# Mathematics, Evolution, and Development Preliminary Program

March 21-27, 2010

#### **Program Outline**

Monday, March 22: Morning	9:00	Opening Ceremony
Monday, March 22. Morning	9:00 9:15	Mechanisms of evolution I
Manday Mansh 22. Afternoon	0.20	Mechanisms of Evolution II
Monday, March 22: Afternoon	13:30	
	15:30	Molecular Evolution I
Tuesday, March 23: Morning	9:00	Signing Ceremonies with U Tübingen and Malahayati U
	9:45	Genome Research
Tuesday, March 23: Afternoon	13:30	Gene Expression, Networks, and their Evolution I
	15:30	Gene Expression, Networks, and their Evolution II
Tuesday, March 23: Evening	20:15	After Dinner Talk
Wednesday, March 24: Morning	9:00	RNA Research: Introduction and Talks
Wednesday, March 24: Afternoon		Free Afternoon
Thursday, March 25: Morning	9:00	Phylogenomics Combinatorics I
Thursday, March 25: Afternoon	13:30	Phylogenomics Combinatorics II
	15:30	Towards Genome Semantics
Thursday, March 25: Evening	18:30	Snacks and Posters
Friday, March 26: Morning	9:00	Molecular Evolution II
Friday, March 26: Afternoon	13:30	Molecular Evolution III
	14:15	Mathematics and Protein Structure
	15:30	Evolution, Genetics and Epigenetics
Saturday, March 27: Morning	9:30	XU Guangqi Lecture
Saturday, March 27: Morning	11:00	Closing Session
Saturday, March 27: Morning	13:00	Farewell

# Monday Morning, March 22, 2010, 9:00-9:15 Opening Ceremony

JIN Li PICB, Shanghai, PR China	9:00-9:05	Welcome
SuzanneBertrand,Johanna PierreMoRST NZ	9:05-9:15	Greetings from NZ

# Monday Morning, March 22, 2010, 9:15-12:00 Mechanisms of evolution I: 3 talks

David Penny The AWC for Mol. Ecol. & Evol., Massey U, Palmerston North, NZ	9:15-9:45 9:45-10:00	Integrating molecular evolution over different time scales Discussion
<b>Burkhard Morgenstern</b> Dept. of Bioinformatics, U Göttingen, Germany	10:00-10:30 10:30-10:45	Phylogenomics sheds light on deep metazoan phylogeny Discussion
	10:45-11:15	Coffee Break
<b>Bai-Lian (Larry) Li</b> Ecol. Complexity Laboratory, UC Riverside, USA	11:15-11:45 11:45-12:00	Self-Sustained Optimality: A Novel Concept of Life's Supply/Demand Scaling Discussion
	12:00-12:05	Group Photo in front of SIBS main building
	12:05-13:30	Lunch Break

#### Monday Early Afternoon, March 22, 2010, 13:30-15:00 Mechanisms of evolution II: 2 more talks

<b>Pete Lockhart</b> The AWC for Mol. Ecol. & Evol., Massey U, Palmerston North, NZ	13:30-14:00 14:00-14:15	Using studies of gene expression to investigate adaptive diversification of New Zealand alpine plants Discussion
Hamish G. Spencer Dept of Zoology, U of Otago, Dunedin, NZ	14:15-14:45 14:45-15:00	The Evolution of Phenotypic Plasticity: Untangling Waiting and Development Times Discussion
	15:00-15:30	Coffee Break

# Monday Late Afternoon, March 22, 2010, 15:30-17:45 Molecular evolution I: 3 talks

Naruya Saitou Saitou Lab, Nat. Inst. of Genetics, Mishima, Japan	15:30-16:00 16:00-16:15	Blurring distinction between ortholog and paralog through frequent duplog homogenization Discussion
Mike Steel Biomath. Research Centre, U of Canterbury, CHC, NZ.	16:15-16:45 16:45-17:00	The combinatorics of gene content and their role in phylogenetics Discussion
Kenta Sumiyama Saitou Lab, Nat. Inst. of Genetics, Mishima, Japan	17:00-17:30 17:30-17:45	Evolutionary study of the vertebrate cis-regulatory elements in development with special reference to the Dlx gene system Discussion

# Tuesday Morning, March 23, 2010, 9:00-9:45 Signing of Cooperation Agreements

XU Zhang, SIBS	9:00-9:05	Signing of the Cooperation Agreement with U of Tübingen, Germany
Shanghai, PR China		Signing of the Cooperation Agreement with Malahayati U, Lampung,
JIN Li, PICB	9:05-9:10	Indonesia
Shanghai, PR China		
Hermann Schlüsener	0.10.00.00	The Shanghai-Tübingen Research School for Computational Methods in
Inst. for Brain Research,	9:10-09:30	the Life Sciences and Medicine: A vision for the future
Med. School, U of Tübingen, Germany	09:30-09:45	Discussion
ruomgen, cermany	09.50-09.45	

# Tuesday Morning, March 23, 2010, 9:45-12:15 Genome Research: 3 talks

<b>FENG Jianfeng</b> U of Warwick, UK, and Comp. Systems Biology,	09:45-10:15	Transcription Dynamics in Nucleus
Fudan U, Shanghai	10:15-10:30	Discussion
	10:30-10:45	Coffee Break
Hon Wai Leong Dept of CS, NUS,	10:45-11:15	Spatial Gene Clusters: Two Models and Applications
Singapore	11:15-11:30	Discussion
Yukuto Sato Saitou Lab, Nat. Inst. of	11:30-12:00	Molecular evolutionary study with pathway simulation on gene dosage effect
Genetics, Mishima, Japan	12:00-12:15	Discussion
	12:15-13:30	Lunch Break

# Tuesday Early Afternoon, March 23, 2010, 13:30-15:00 Gene Expression, Networks, and their Evolution I: 2 talks

Martin Vingron MPI for Mol. Genetics, Berlin-Dahlem, Germany, and PICB, Shanghai	13:30-14:00 14:00-14:15	Transcription factor binding site analysis and histone modifications point at two classes of promoters Discussion
Lesley Collins The AWC for Mol. Ecol. & Evol., Massey U, Palmerston North, NZ	14:15-14:45 14:45-15:00	Investigating ancestral RNA-Protein networks in eukaryotes Discussion
	15:00-15:30	Coffee Break

#### Tuesday Late Afternoon, March 23, 2010, 15:30-17:45 Gene Expression, Networks, and their Evolution II: 3 more talks

Xinguang ZHU PICB, Shanghai, PR China	15:30-16:00 16:00-16:15	Evolution of plant metabolic network revealed through genomic and expression analysis Discussion
YAN Jun PICB, Shanghai, PR China	16:15-16:45 16:45-17:00	Flux analysis of metabolism and inter-organ transport in mammals Discussion
Olgert Denas Dept. of Math. & CS, Emory U, Atlanta, USA	17:00-17:30 17:30-17:45	Adapting the maximum b-matching problem to gene network reconstruction Discussion
	18:00-20:00	Conference Dinner

#### Tuesday Evening, March 23, 2010, 20:15 After-Dinner talk

<b>David Penny</b> The AWC for Mol. Ecol.	20:15-21:00	The continuity of mind, from great apes to humans
& Evol., Massey U, Palmerston North, NZ	21:00	Discussion

#### Wednesday, March 24, 2010, 9:00-13:15 RNA Research: Introduction and 5 talks

Axel Mosig PICB, Shanghai, PR China	9:00-9:15	Introduction
Ming Chen Zhejiang U, Hangzhou, PR China	9:15-9:45 9:45-10:00	Computational analysis of RNA editing and small RNA characterization in plants Discussion
<b>Lihui Lai</b> East China Normal University, Shanghai, PR China	10:00-10:30 10:30-10:45	MicroRNA-145 modulates tumor growth and angiogenesis in breast cancer through translational repression of DDR1, NRAS and VEGF Discussion
	10:45-11:00	Coffee Break
Christian Reidys Center for Comb., Nankai U, Tianjin, PR China	11:00-11:30 11:30-11:45	Robustness and Plasticity in the RNA world Discussion
Ming Zhang Theor. Biology and Biophys., Los Alamos National Laboratory	11:45-12:15 12:15-12:30	Interplay between microRNAs and cis-antisenses Discussion
Lesley Collins The AWC for Mol. Ecol. & Evol., Massey U, Palmerston North, NZ	12:30-13:00 13:00-13:15	Graph theoretical and algorithmic problems related to genome research, with particular emphasis on RNA (secondary) structure and RNA interaction. Discussion
	13:15-14:15	Lunch Break

#### Wednesday Afternoon, March 24, 2010, 14:15

Free Afternoon — Options: Visiting the Shanghai Museum or the Shanghai Botanical Garden or ..., perhaps also the Peking Opera House in Shanghai in the evening presenting the Shaoxing Opera "Fish Pursue"

# Thursday Morning, March 25, 2010, 9:00-12:15 Phylogenetic combinatorics I: 4 talks

David Bryant Math. Dept, U of Auckland, NZ	9:00-9:30 9:30-9:45	The tight span of Boolean functions Discussion
Vincent Moulton School of CS, U of East Anglia, Norwich, UK	9:45-10:15 10:15-10:30	Phylogenetic networks and multi-labeled trees Discussion
	10:30-10:45	Coffee Break
<b>Stefan Gruenewald</b> PICB, Shanghai, PR China	10:45-11:15 11:15-11:30	A Quartet Version of Split Decomposition Discussion
Katharina Huber School of CS, U of East Anglia, Norwich, UK	11:30-12:00 12:00-12:15	Recent results on phylogenetic networks of bounded level Discussion
	12:15-13:30	Lunch Break

# Thursday Early Afternoon, March 25, 2010, 13:30-15:00 Phylogenetic combinatorics II: 2 more talks

Jack Koolen Math. Dept, POSTECH, Pohang, Korea	13:30-14:00 14:00-14:15	On the minimal number of quartets in a maximal 2-compatible split system Discussion
Mike Steel Biomath. Res. Centre, U of Canterbury, CHC, NZ.	14:15-14:45 14:45-15:00	Extracting evolutionary signal from patchy genomic data Discussion
	15:00-15:30	Coffee Break

#### Thursday Late Afternoon, March 25, 2010, 15:30 -17:45 Towards Genome Semantics: 3 talks

Alberto Apostolico College of Comp., Georgia Tech, Atlanta, USA	15:30-16:00 16:00-16:15	The Quest for Succinct Pattern Generators Discussion
Raymond Chan Math. Dept, The Chinese University of Hong Kong	16:15-16:45 16:45-17:00	Maximum Entropy Principle for Composition Vector Method in Phylogenetics Discussion
<b>JIN Li</b> PICB, Shanghai, PR China	17:00-17:30 17:30-17:45	Identifying mutational hot spots of copy number variations in human genomes: A phylogeny approach Discussion

# Thursday Evening, March 25, 2010, 18:30

#### **Snacks and Posters**

### Friday Morning, March 26, 2010, 9:00-12:30 Molecular evolution II: 4 talks

Mick Roberts IIMS & NZIAS, Massey U, Auckland, NZ	9:00-9:30 9:30-9:45	Virus evolution - escaping immunity Discussion
Alex Grossmann Lab Statistique et Genome, Evry, France	9:45-10:15 10:15-10:30	Words, words, words Discussion
	10:30-11:00	Coffee Break
James Degnan Dept. of Math. and Stat., U of Canterbury, CHC,	11:00-11:30	Identifiability of rooted species trees from unrooted gene tree distributions
NZ	11:30-11:45	Discussion
Louxin Zhang Math. Dept., NUS, Singapore	11:45-12:15 12:15-12:30	Reconciling gene trees with the deep coalescence cost Discussion
	12:30-13:30	Lunch Break

#### Friday Early Afternoon, March 26, 2010, 13:30-14:15 Molecular evolution III: 1 talk

Wim Hordijk World Wanderer (currently Comp.	13:30-14:00	Adventures in Phylogenetics
Systems Biology, Fudan U, Shanghai)	14:00-14:15	Discussion

# Friday Early Afternoon, March 26, 2010, 14:15-15:00 Mathematics and Protein Structure: one talk

<b>Deok-Soo Kim</b> Voronoi Diagram Research Center,	14:15-14:45	BioGeometry and Beta-complexes: What, and how useful, are they?
Hanyang U, Seoul, Korea	14:45-15:00	Discussion
	15:00-15:15	Coffee Break

# Friday Late Afternoon, March 26, 2010, 15:15-18:30 Evolution, Genetics, and Epigenetics: 4 talks

Philipp Khaitovich PICB, Shanghai, PR China, khaitovich@eva.mpg.de	15:15-15:45 15:45-16:00	A molecular survey across lifespan: the role of developmental timing in human brain evolution and aging Discussion
<b>FU XinYuan</b> Dept Biochem NUS, Singapore, Dept Microbiol. and Immun., Indiana U, Indianapolis, USA	16:00-16:30 16:30-16:45	Evolution and Functional Analysis of Networks Regulated by the STAT Signaling Pathway Discussion
	16:45-17:00	Coffee Break
Liping Zhao Systems Biomedicine Center, Shanghai Jiao Tong U, Shanghai, PR China	17:00-17:30 17:30-17:45	Whole-Body Systems Biology Discussion
<b>XU Shuhua</b> PICB, Shanghai, PR China	17:45-18:15 18:15-18:30	Chromosome-Wide Haplotype Sharing: Integrating Recombination Information into Studies on Human Population Genetic Relationship Discussion

# Saturday, March 27, 2010, 9:30-11:00 Evolutionary Medicine: A XU Guangqi Lecture

Sir Peter Gluckmann, FRS The Liggins Institute, and	9:30-10:30	Echoes of the past - an evolutionary perspective on medicine and its relevance to public health in Asia
the NZ Prime Minister's Chief Science Advisor	10:30 -11:00	Discussion

#### Saturday, March 27, 2010, 11:00 - 13:15 Closing Session

All Participants from all over this planet	11:00-13:00	A general discussion on how to forge national and international alliances to meeting current challenges in mathematics arising in the study of the molecular evolution of genes and genomes, their control, and their correlation with metabolic, developmental, morphological, and environmental issues.
<b>Andreas Dress</b> PICB, Shanghai, PR China	13:00-13:15	Farewell