



China-South Africa Mini Symposium on Plant Omics

June 4th, 2015, Hangzhou, China

Venue: Meeting room 234, Main building of College of Life Sciences, Zhejiang University, Zijingang Campus

Time	Speaker	Topic
9:00-9:15	Ming Chen, Nicola Illing	Openning
9:15-9:45	Stephen Schlebusch (University of Cape Town)	Xerophyta genome
9:45-10:15	Rafe Lyall (University of Cape Town)	Transcriptome assemblies
10:15-10:45	Yuriy Orlov (Institute of Cytology and Genetics, Russia)	Computer analysis of gene location and expression patterns based on ChIA-PET and Hi-C data
10:45-11:15	Chunhui Yuan (Zhejiang University)	PceRBase: a comprehensive competing endogenous RNA database in plants
11:30-14:00	Lunch break	
14:00-14:30	Robert Ingle (University of Cape Town)	Clock regulation of innate immunity in Arabidopsis
14:30-15:00	Evan Milborrow (University of Cape Town)	Strategy to characterize long non-coding RNA
15:00-15:30	Jingjing Wang (Zhejiang University)	Genome-wide analysis of the distinct types of chromatin interactions in arabidopsis thaliana
15:30-16:00	Youjun Huang (Zhejiang A&F University)	Use of transcriptome sequencing to understand the pistillate flowering and oil accumulation in hickory
16:00-16:30	Ming Chen (Zhejiang University)	Organelle-focused proteomes and interactomes in rice
16:30-17:00	Discussion	

Organizers:

Ming Chen, Zhejiang University; Bioinformatics Society of Zhejiang Province

Nicola Illing, University of Cape Town