Chuan-Hsiung Chang (張傳雄)

Contact Information

Address: Institute of Biomedical Informatics, National

Yang-Ming University, Taipei 11221, Taiwan

Phone: +886-2-2826-7316 (office)

Fax:+886-2-2820-6754 (office)

Email: cchang@ym.edu.tw Lab web site: http://gel.ym.edu.tw/

Current Position

Assistant Professor and Director of Bioinformatics Program, Center for Systems and Synthetic Biology,

Institute of Bioinformatics, National Yang-Ming University, Taiwan

Taiwan International Graduate Program (TIGP) – Bioinformatics Program,

Academia Sinica, Taiwan

Graduate Education

University of Southern California, Los Angeles, CA, USA. (Ph.D. in Molecular Biology)

Professional Experience

Postdoctoral Fellow, Cancer Research Center, University of California, San Francisco, CA, USA Research Fellow, Genomics Department, Biomedical Engineering Center,

Industrial Technology Research Institute, Taiwan

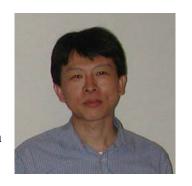
Assistant Professor, Institute of Genetics, National Yang-Ming University, Taiwan

Research Interests

- 1. Forward & reverse engineering of genome complexity through cross-genome comparisons
- 2. Reconstruction and comparison of integrated genome-scale interaction networks
- 3. In silico genome modeling & design for genome engineering and synthetic biology

Selected Publications

- 1. Chih-Hsien Yang, Wei-Chao Liao, Yi-Feng Chang, Li-Hsiang Yen, Te-Ming Lin, Hsin-Yen Wang, Ying-Chun Lo, Ruo-Hsuan Chen, Min-Lun Li, Chun-Ju Yang, Yu-Hsien Lee, Shiang-Jiun Chen, Yi-Hsien Lin, Ueng-Cheng Yang, and Chuan-Hsiung Chang. (2007) A Synthetic Biology Approach for Designing and Engineering BioRobot Systems with Embedded Insulin Auto Feedback and Control Mechanisms. International Conference on Bionano Science.
- 2. Ying-Hsueh Huang, Chien-chi Lo, Wen-Ching Chan, Wei-Xuan Lo, Chih-Hsien Yang, Chi Yang, Ching-Hung Tzeng, Tien-Chueh Kuo and Chuan-Hsiung Chang. (2007) INCO: An Integrated



- Interaction Network of Escherichia coli K-12 MG1655. Microbial Genomes 2007, p.12.
- 3. Chang, C. J., Chiu, J. H., Tseng, L. M., Chang, C. H., Chien, T. M., Wu, C. W. & Lui, W. Y. (2006) Modulation of HER2 expression by ferulic acid on human breast cancer MCF7 cells. European Journal of Clinical Investigation 36 (8), 588-596.
- 4. Chun-Ju Chang, Jen-Hwey Chiu*, Ling-Ming Tseng, Chuan-Hsiung Chang, Tsu-Ming Chien, Chien-Chih Chen, Chew-Wun Wu, Wing-Yiu Lui. Si-Wu-Tang and Its Constituents Promote Mammary Duct Cell Proliferation by Upregulation of HER-2-signaling. (2006) Menopause. 13(6):967-976.
- 5. Lin Yu-Hsuan and Chang Chuan-Hsiung. Phylogenetic Analysis with Metabolic Profiles. (2006) 10th Annual International Conference on Research in Computational Molecular Biology (RECOMB).
- 6. Liao Wei-Chao, Ng Wailap Victor, and Chang Chuan-Hsiung. Distinct Grouping and Genome Organization among *Escherichia coli* Bacteriophages. (2006) 10th Annual International Conference on Research in Computational Molecular Biology (RECOMB).
- 7. Lo, Eric Wei-Xuan and Chang Chuan-Hsiung. Conservation of mitochondrial regulatory elements in sequenced mitochondrial genomes. (2006) 10th Annual International Conference on Research in Computational Molecular Biology (RECOMB).
- 8. Chang, Y.F. Chen, C.Y. Chen, H.W. Lin, I.H. Luo, W.X. Yang, C.H. Lin, Y.H. Chang, C.H. (2005) Bioinformatics analysis for genome design and synthetic biology. Emerging Information Technology Conference. On page(s): 2 pp.
- 9. Chang Y-F and Chang C-H. Identification and Characterization of Conserved Overlapping Genes in Vibrio Genomes. (2005) 9th Annual International Conference on Research in Computational Molecular Biology (RECOMB).
- 10. Lo W-X, Yamada1 T, Tanaka1 M, Hattori M, Goto S, Kanehisa M, and Chang C-H. A method for customized cross-species metabolic pathway comparison. (2004) The 15th International Conference on Genome Informatics.
- 11. Chen H-W, Chang Y-C, Liou D-M, and Chang C-H. Conservation profiling of functionally-coupled gene clusters through multiple genome comparison. (2004) 12th International Conference on Intelligent Systems for Molecular Biology (ISMB).
- 12. Chen C-Y and Chang C-H. CAMP a computational system for Comparative Analysis of Metabolic Pathways. (2004) 8th Annual International Conference on Research in Computational Molecular Biology (RECOMB).
- 13. Chen C-Y, Wu K-M, Liao T-L, Chang Y-C, Chang C-H, Tsai H-C, Liu Y-M, Tsen T-Y, Chen H-J, Li J-C, Su T-L, Su J-S, Chen T-T, Shen Arthur B-T, Shao C-P, Lee C-T, Hor L-I, and Tsai S-F. Comparative genome analysis of *Vibrio vulnificus*, a marine pathogen. (2003) Genome Research 13: 2577-2587.
- 14. Chang Y-C and Chang C-H. Common repeat sequences in bacterial genomes. (2003) Journal of Medical and Biological Engineering 23: 65-72.
- 15. Neve RM, Ylstra B, Chang C-H, Albertson DG, and Benz CC. ErbB2 activation of ESX gene expression. (2002) Oncogene 30: 3934-3938.
- 16. Pan CC, Lin JH, Wu JD, Chang YS, Chang CJ, Tyan MC, Leu CY, Chang YC, Huang JL, and Chang CH. FeverChip for fever-causing pathogen detection and identification. (2001) IBC CHIPS TO HITS

CONFERENCE.

- 17. Scott GK, Chang C-H, Erny K, Xu F, Fredericks WJ, Rauscher FJ, Thor AD, and Benz CC. Ets regulation of the erbB2 promoter. (2000) Oncogene 19: 6490-6502.
- 18. Lin CS, Lau A, Yeh CC, Chang C-H, and Lue TF. Upregulation of L-plastin gene by testosterone in breast and prostate cancer cells: identification of three cooperative androgen receptor-binding sequences. (2000) DNA Cell Biol. 19: 1-7.
- 19. Chao Y, Walther A, Baldwin MA, Chang C-H, Scott GK, Wold MS, and Benz CC. An Epithelium-restricted ETS factor, ESX/ELF3, can assemble both transcription and replication complexes. (2000) Amer. Assoc. Cancer Res. 41:3976.
- 20. Chang C-H, Scott GK, Baldwin MA, and Benz CC. Exon 4-encoded acidic domain in the epithelium-restricted Ets factor, ESX, confers potent transactivating capacity and binds to TATA-binding protein (TBP). (1999) Oncogene 18: 3682-3695.
- 21. Chang C-H, Scott GK, Semyonov J, Chao Y, and Benz CC. Exon4 and 7 encode separte transactivating and chromatin localizing domains in ESX, an Ets transactivator linked to mammary gland development and tumorigenesis. (1999) Amer. Assoc. Cancer Res. 40: 131.