

Program of RAP6 Workshop

International Scientific Organizing Committee

- Takashi Gojobori, National Institute of Genetics, Japan
- Takuji Sasaki, National Institute for Agrobiological Sciences, Japan
- Rod Wing, Arizona Genomics Institute, USA
- Olivier Panaud, University of Perpignan, France
- Tatiana Tatusova, National Center for Biotechnology Information, USA
- Pankaj Jaiswal, Oregon State University, USA
- Richard Bruskiewich, International Rice Research Institute, Philippines

Sunday, 15th November 2009

07:30 – 08:30	Registration package pickup by delegates	
08:30 – 08:55	RAP6 Opening Addresses	Chair: Kenneth McNally
	· Welcome from IRRI (Hei Leung)	
	· Welcome by Takuji Sasaki	
	<i>Plenary Session I – Rice Genome Assembly and Annotation</i>	
08:55 – 09:00	Introduction of 1 st Plenary Session	Chair: Takuji Sasaki
09:00 – 09:10	Takashi Matsumoto, NIAS, Japan	Towards a Consensus Nipponbare Rice Genome Assembly
09:10 – 09:30	Tsuyoshi Tanaka, NIAS, Japan	Improvements to the RAPdb Rice Genome Annotation

09:30 – 9:50	Kevin Childs, Michigan State University, USA	Improvements to the MSU Rice Genome Annotation
9:50 – 10:10	Pankaj Jaiswal	Gramene Database: A Resource for Comparative Plant Genomics
10:10 – 10:25	<i>Coffee Break</i>	
	<i>Plenary Session II – Comparative Genomics</i>	
10:25 – 10:30	Introduction of 2 nd Plenary Session	Chair: Olivier Panaud
10:30 – 11:00	Jo Messing, Rutgers University, USA	“The prolamins of rice and their orthologs in other cereals”
11:00 – 11:30	Thomas Bureau, McGill University, Canada	“Bridging comparative, population and functional genomics to identify and experimentally validate novel regulatory regions and genes for crop improvement”
11:30 – 12:00	Klaus Mayer, MIPS, Germany	“Tailoring comparative plant genomics: sCISsors and Zippers”
12:00 – 13:25	<i>Buffet Lunch</i>	
	<i>Plenary Session III – Post-Sequencing Era</i>	
13:25 – 13:30	Introduction of 3 rd Plenary Session	Chair: Takuji Sasaki
13:30 – 15:00	IRGSP Alumni: Takuji Sasaki, Japan	Invited brief presentations by IRGSP alumni on post-sequencing

	<ul style="list-style-type: none"> · Takashi Matsumoto, Japan · Rod Wing, USA · Yue-Ie Hsing, Taiwan · Antonio de Oliveira, Brazil · Apichart Vanavichit, Thailand · Nagendra Singh, India 	<p>era genomic activities in their respective countries</p>
15:00 – 15:15	<i>Coffee Break</i>	
	<i>Plenary Session IV – Functional Annotation of the Rice Genome</i>	
15:15 – 15:20	Introduction of 4 th Plenary Session	Chair: Ken McNally
15:20 – 15:50	Shoshi Kikuchi, NIAS	<p>“Comparative analysis of transcription factor genes from the 32 organisms revealed the distinct evolution procedures of TF genes in plants and animals”</p>
15:50 – 16:10	Shamala Tirimanne, Sri Lanka	<p>“Use of nucleic acid databases and protein structure databases to characterize protein families: the 10-member rice hexokinase family”</p>
16:10 – 16:30	Hei Leung, Ken McNally, Ramil Mauleon, IRRI	<p>“Needs for high quality rice gene annotation for large-scale SNP discovery projects”</p>

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- 16:30 – 17:30 - *Back to the Future (of RAP)* - Chair: Rod Wing -
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