# High-resolution field phenotyping: exploiting rice genetic diversity

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International Rice Research Institute Los Baños, Laguna, Philippines

IRRI

IR8 – 3x yield 1962, Peter Jennings

PETA

PETA

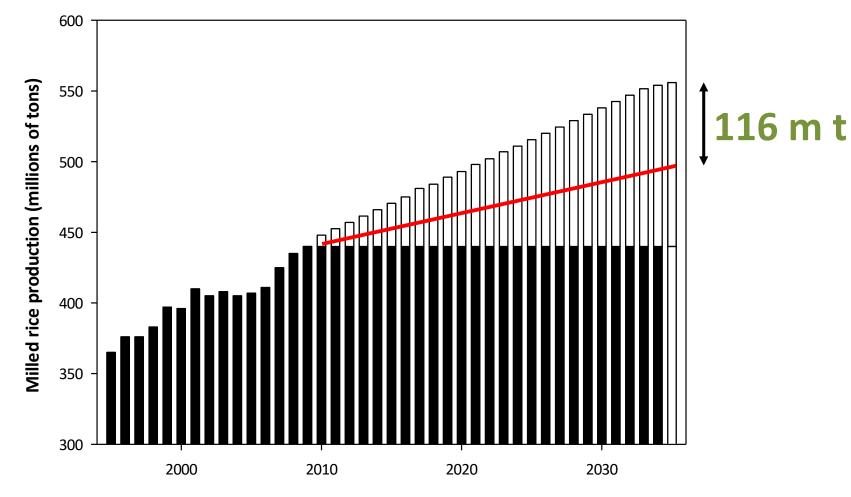
**IR 8** 

IRS

DGWG

DEWE

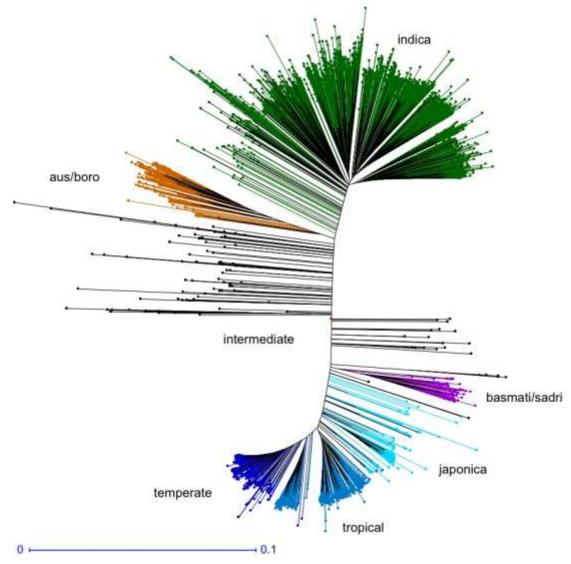
#### **Yield Gap**



Year

The International Rice Genebank 127,000 rice accessions and wild relatives

### The 3,000 rice genomes project



The 3,000 rice genome project 2014, GigaScience, 3:7.

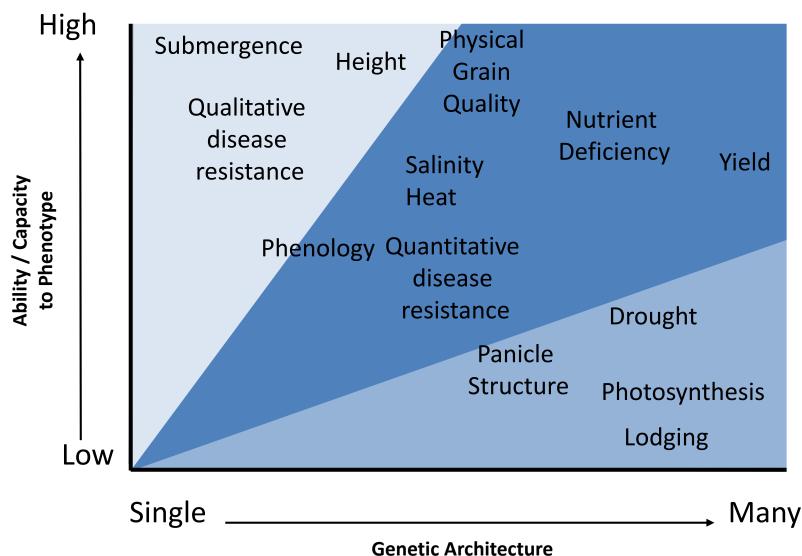
### Genotype to Phenotype



#### **Traditional Methods**

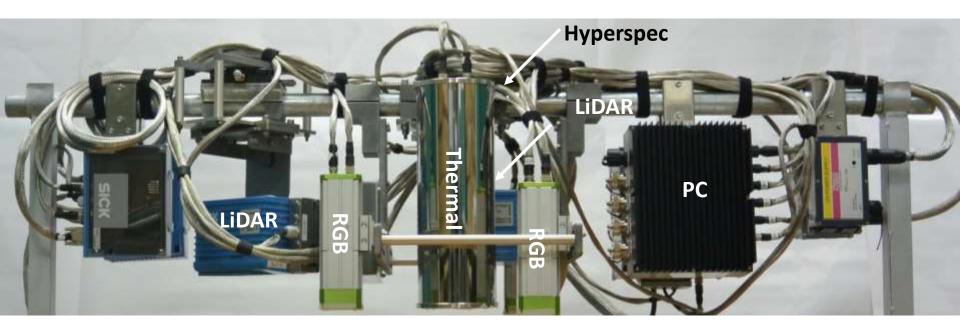


### **Trait Selection**



#### (number of genes controlling trait)

# **High-resolution Phenotyping**





# Paddy Field

a as as a second of the second **IRRI** Zeigler Experiment Station

# High-throughput phenotyping







### Paddy 'Field'



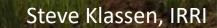
#### Automation and Control





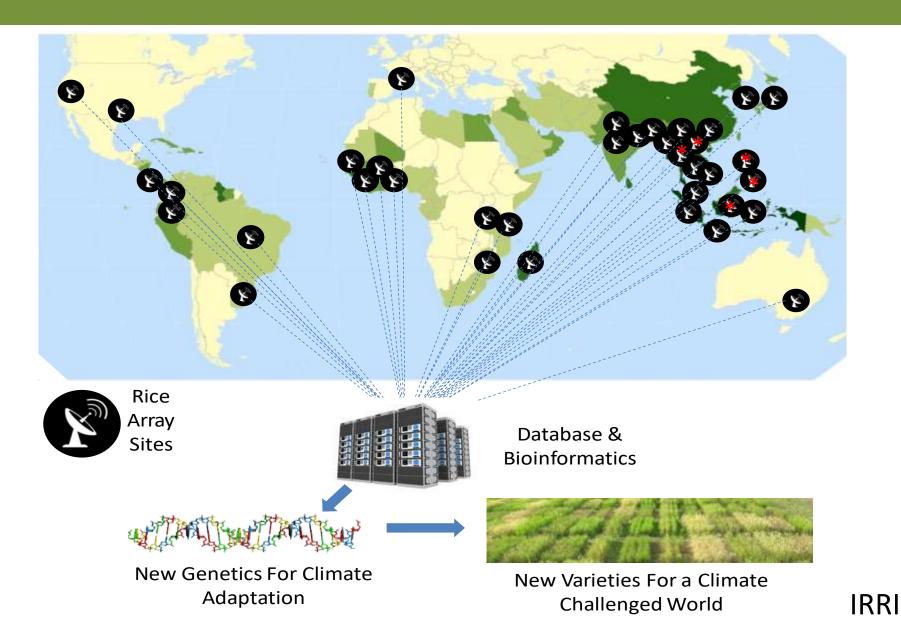


#### Scaling Up: IRRI Farm



IRRI Zeigler Experiment Station (Upland Farm)

### Scaling Up: Global Rice Array

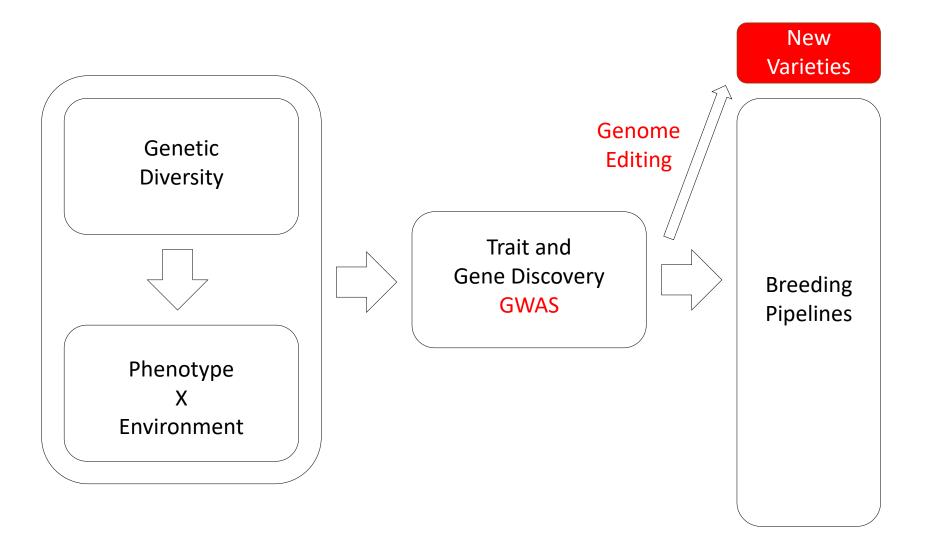


### **Future Challenges**

IRRI, Rwanda, East Africa



# Begin with the end in mind...

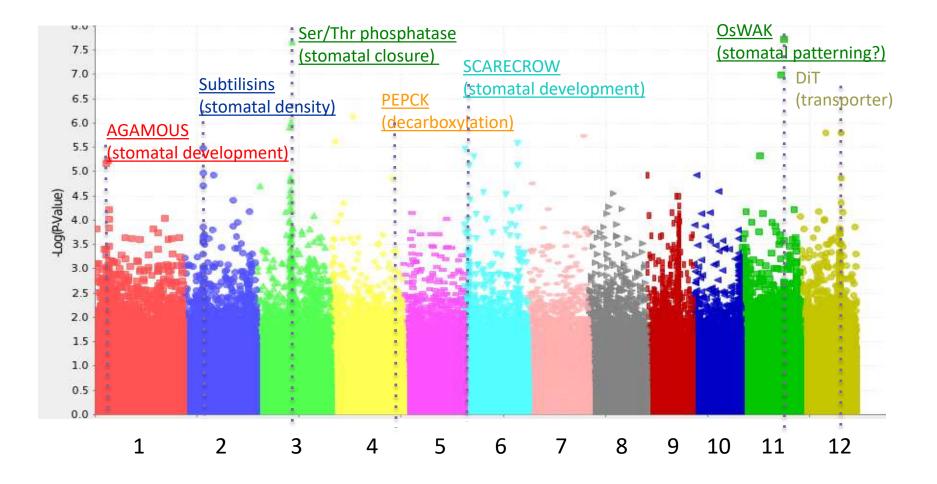


### Natural variation in photosynthesis



47 leaf photosynthesis and anatomy traits

# Intercellular CO<sub>2</sub> Concentration



TASSEL (MLM: PCA+Kinship)



# • Find out more about who we are and what we do: $C_4$ Rice Centre



# Acknowledgements

#### C<sub>4</sub> Rice Centre

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