



# CSB2004 REGISTRATION FORM

## ATTENDEE INFORMATION (PLEASE PRINT CLEARLY)

First Name \_\_\_\_\_ Middle Name \_\_\_\_\_ Last / Family Name \_\_\_\_\_

Company / School Affiliation \_\_\_\_\_

Position \_\_\_\_\_

Email \_\_\_\_\_

Telephone \_\_\_\_\_ Fax \_\_\_\_\_

Address \_\_\_\_\_

City \_\_\_\_\_ State / Region \_\_\_\_\_ Zip / Postal Code \_\_\_\_\_

Country \_\_\_\_\_

## CONFERENCE REGISTRATION (TUES, 17 AUG 2004 TO THURS, 19 AUG 2004) **NOTE: EARLY REGISTRATION IS CLOSED.**

	<del>On or before Monday, 19 July 2004</del>	After Monday, 19 July 2004	Total Due
Member *	\$ 400	\$ 480	US\$ _____
Non-Member	\$ 500	\$ 600	US\$ _____
Student **	\$ 120	\$ 150	US\$ _____

## TUTORIAL REGISTRATION (MONDAY, 16 AUG 2004)

	<del>On or before Monday, 19 July 2004</del>	After Monday, 19 July 2004	Total Due
Member *	\$ 100	\$ 120	US\$ _____
Non-Member	\$ 125	\$ 150	US\$ _____
Student **	\$ 100	\$ 120	US\$ _____
			Total: US\$ _____

If you are registering for tutorial sessions, please indicate your session choices below by    
Please register for at most one morning and one afternoon session. Multiple morning or afternoon sessions not permitted.

### MORNING SESSIONS (8:30AM TO 12:00PM)

- AM 1** Introduction to Evolutionary and Functional Genomic Analysis - Cristian Castillo-Davis, Post-doc, Harvard University
- AM 2** Tandem Mass Spectrometry in Proteomics - Bing Ma, Assist Prof., Univ. of Western Ontario
- AM 3** How to Use Genome Browsers to Get the Most Out of Public Genomes - Daryl Thomas, PhD Candidate UC Santa Cruz
- AM 4** Computational Genetics: Haplotype Inference and Applications in Human Disease Gene Mapping, Tianhua Niu, Harvard Univ.
- AM 5** Introduction to Dynamic Programming and Its Applications to Bioinformatics - Robert Edgar, Visiting Scholar, UC Berkeley

### AFTERNOON SESSIONS (1:30PM TO 5:00PM)

- PM 6** Bioinformatics: The Machine Learning Approach - Pierre Baldi, Professor, UC Irvine
- PM 7** Using dChip for Microarray and SNP Chip Data Analysis - Yu Guo, Ph.D. candidate, Harvard School of Public Health
- PM 8** Discovering Regulatory Networks from Gene Expression and Promoter Sequence - Eran Segal, PHD candidate, Stanford Univ.
- PM 9** Computational Methods in Phylogenetics - Tandy Warnow, Professor, University of Texas, Austin
- PM 10** From Sequence to Structure: Protein Structure Prediction - Juntao Guo, Univ. of Georgia

### PAYMENT METHOD

- Check/Money Order** payable to 'CSB2004' must be mailed to **1579 Fairway Dr, Los Altos, CA 94024** or faxed to: **+1-650-851-8643**
- Credit Card:**  Visa  MasterCard  American Express

Credit Card Number \_\_\_\_\_ Expiration Date \_\_\_\_\_

Print Name as it Appears on Card \_\_\_\_\_

Signature \_\_\_\_\_ Date \_\_\_\_\_

\* Member - IEEE or IEEE-CS member card will be required at check-in to receive the member rate

\*\* Student - A valid student ID card will be required at check-in to receive the student rate