

# Update on high throughput marker facility and genotyping services at IRRI

**Dr. Michael J. Thomson**

Molecular Marker Applications Lab (MMAL)

Plant Breeding, Genetics and Biotechnology Division, IRRI

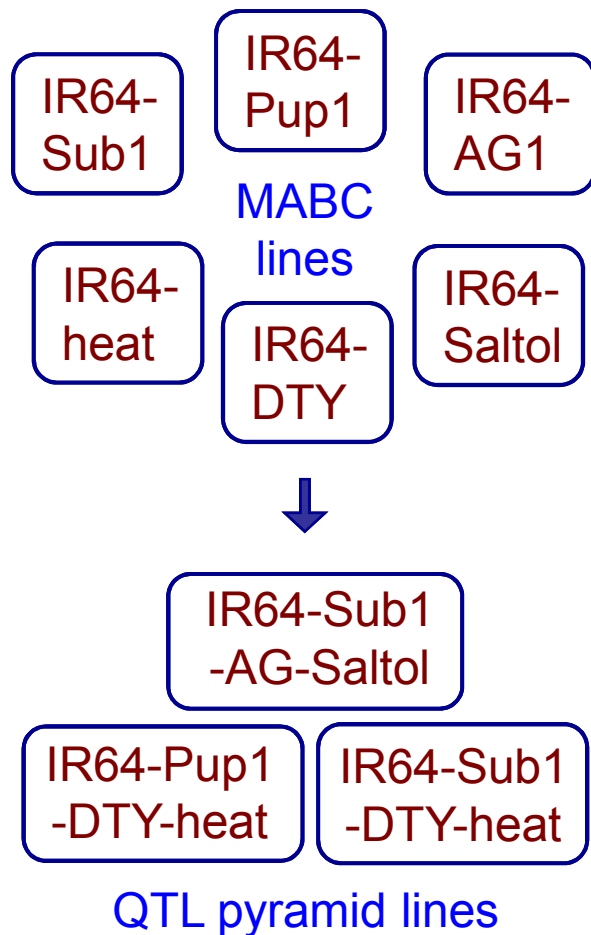
2013 HRDC Annual Meeting, IRRI      2 April 2013

**IRRI**

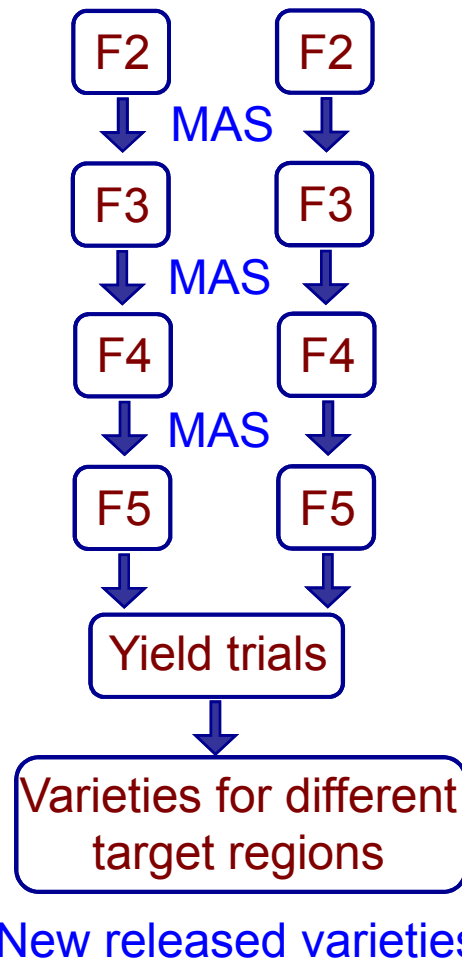


# Integrating markers into breeding programs

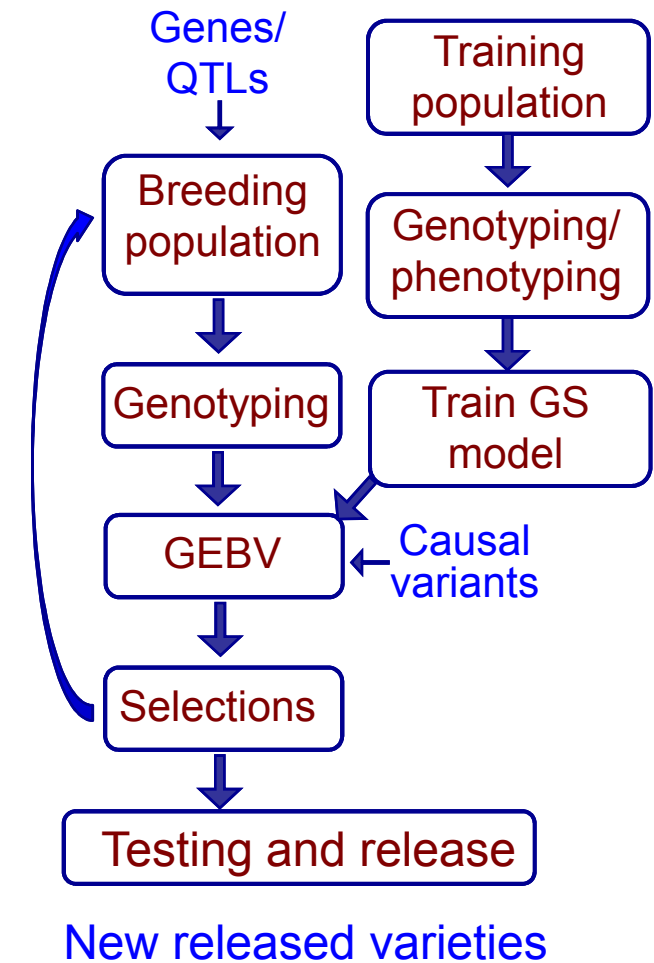
## MABC and QTL pyramiding



## MAS in the pedigree breeding programs

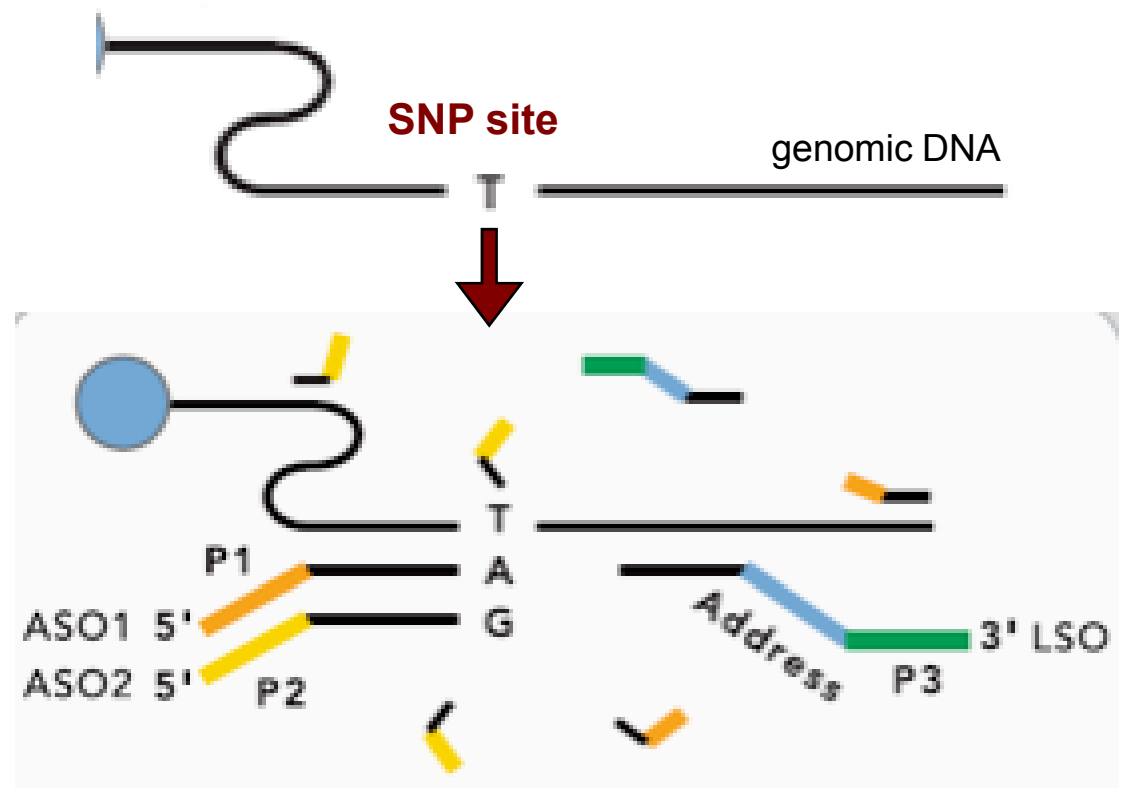


## Genomic selection



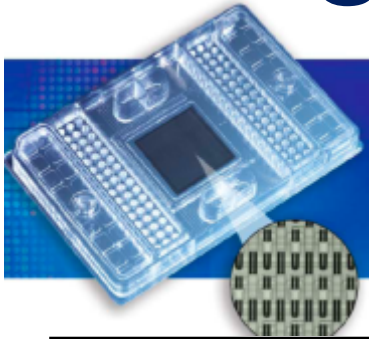
# Advantages of SNP markers

- Millions of SNP loci across the genome
- Most SNP markers are bi-allelic
- SNP data can be easily merged in a database
- Rapid high-throughput SNP genotyping systems are available
- SNP haplotypes can track specific alleles



Single nucleotide polymorphism (SNP) marker

# SNP marker platforms



**Fludigm** Fine-mapping, trait-based  
24 and 96 SNPs SNPs for breeding, MABC



**BeadXpress** Diversity analysis,  
384 SNPs fingerprinting, MABC,  
QTL mapping



**GBS** High resolution  
3k-20k SNPs mapping,  
genomic selection,  
association studies



**SNP chips**  
6k-50k SNPs



# New initiative under the Global Rice Science Partnership (GRiSP)

- **Product 2.1.3 High-throughput SNP genotyping platform for breeding applications:**
  - **Facilities for high-throughput SNP genotyping** set up at IRRI and CIAT with access to GRiSP partners
  - **SNP fingerprinting platform** in place to support activities for variety identification, quality assurance and characterizing breeding lines and released varieties
  - **Trait-based SNP markers** for key traits validated, optimized, and deployed in breeding programs

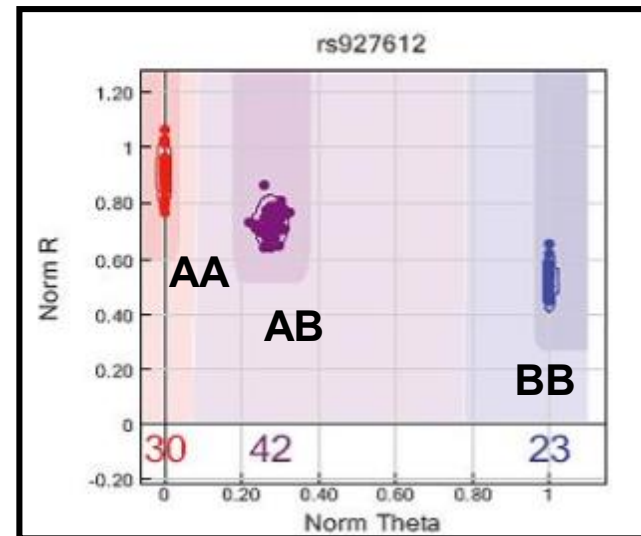
# 384-SNP marker genotyping at IRRI



- Illumina BeadXpress Reader at IRRI
- 96 samples x 384 SNP markers per run

## Breeding applications:

- Diversity analysis
- DNA fingerprinting
- QTL mapping
- Marker-assisted selection



Over 16,000 rice samples  
(6 million data points) run  
at IRRI in past 3 years

Thomson et al. 2012 (Mol. Breed. 29:875)

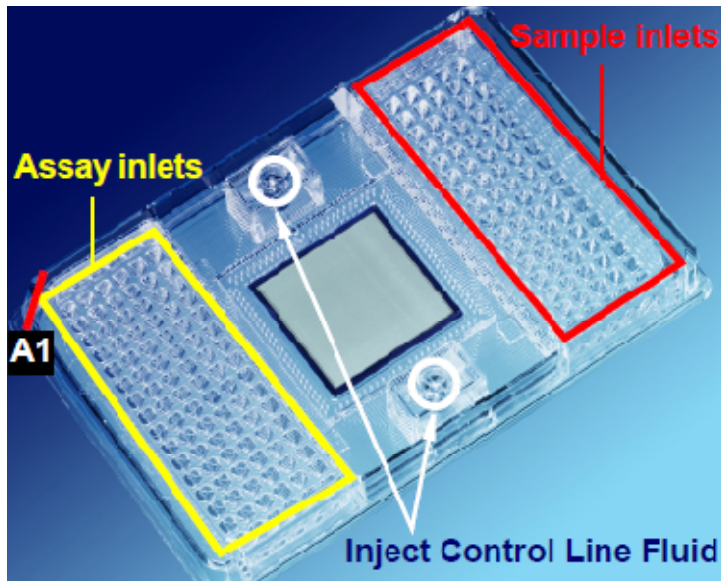
# Seven custom VeraCode OPA sets

Rice SNP set	Illumina OPA ID	No. of SNPs	Description
RiceOPA1.0	VC0011438-OPA	96	Quality control
RiceOPA2.1	GS0011861-OPA <sup>a</sup>	384	indica/indica
RiceOPA3.1	GS0011862-OPA <sup>b</sup>	384	indica/japonica
RiceOPA4.0	VC0013043-OPA	384	japonica/japonica
RiceOPA5.0	GS0011972-OPA	384	indica/ <i>O. rufipogon</i>
RiceOPA6.0	VC0011530-OPA	384	japonica/ <i>O. rufipogon</i>
RiceOPA7.0	VC0011440-OPA	384	indica/japonica

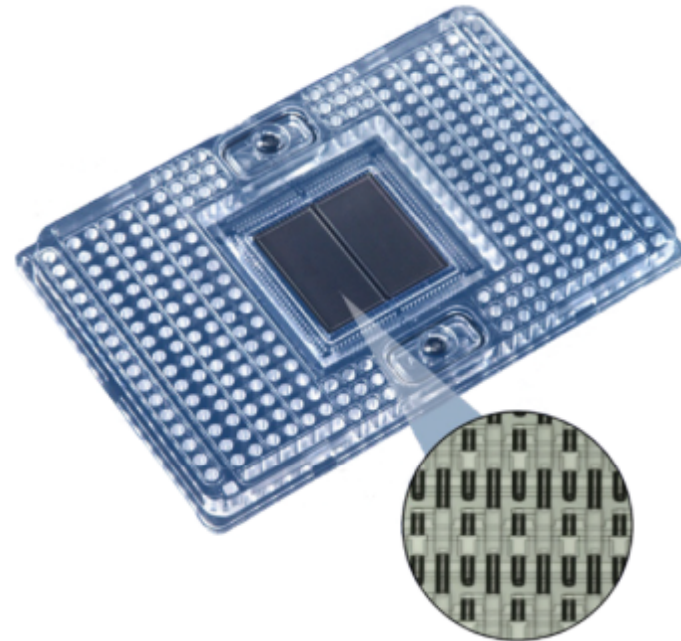
Thomson et al. 2012 (Molecular Breeding 29:875–886)

# Fluidigm EP1 system for high sample throughput applications

- **Fluidigm Dynamic Arrays** provide nano-liter size reactions for rapid genotyping at a low cost per sample



96 SNPs x 96 samples  
(9,216 reactions)



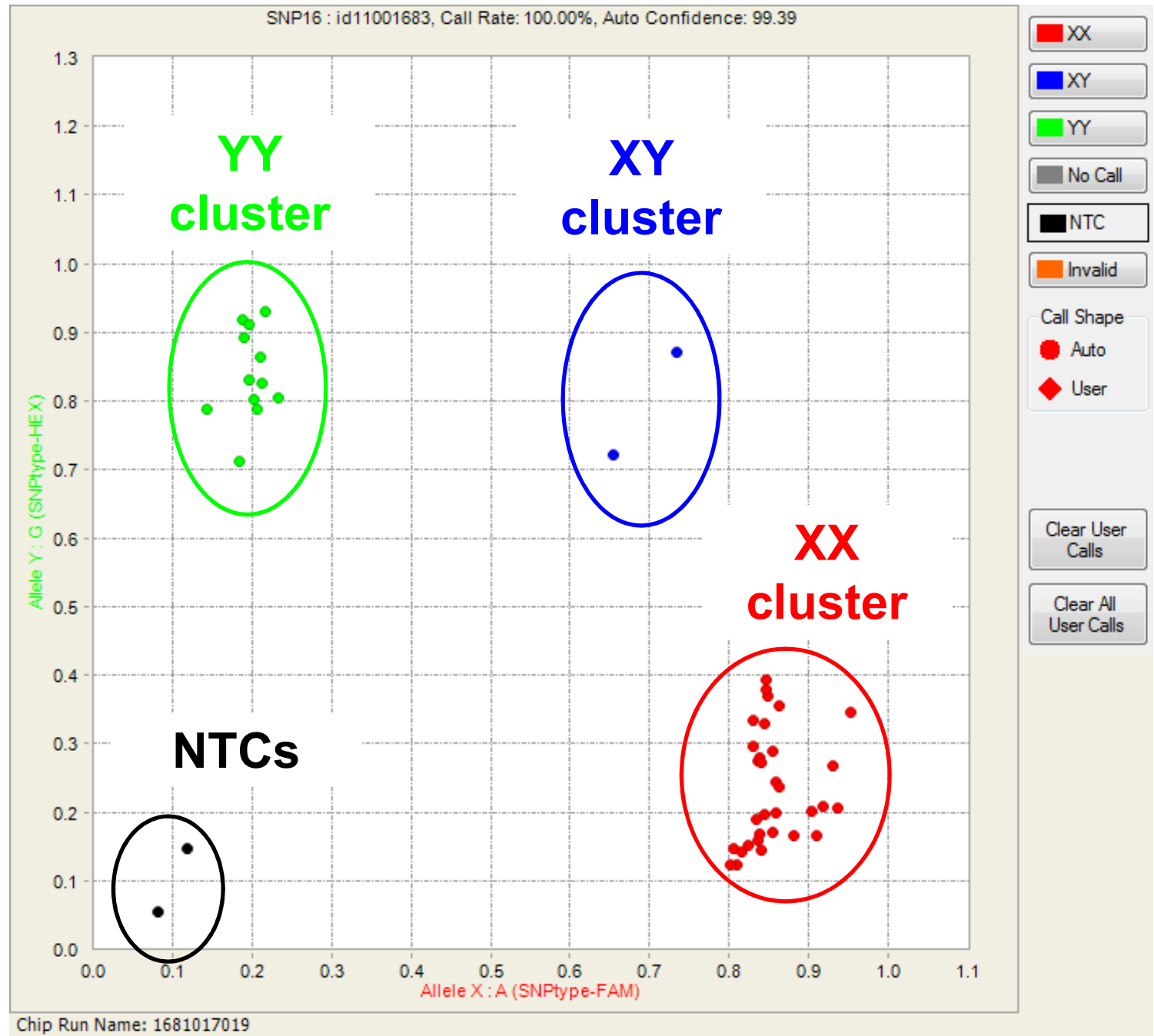
24 SNPs x 192 samples  
(4,608 reactions)

- Different sets of custom SNPs can be used for each chip run

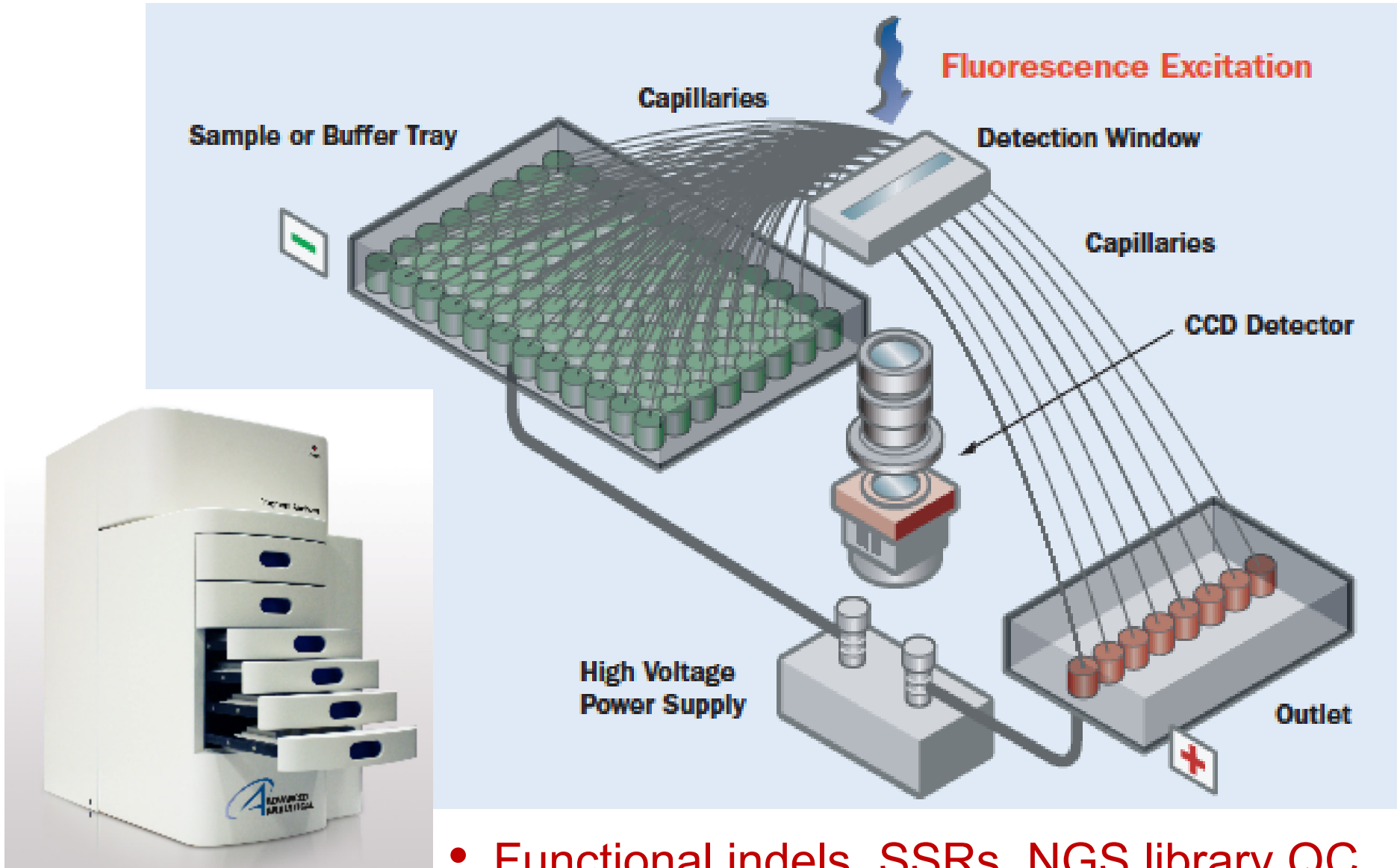
SNP genotyping services now offered at IRRI starting Mar. 2013

**Scatter-plot view for a single SNP**

**Well defined clusters**

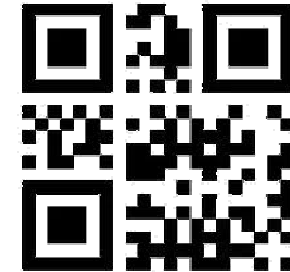
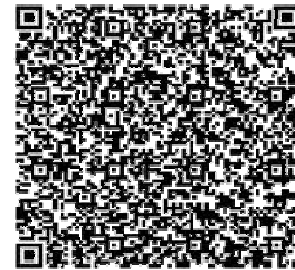


# 96-capillary Fragment Analyzer

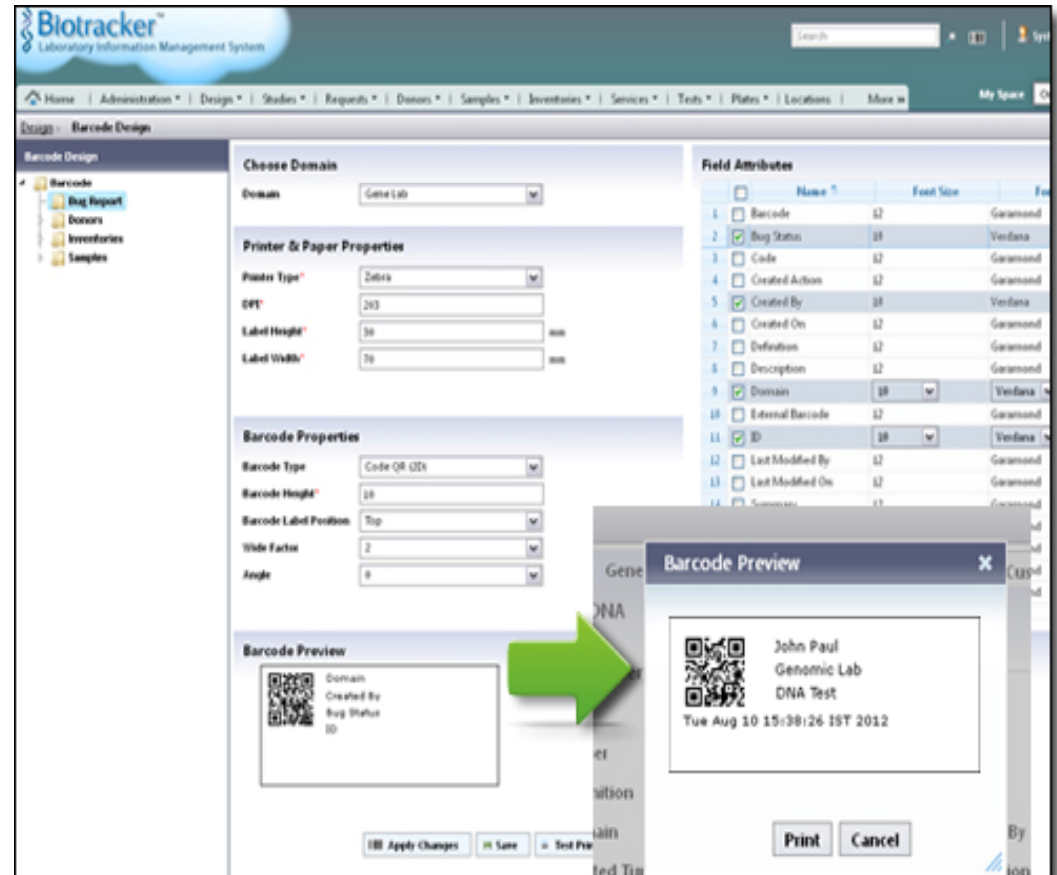


- Functional indels, SSRs, NGS library QC

# LIMS and barcoding for lab data management



- Barcoding for plots/individual plants, tissue and DNA samples, SNP assays
- Integrating with web-based Biotracker 5.0 LIMS (Ocimum Biosolutions)





# Genotyping Services Lab at IRRI

## External for-profit customers (private sector)

Service	Costs - per set	Samples per set	Costs - per sample	Costs - per SNP	Fixed or custom	Shared or exclusive
BXP-384 SNPs	\$1,608	24	\$67.00	\$0.17	Fixed	Shared
Fluidigm-96 SNPs	\$2,065	94	\$21.97	\$0.23	Fixed	Shared
Fluidigm-96 SNPs	\$2,362	94	\$25.13	\$0.26	Custom	Shared
Fluidigm-96 SNPs	\$2,488	94	\$26.47	\$0.28	Custom	Exclusive
Fluidigm-24 SNPs	\$1,400	188	\$7.45	\$0.31	Fixed	Shared
Fluidigm-24 SNPs	\$1,502	188	\$7.99	\$0.33	Custom	Shared
Fluidigm-24 SNPs	\$1,574	188	\$8.37	\$0.35	Custom	Exclusive
DNA extraction	\$348	94	\$3.70		Fixed	Shared

Services now offered for DNA extraction (from lyophilized leaf tissue), Fluidigm 24 and 96 SNP sets (fixed and custom), and BeadXpress 384-SNP sets (fixed)



# GSL price list (public sector partners)

## External non-profit (Universities, NARES, other CG)

Service	Costs - per set	Samples per set	Costs - per sample	Costs - per SNP	Fixed or custom	Shared or exclusive
BXP-384 SNPs	\$1,296	24	\$54.00	\$0.14	Fixed	Shared
Fluidigm-96 SNPs	\$1,666	94	\$17.72	\$0.19	Fixed	Shared
Fluidigm-96 SNPs	\$1,906	94	\$20.28	\$0.21	Custom	Shared
Fluidigm-96 SNPs	\$2,009	94	\$21.37	\$0.22	Custom	Exclusive
Fluidigm-24 SNPs	\$1,128	188	\$6.00	\$0.25	Fixed	Shared
Fluidigm-24 SNPs	\$1,213	188	\$6.45	\$0.27	Custom	Shared
Fluidigm-24 SNPs	\$1,270	188	\$6.76	\$0.28	Custom	Exclusive
DNA extraction	\$282	94	\$3.00		Fixed	Shared

# Training and capacity building for molecular breeding in rice



Molecular Breeding Course

- **Molecular Breeding course**

Case studies in using markers to accelerate breeding efforts, including SNP genotyping.

September 2-13, 2013

- **SNP Data Analysis Course**

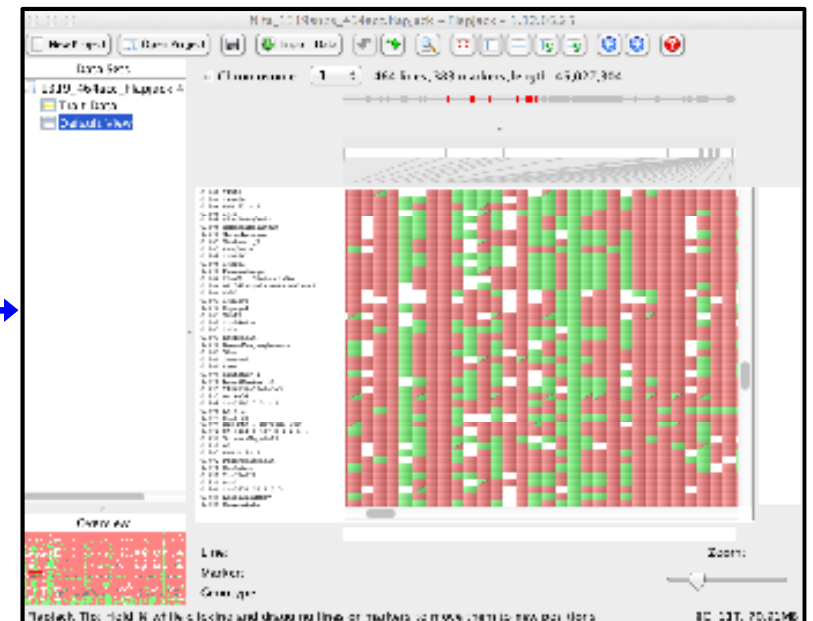
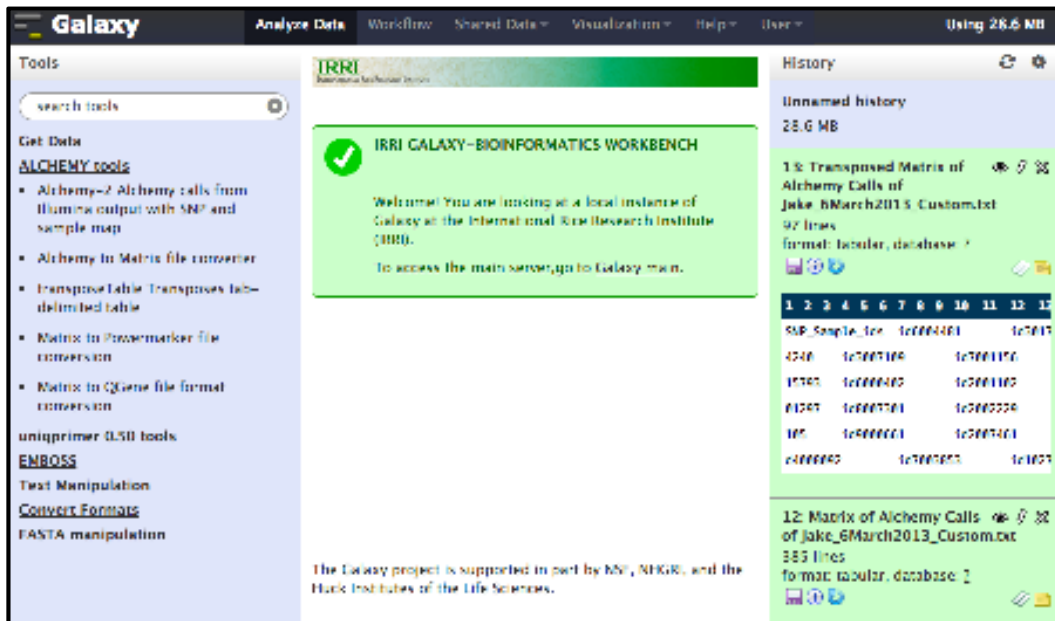
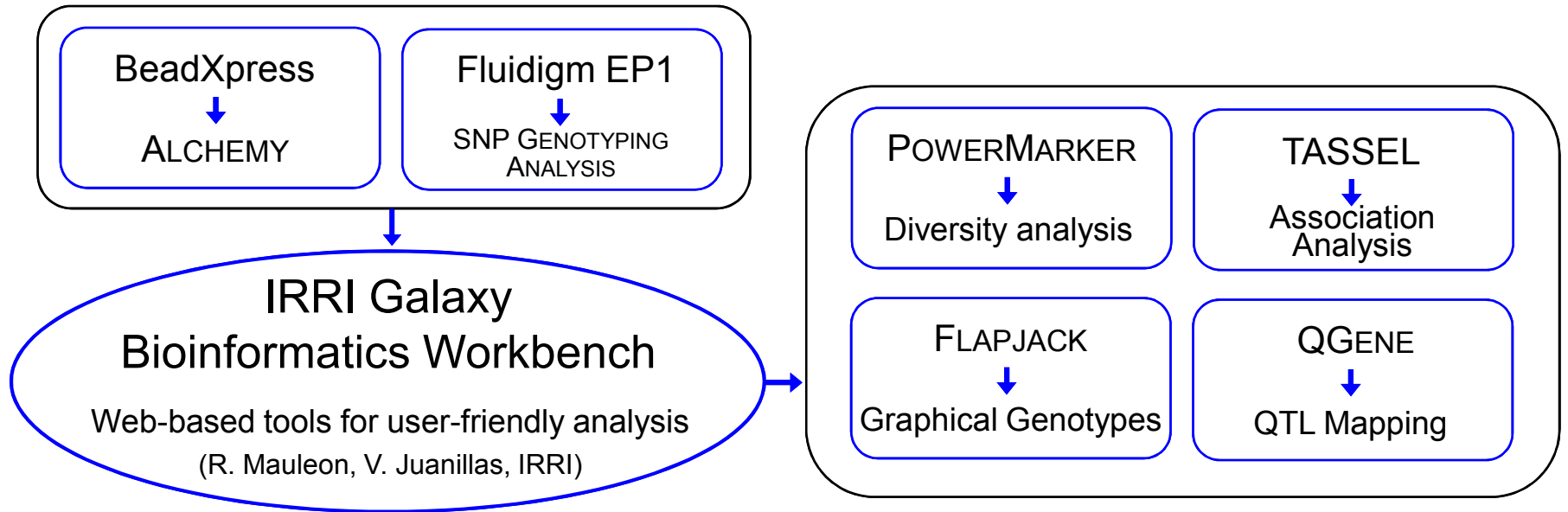
SNPs for genetic diversity analysis, QTL mapping, graphical genotyping, association mapping and MAS.

April 22-27, 2013



SNP Data Analysis

# SNP data analysis workflow



# Acknowledgments

## **MMAL-SNP team at IRRI:**

- Ma. Ymber Reveche
- C. Jade Dilla-Ermita
- Erwin Tandayu
- Dennis Lozada
- Socorro Carandang
- Grace Cariño
- Crisostomo Dizon
- Venice Juanillas

Joong Hyoun Chin

- Chenie Zamora
- Katreena Titong

Bertrand Collard, Ramil Mauleon, Casiana Vera Cruz, Hei Leung, Ken McNally, Ed Redoña, Parminder Virk, Abdelbagi Ismail



## **Cornell University**

Susan McCouch, Chih-Wei Tung, Mark Wright,  
Andy Reynolds, Hyunjung Kim

## **USDA-ARS**

Brian Scheffler (GBRU, Mississippi)  
Georgia Eizenga, Anna McClung (DBNRRRC, Arkansas)

## **Funding sources:**

U.S. NSF, USDA, Government of Japan, GRiSP,  
Germany BMZ/GTZ, Syngenta-IRRI SKEP, GCP



# Thank You!

