WTBG@XTBG

Workshop on Tropical Biodiversity and Genomics [at] Xishuangbanna Tropical Botanical Gardens June 4th-July 7th, 2012

June 3rd (Sunday)

Participants arrive from abroad and Kunming

Registration – evening welcome mixer (Green Bamboo Bar), 8.00pm

June 4th(Monday)

0800-0930	Final registration. Coffee or tea at foyer (new research center-East Campus).
0930-1030	Introductions
Director of XTBG	Welcome Remark
Organization Committee : Kua Chai-Shian	Basic logistics and workshop information
Scientific Committee : Chuck Cannon	Objectives and Agenda, self introduction of participants
1030-1100	coffee-tea break
1100-1200 Chuck Cannon	Diversification of Tropical Trees
1200-1400	Lunch/Siesta/Working Groups
Chair: Song Yu	Evolutionary genomics
1400-1450 Wu Chung-I	The Ultra-microevolutionary process - Natural selection within multicellular organisms
1450-1540 Graham Stone	The Gallwasp Multitrophic System - a very convenient system for a range of cool questions!
1540-1600	coffee-tea break
Chair: Fan Huan	Bioinformatics
1600-1620 Jack Hearn	Dissecting an extended phenotype: gall induction by cynipid gallwasps
1620-1640 Ye Chengxi	Exploiting Sparseness in de novo Genome Assembly
1640-1700 He Ziwen	Estimating Level of DNA Polymorphism from Next Generation Sequencing Data with High Error Rate
1620-1640 Zhang Ticao	Genome sequence of Anabaena ssp.: insights into adaptation to the Qinghai-Tibet Plateau environment
1720-1800 Gao Jie	Demography and speciation history of the homoploid hybrid pine Pinus densata on the Tibetan Plateau.
1800-2100	Welcome Dinner Party at Rainforest Lodge

June 5th (Tuesday)

Chair : Chen Si Chong	Comparative Genomics
930-1000 Kua Chai-Shian	Comparative Genomics Without Reference genomes
Chair : Marie Fougère	Transcritptomics
1000-1030 Yann Surget-Groba	Analysis of RNA-seq data
1030-1100	coffee-tea break
1100-1130 Chow Keng See	A Decade of Rubber Genomics
1130-1200 Yang Ming	Inferring the invasion history of <i>Mikania micrantha</i> by RNA-seq
1200-1400	Lunch/Siesta/Working Groups
Chair: Warin Wannaprapho	Population Genomics
1400-1500 Shi Shuhua	Fine-scale local adaptation in mangroves revealed by population genomic analysis
1500-1540 Yann Surget-Groba	Rad sequencing as a rapid biodiversity assessment tool
1540-1600	coffee/tea break
1600-1700 Graham Stone	XTBG Seminar
1800-1900	Dinner
1900-2100 John Harting/Chuck Cannon/	reference free pipeline via Skype / working groups

June 6th (Wednesday)

Field Trip 730	Rubber plantation/ Forest
1200	Lunch
evening	Working groups

June 7th (Thursday)

930-1200	
Participants	Reports
Chuck Cannon	Final synthesis and discussion for the workshop
	Farewell lunch

June 8th (Friday)

additional working day for those who want to work on data analysis and paper development.

June 9th (Saturday)

additional working day for those who want to work on data analysis and paper development.