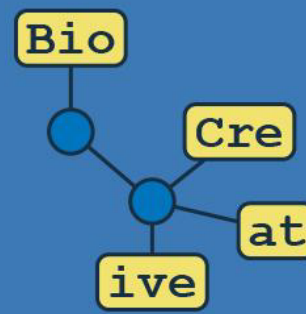


BioCreative 5 Challenge Workshop



Sevilla, Spain
9-11 Sept 2015

First name	Last name	Ref.	Title
Yung-Chun	Chang	A-01	Protein-protein Interaction Passage Extraction Using the Interaction Pattern Kernel Approach for the BioCreative 2015 BioC Track
FERHAT	AYDIN	A-02	Retrieving Passages Describing Experimental Methods using Ontology and Term Relevance based Query Matching
Rezarta	Islamaj Dogan	A-03	Identifying genetic interaction evidence passages in biomedical literature
Sérgio	Matos	A-04	Neji – Heterogeneous biomedical concept recognition
Yifan	Peng	A-05	Extended dependency graph for BioC-compatible protein-protein interaction (PPI) passage detection in full-text articles
Riza	Batista-Navarro	A-06	Development of bespoke machine learning and biocuration workflows in a BioC-supporting text mining workbench
Hong-Jie	Dai	A-07	NTTMUNSW BioC Modules for Recognizing and Normalizing Species and Gene/Protein Mentions in Full Text Articles
Thaer	Moustafa Dieb	B-01	Comparison of different strategies for utilizing two CHEMDNER corpora
Ruben	Rodrigues	B-02	BioTML: a machine learning framework for biomedical text mining
Daniel	Lowe	B-03	Recognition of chemical entities in patents using LeadMine
Sérgio	Matos	B-04	Identification of chemical and gene mentions in patent texts using feature-rich conditional random fields
Riza	Batista-Navarro	B-05	Adapting ChER for the recognition of chemical mentions in patents
Hsin-Chun	Lee	C-01	An enhanced CRF-based system for disease named-entity recognition and normalization
Dingcheng	Li	C-02	Constructing a UIMA-based Chemical Induced Disease Detection Pipeline with Rich Features
Christian	Hänig	C-03	ExB Medical Text Miner
Alberto	Lavelli	C-04	A Knowledge-Poor Approach to BioCreative V DNER and CID tasks
Tong Shu	Li	C-05	Extracting structured chemical-induced disease relations from free text via crowdsourcing
Daniel	Lowe	C-06	Use of Wikipedia as a resource for disease identification and concept mapping
Nigel	Collier	C-07	The UET-CAM System in the BioCreative V CDR Shared Task
Àlex	Bravo	C-08	Combining crowdsourcing; expert knowledge and machine learning to detect chemical-side effect relationships
TILIA	Ellendorff	C-09	Ontogene Term and Relation Recognition for CDR
Riza	Batista-Navarro	C-10	Augmenting the Medical Subject Headings vocabulary with semantically rich variants to improve disease mention normalisation
Po-Ting	Lai	D-01	NCU-IISR System for BioCreative BEL Task 1
Yalbi I.	Balderas-Martínez	E-01	Biocuration of microRNAs in idiopathic pulmonary fibrosis
Evangelos	Pafilis	E-02	EXTRACT: Interactive extraction of environment metadata and term suggestion for metagenomic sample annotation
Sérgio	Matos	E-03	Egas - Biomedical Annotation as a Service
Cecilia	Arighi	E-04	User Advisory Group (UAG) Activities in BioCreative
Sumit	Madan	E-05	BELIEF Dashboard – a Web-based Curation Interface to Support Generation of BEL Networks
Riza	Batista-Navarro	E-06	Semi-automatic curation of chronic obstructive pulmonary disease (COPD) phenotypes using Argo
Hong-Jie	Dai	E-07	MET Pathway in PubMed: A Pathway Visualization and Curation System
Claire	Nedellec	F-01	A knowledge model of regulatory networks involved in Arabidopsis seed development for information extraction and integration from text
Mary	Schaeffer	F-02	Literature Curation at MaizeGDB
Hugo	Costa	F-03	A text mining approach for generating transcriptional regulatory networks
Hugo	Costa	F-04	@Note2 – open source computational tools for biomedical text mining
shruti	rao	F-05	Enhancing evidence for personalized cancer treatments by curating public literature on biomarkers that predict response to chemotherapy
Nancy	George	F-06	SourceData: making data discoverable
Cecilia	Arighi	F-07	Text Mining Tools for Biocuration in the iProLINK Web Portal
Kimberly	Van Auken	F-08	TextpressoCentral: A System for Integrating Full Text Literature Curation with Diverse Curation Platforms