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- Sent: Thursday, November 30, 2006 4:34 PM

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Subject: RE: Activities of CBL members in October

Hi CBL Folks,

Here are some interesting updates for November ...

1/ Three of the bioinformatics courses supported by us have new names now, which make their content/emphasis more explicit. They are

- CS3225 --> Combinatorial Methods in Bioinformatics (previously: Post-Genome Informatics)
- CS4220 --> Knowledge Discovery Methods in Bioinformatics (previously: Computational Analysis of Biological Data)
- CS5238 --> Advanced Combinatorial Methods in Bioinformatics (previously: Combinatorial Methods in Bioinformatics)

2/ Congrats also to Vipin, Ken, & Hugo for publishing the papers below. Looks like we have papers in bioinformatics conferences/journals every month! I hope we also get citations to our papers every month.

- Vipin Narang, Wing-Kin Sung, Ankush Mittal. "LocalMotif an in silico tool for detecting localized motifs in regulatory sequences," 18th IEEE International Conference on Tools with Artificial Intelligence (ICTAI 2006), pages ???--???, Washington D.C., Nov 2006.
- Tie-Fei Liu and Wing-Kin Sung, "Learning Gene Network Using Conditional Dependence," 18th IEEE International Conference on Tools with Artificial Intelligence (ICTAI 2006), pages ???--???, Washington D.C., Nov 2006.
- Soon-Heng Tan, Willy Hugo, Wing-Kin Sung, See-Kiong Ng. "A correlated motif approach for finding short linear motifs from protein interaction networks", *BMC Bioinformatics*, 7:502, November 2006.

3/ The following talks were given by us and our friends this month:

• Limsoon Wong. "Guts of Dragon Promoter Finder and Mapper". Invited talk at 5th East Asian Biophysics Symposium & 44th Annual Meeting of the Biophysical Society of Japan, Okinawa, Japan, 14 November 2006.

- Vipin Narang, LocalMotif An In-Silico Tool for Detecting Localized Motifs in Regulatory Sequences, SOC Postgraduate Seminar, 30 November 2006.
- Liu Tiefei, Learning Gene Network Using Conditional Dependence, SOC CS Seminar, 23 November 2006.
- Eric Xing, Reasoning in Open Possible World: On a New Class of Nonparametric Bayesian Models for Haplotype Phasing, LD Modeling, and Demographic Inference in Open Ancestral Space, SOC CS Seminar, 16 November 2006.
- Wing-Kin Sung, Annotating the Genome Using Paired-End-diTag, SOC CS Seminar, 9 November 2006.
- Quaid Morris, GeneMANIA: Real-Time Integration of Functional Genomic Datasets for Gene Function Prediction, SOC CS Seminar, 8 November 2006.

If I have left out anything, please do let me know!

4/ No seminar is arranged for December. I hope you all have a nice Xmas.

Limsoon