Wong Lim Soon

From: Wong Lim Soon [limsoon@nus.edu.sg]
Sent: Sunday, 7 August, 2011 2:07 PM
To: David Hsu Xu Ye (SoC); Wynne Hsu (SoC); leeml@comp.nus.edu.sg; leonghw@comp.nus.edu.sg; Ooi Beng Chin (SoC); ksung@comp.nus.edu.sg; Tan Kian Lee (SoC); thiagu@comp.nus.edu.sg; atung@comp.nus.edu.sg; liugm@comp.nus.edu.sg; lijy@comp.nus.edu.sg; liubing@comp.nus.edu.sg; hugowill@comp.nus.edu.sg; chongket@comp.nus.edu.sg; tanhuiyi@comp.nus.edu.sg; melvin@comp.nus.edu.sg; yenan@comp.nus.edu.sg; srigsri@comp.nus.edu.sg; a0066309@nus.edu.sg; Nguyen Nam Ninh; Nguyen Phi Vu; suchee@comp.nus.edu.sg; kangism@gmail.com; ben.gyori@gmail.com; xxqqxiao@gmail.com; Sucheendra Kumar Palaniappan; th.chiang@gmail.com; dcsgv@nus.edu.sg; Won Kok Sung; Bai Haoyu; kevinl@comp.nus.edu.sg; zhouhufeng@gmail.com; iamwym@gmail.com; zhanghao@comp.nus.edu.sg; 89.andre@gmail.com; misias@mimuw.edu.pl; kohchuanhock@gmail.com; dongdife@comp.nus.edu.sg; g0600208@nus.edu.sg; bowenli37@hotmail.com; Chua Gek Huey; Fan Mengyuan; fayue1015@gmail.com; Yong Chern Han; jin-jing@comp.nus.edu.sg; yusufif@comp.nus.edu.sg; hoang@comp.nus.edu.sg; g0305719@nus.edu.sg; hugowill@comp.nus.edu.sg; drcyber@gmail.com; zhizhuo@comp.nus.edu.sg; Lim Jing Quan; rwenang@comp.nus.edu.sg; Wang Jianxing; kangism@gmail.com
Cc: wongls@comp.nus.edu.sg
Subject: CBL update - July 2011

Dear CBL Members,

1/
Congratulations to Chong Ket Fah. He successfully defended his PhD thesis on "New Models and Algorithms for De Novo Peptide Sequencing of Multicharge MS/MS Spectra" last week. His thesis was supervised by A/P Leong Hon Wai.

2/
The following four papers were published this month:

- S. Guillemot, J. Jansson, W. K. Sung. **Computing a Smallest Multi-labeled Phylogenetic Tree from Rooted Triplets**. *IEEE/ACM Trans Comput Biol Bioinform*, 8(4):1141--1147, July 2011.

- Z. Zhang, C. W. Chang, W. L. Goh, W. K. Sung, E. Cheung. **CENTDIST: Discovery of Co-Associated Factors by Motif Distribution**. *Nucleic Acids Research*, 39(Suppl 2):W391--W399, July 2011.

- Zhenhua Li, Limsoon Wong, Jinyan Li. **DBAC: A simple prediction method for protein binding hot spots based on burial levels and deeply buried atomic contacts**. *BMC Systems Biology*, 5(Suppl 1):S5, June 2011.

- Michal Wozniak, Limsoon Wong, Jerzy Tiuryn. **CAMBer: An appproach to support comparative analysis of multiple bacterial strains**. *BMC Genomics*, 12(Suppl. 2):S6, July 2011.

3/
And we hosted the following two seminars:

- [CS Seminar: Investigating Small RNA Guided Regulations with a Combined Approach of High-throughput Sequencing and Bioinformatics](#)
  Yun Zheng, Fudan University
  3 August 2011 at Video Conference Room (COM1-02-13)

- [CS Seminar: Coupled Objects and Behaviors Analysis](#)
  Longbing Cao, University of Technology Sydney
  28 July 2011 at Video Conference Room (COM1-02-13)

4/
Lastly, here is an ad from a friend:

[Bioinformatics postdoc and RA positions, NUS Dept of Statistics. Received from Yik Ying Teo, 29 June 2011.](http://www.comp.nus.edu.sg/~wongls/bp/openings/yyteo-jun2011.pdf)
http://www.comp.nus.edu.sg/~wongls/bp/openings/yyteo-jun2011.pdf

Limsoon

PS. I may have missed out a thing or two…. Please drop me a note.