autoencoder Approach for Document Ranking and Query Refinement in Pharmacogenomic Information Retrieval Jonas Pfeiffer, Samuel Broscheit, Rainer Gemulla and Mathias Göschl
- 14:45–15:00 Biomedical Event Extraction Using Convolutional Neural Networks and Dependency Parsing Jari Björne and Tapio Salakoski
- 15:00–15:15 BioAMA: Towards an End to End BioMedical Question Answering System Vasu Sharma, Nitish Kulkarni, Srividya Pranavi, Gabriel Bayomi, Eric Nyberg and Teruko Mitamura
- 15:15–15:30 Phrase2VecGLM: Neural generalized language model-based semantic tagging for complex query reformulation in medical IR
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15:30–16:00 Coffee Break

16:00–16:15 Invited Presentation: "A Corpus with Multi-Level Annotations of Patients, Interventions and Outcomes to Support Language Processing for Medical Literature" – Ben Nye

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Convolutional neural networks for chemical-disease relation extraction are improved with character-based word embeddings Dat Quoc Nguyen and Karin Verspoor

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