Programme

Day 1 (Oct. 19, 2012)

Morning Session Chair: Luhua LAI			
Time	People	Activities	
9:00-9:10	Chao TANG Howard LIPSHITZ Yuxin YIN	Opening Remarks	
9:10-10:00	Charlie BOONE The Genetic Landscape of a Cell		
10:00-10:20	Coffee Break		
10:20-11:10	Chao TANG	How do yeast cells decide to divide or not to divide	
11:10-12:00	Hao LI	Understanding the Molecular Mechanisms of Aging: Modern Approaches to an Ancient Problem	
12:00-13:30	Lunch		
Afternoon Session Chair: Howard LIPSHITZ			
13:30-14:20	Yuxin YIN Network of P53 and PTEN Tumor Suppressors		
14:20-15:10	Anne-Claude GINGRAS	Cancer Interactomes: Towards a Dynamic View of Signalling	
15:10-15:30	Coffee Break		

15:30-16:20	Qi Ouyang	Nonlinear Dynamics Analysis of P53 Pathway
16:20-17:10	Benjamin BLENCOWE	Systematic Analysis of the Function and Regulation of Alternative Exon Networks
18:30	Dinner	

Day 2 (Oct. 20, 2012)

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Morning Sessio		
Time	People	Activities
9:00-9:50	Brenda ANDREWS	From phenotypes to pathways: global exploration of cellular networks using yeast phenomics
9:50-10:40	Minghua DENG	Integrating Phosphorylation Network with Transcriptional Network Reveals Novel Functional Relationships
10:40-11:10	Coffee Break	
11:10-12:00	Timothy HUGHES	Systematic analysis of the sequence specificity of eukaryotic DNA and RNA binding proteins
12:00-14:30	Lunch & Poster Session	
Afternoon Sess Chair: Qi Ouyar		
14:30-15:20	Qinghua CUI	Understanding the Interactions Between miRNA and Environmental Factors and Their Relationships with Human Disease

15:20-16:10	Zhaolei ZHANG	Evolution and Genetic Variation of microRNA Mediated Gene Regulation in Humans
16:10-16:30	Coffee Break	
16:30-17:20	Yuhai TU	Adaptation: Feedback Control and Its Energy Cost
18:30	Dinner	

Day 3 (Oct. 21, 2012)

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Chair: YuxinYIN

Time	People	Activities
9:00-9:50	Luhua LAI	Human Arachidonic Acid Metabolic Network Analysis and Drug Discovery
9:50-10:40	Quaid MORRIS	RNA-binding Proteins and their Binding Sites
9:50-11:10	Coffee Break	
11:10-12:00	Jianyuan LUO	Regulation of protein acetylation in DNA repair pathway
12:00-14:30	Lunch & Poster Session	

Afternoon Session

Chair: Chao TANG

14:30-15:20	Howard LIPSHITZ	System-wide Analysis of
		Posttranscriptional Regulation during Early Embryogenesis

15:20-16:10	Fuchou TANG	Tracing the pluripotency of early human embryos and embryonic stem cells by single-cell RNA-Seq analysis
16:10-16:30	Coffee Break	
16:30-17:20	Chi-chung HUI	Genetics of systems biology: the yin and yang of Hedgehog signaling".
18:30	Dinner	