

BioCreative III Meeting Agenda
September 13 - 15, 2010
Double Tree Hotel
Bethesda, Maryland, USA

Meeting Location: Second level
Registration: Outside Ballroom A/B
All presentations: Grand Ballroom A/B
Demo: Grand Ballroom A/B
Posters and Reception: Ballroom C

September 13, 2010 (Monday)	
7:30 AM – 5:00 PM	Registration
7:30 AM – 8:30 AM	Breakfast (provided, outside of Ballroom A/B in the EMC Foyer)
8:30 AM – 8:40 AM	Workshop Opening – Cathy Wu
8:40 AM – 9:10 AM	Discussion of systems results (GN) – Zhiyong Lu, NCBI
9:10 AM – 10:40 AM	Challenge Participant systems 1 (GN) – chaired by John Wilbur Team 63 University of Colorado by Karin Verspoor Team 65 University of Zurich by Fabio Rinaldi Team 68 Arizona State University by Martin Gerner Team 70 Universidade de Aveiro, IEETA by Sergio Matos Team 74 Institute of Information Science, Academia Sinica, Taiwan by Chun-Nan Hsu
10:40 AM – 11:00 AM	Break
11:00 AM – 1:00 PM	Challenge Participant systems 2 (GN) – chaired by Zhiyong Lu Team 78 University of Iowa by Sanmitra Bhattacharya Team 83 NCKU IKMLab by Chih-Hsuan Wei Team 89 University of Wisconsin-Milwaukee by Shashank Agarwal Team 93 University of Tokyo by Naoaki Okazaki Team 97 Georgetown University by Hongfang Liu Team 98 Tsinghua University by Minlie Huang Team 101 Yuan Ze University by Richard Tzong-Han Tsai
1:00 PM – 2:30 PM	Lunch (provided, hotel Restaurant)
2:30 PM – 3:30 PM	Keynote: “Communicating Scientific Knowledge: Is progress lagging?” David Lipman , Director of NCBI, NIH, USA
3:30 AM – 3:45 PM	Break
3:45 PM – 5:45 PM	Introduction by workshop sponsor Peter McCartney: Program Director, Division of Biological Infrastructure, NSF Panel Discussion: Challenges and Opportunities for Text Mining in Biology: Government/Publisher Perspectives <ul style="list-style-type: none"> • Government panel (Moderator: Cathy Wu) Valerie Florance: Director, Extramural Programs, NLM, NIH Karin Remington: Director, Center for Bioinformatics & Computational Biology, NIGMS, NIH Susan Gregurick: Program Manager, Computational Biology, DOE Sylvia Spengler: Program Director, Division of Information and Intelligent Systems, NSF Dietrich Rebholz-Schuhmann, Research Group Leader, Literature Analysis, European Bioinformatics Institute (EBI), UK [ELIXIR] • Publisher panel (Moderator: Lynette Hirschman) Anita de Waard: Disruptive Technologies Director, Elsevier Labs Virginia Benson Chanda, Editorial Manager, John Wiley & Sons Kevin Cohen, Biomedical Text Mining Group Lead, Center for Computational Pharmacology, U. Colorado School of Medicine
5:45 PM – 6:00 PM	Break
6:00 PM – 7:30 PM	Reception and Poster Session

September 14, 2010 (Tuesday)	
7:30 AM – 12:00 PM	Registration
7:30 AM – 8:30 AM	Breakfast (provided, outside of Ballroom A/B in the EMC Foyer)
8:30 AM – 9:00 AM	Discussion of systems results (PPI) – Martin Krallinger, CNIO
9:00 AM – 10:30 AM	Challenge Participant systems 3 (PPI) – chaired by Florian Leitner Team 65 University of Zurich by Fabio Rinaldi Team 69 Arizona State University by Robert Leaman Team 70 Universidade de Aveiro, IEETA by Sérgio Matos Team 73 NCBI/NIH by Sun Kim
10:30 AM – 11:00 AM	Break
11:00 AM – 12:30 PM	Challenge Participant systems 4 (PPI) – chaired by Martin Krallinger Team 81 University of Delaware and Indiana University by Hagit Shatkay and Luis Rocha Team 89 University of Wisconsin-Milwaukee by Shashank Agarwal Team 90 National Centre for Text Mining by Rafal Rak Team 100 NCBI/NLM/NIH by Rezarta Islamaj Dogan
12:30 PM – 1:00 PM	Challenge Discussion Panel
1:00 PM – 2:30 PM	Lunch (provided, hotel Restaurant)
2:30 PM – 3:30 PM	Keynote: “Text Bound Annotation and Evaluation -- The Perspectives of BioNLP Shared Tasks and GENIA” Jun’ichi Tsujii , Professor, Department of Computer Science, University of Tokyo, Japan. School of Computer Science, University of Manchester, UK
3:30 AM – 3:45 PM	Break
3:45 PM – 5:45 PM	Developing Text Mining Systems for Biocuration and Biological Discovery: Developer Perspectives •System Interoperability Gully Burns, Information Science Institute, University of Southern California Florian Leitner, Structural and Computational Biology Group, Spanish National Cancer Research Centre, Spain Andreas Prlic, Protein Data Bank, University of California, San Diego Dietrich Rebholz, European Bioinformatics Institute Karin Verspoor, Center for Computational Pharmacology and the Computational Bioscience Program University of Colorado
5:45 PM – 6:00 PM	Break
6:00 PM – 7:30 PM	Systems Demo Session

September 15, 2010 (Wednesday)	
7:30 AM – 8:30 AM	Breakfast (provided, outside of Ballroom A/B in the EMC Foyer)
8:30 AM – 8:45 AM	Interactive task overview (IAT) – Cecilia Arighi, CBCB, UD
8:45 AM – 10:20 AM	Interactive task session 1 (IAT) – Chaired by Cecilia Arighi Team 61 “MyMiner”, David Salgado, Australian Regenerative Medicine Institute - Monash University Team 65 “ODIN: The OntoGene Document Inspector”, Fabio Rinaldi, University of Zurich Team 68 “Gene View”, Philippe Thomas, Humboldt Universität zu Berlin Team 78 “Online Gene Indexing and Retrieval for BioCreative III at the University of Iowa”, Sanmitra Bhattacharya, University of Iowa Team 89 “Interactive NLP system for normalized gene annotations”, Shakshank Agarwal, University of Wisconsin-Milwaukee Team 93 “GNSuite”, Rune Sætre, University of Tokyo
10:20 AM – 11:00 AM	Interactive task session 2 (IAT) •Report from User Advisory Group – Andrew Chatr-aryamontri, BioGrid, Wellcome Trust Centre for Cell Biology, University of Edinburgh, UK
11:00 AM – 11:30 AM	Break
11:30 AM – 12:30 PM	Developing Text Mining Systems for Biocuration and Biological Discovery: User Perspectives •Discussion session – Led by Phoebe Roberts, Pfizer, USA -Metrics for evaluation User Studies and System Evaluation in Information Retrieval, Ben Carterette, Department of Information and Computer Sciences, University of Delaware
12:30 PM – 1:30 PM	Lunch (provided, hotel Restaurant)
1:30 PM – 2:30 PM	Biocreative III and BioCreative IV (Lynette Hirschman and Alfonso Valencia) •Special issue •Future task challenges
2:30 PM	Workshop closing