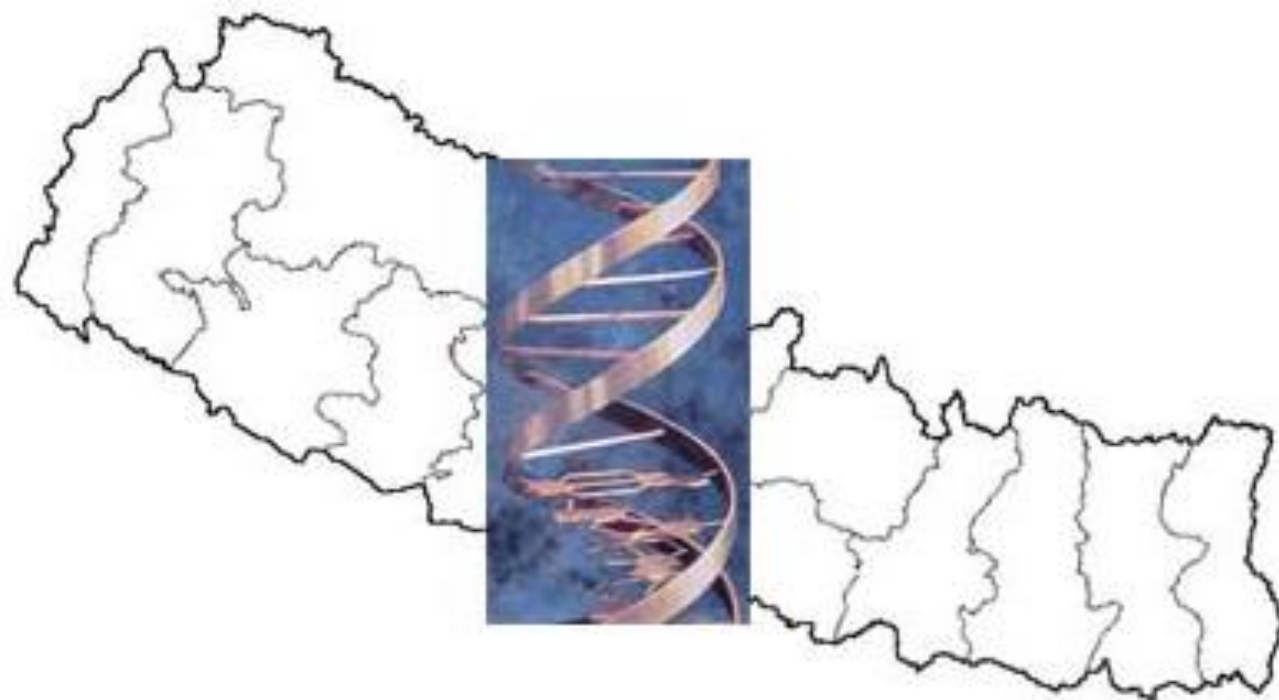


Biochemical and Molecular Markers in Plant Breeding

Progress and Prospect



Bal K. Joshi
Feb 2008

Plant Breeding

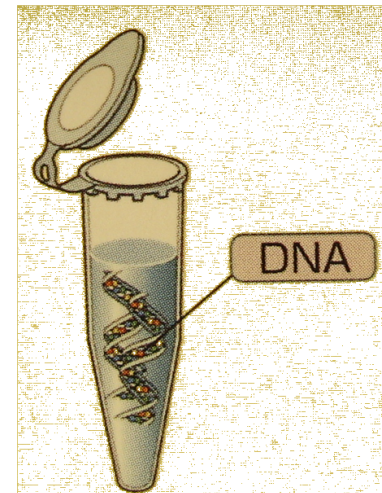
- The science, art and business of improving plants for the human benefit

Plant Breeding Era

Era 1. Domestication and phenotypic mass selection

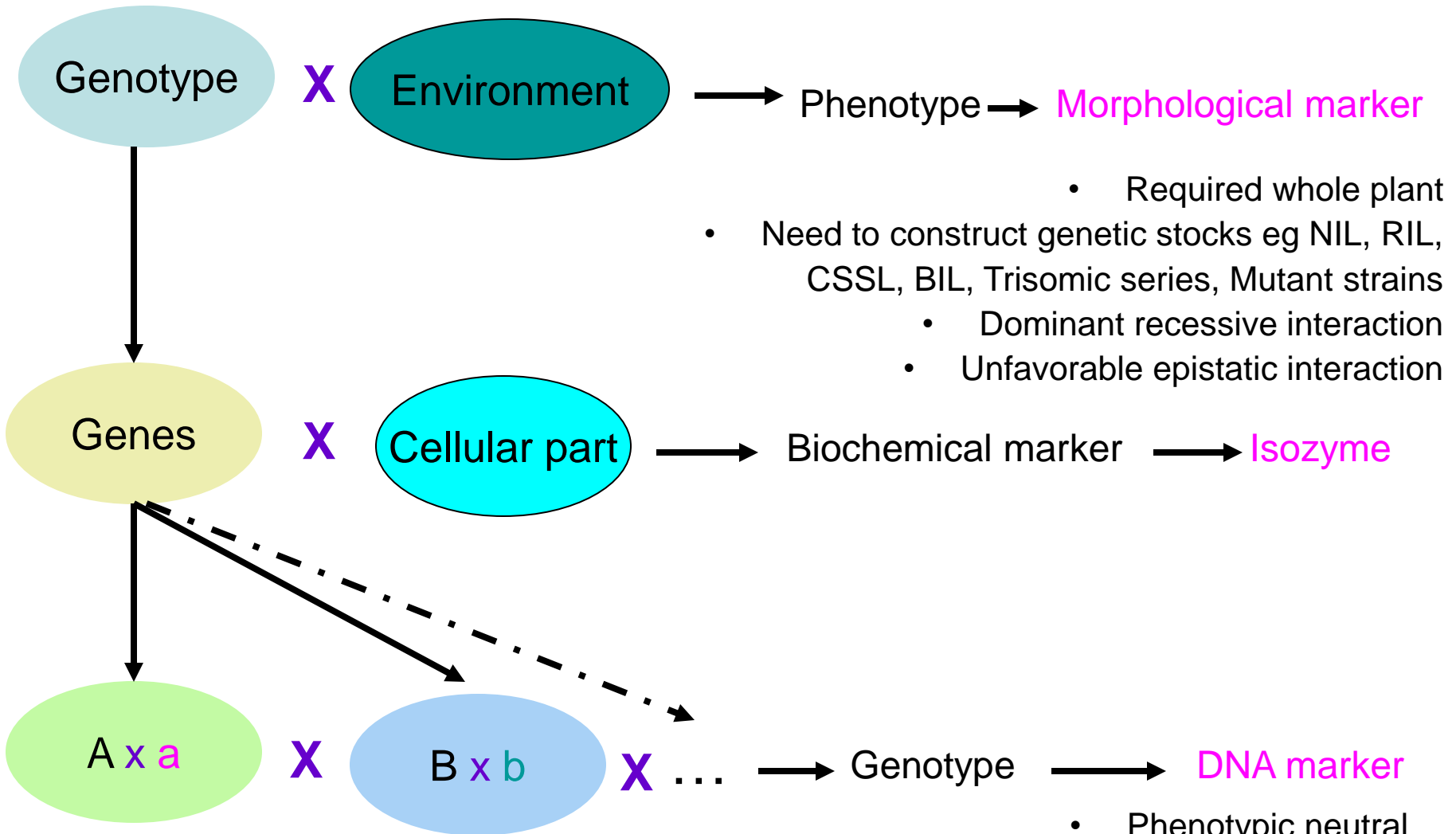
Era 2. Replicated progeny selection

Era 3. Direct genotypic selection



Marker systems

- Gene tagging
- Indirect selection
- Identification etc

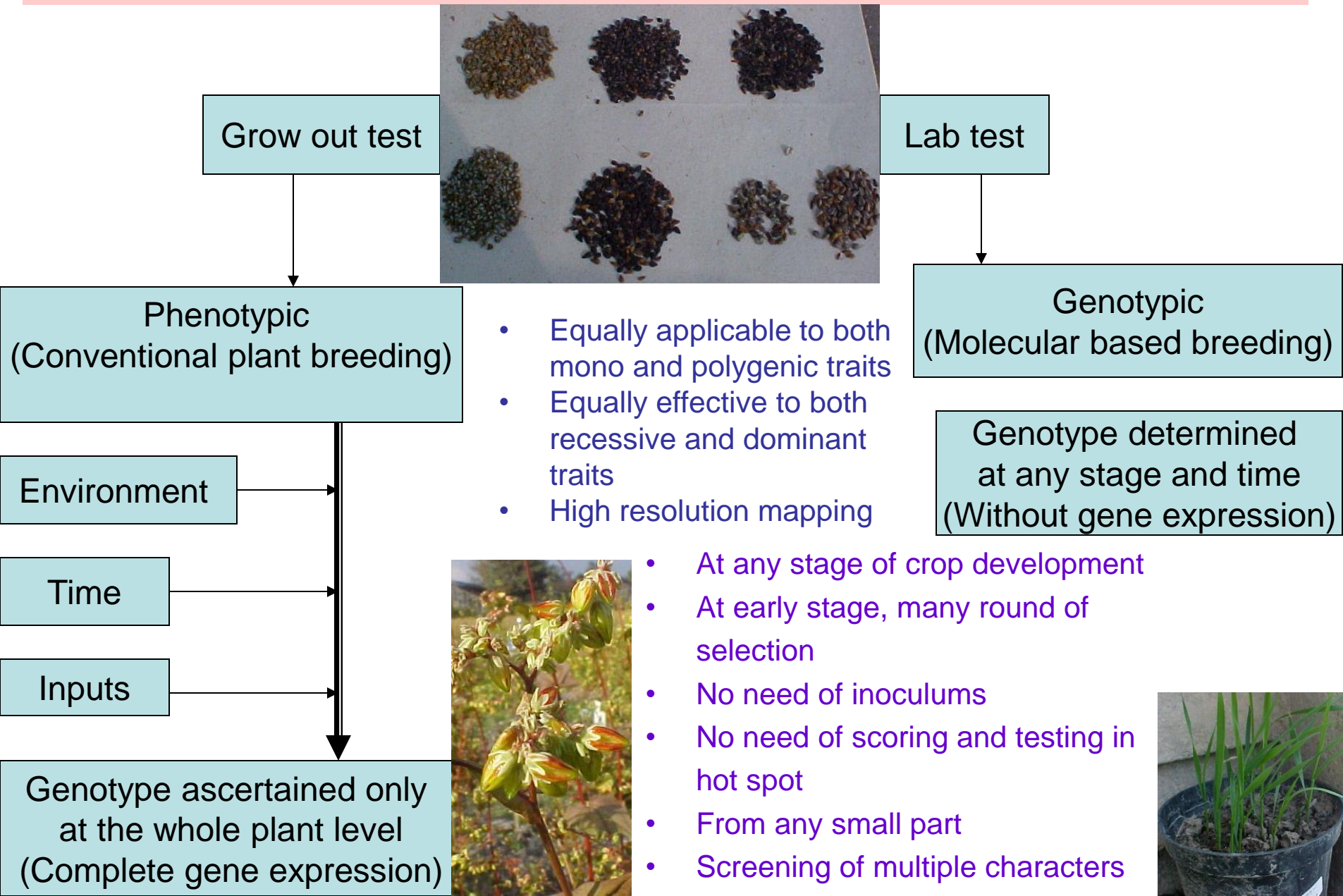


- Required whole plant
- Need to construct genetic stocks eg NIL, RIL, CSSL, BIL, Trisomic series, Mutant strains
 - Dominant recessive interaction
 - Unfavorable epistatic interaction

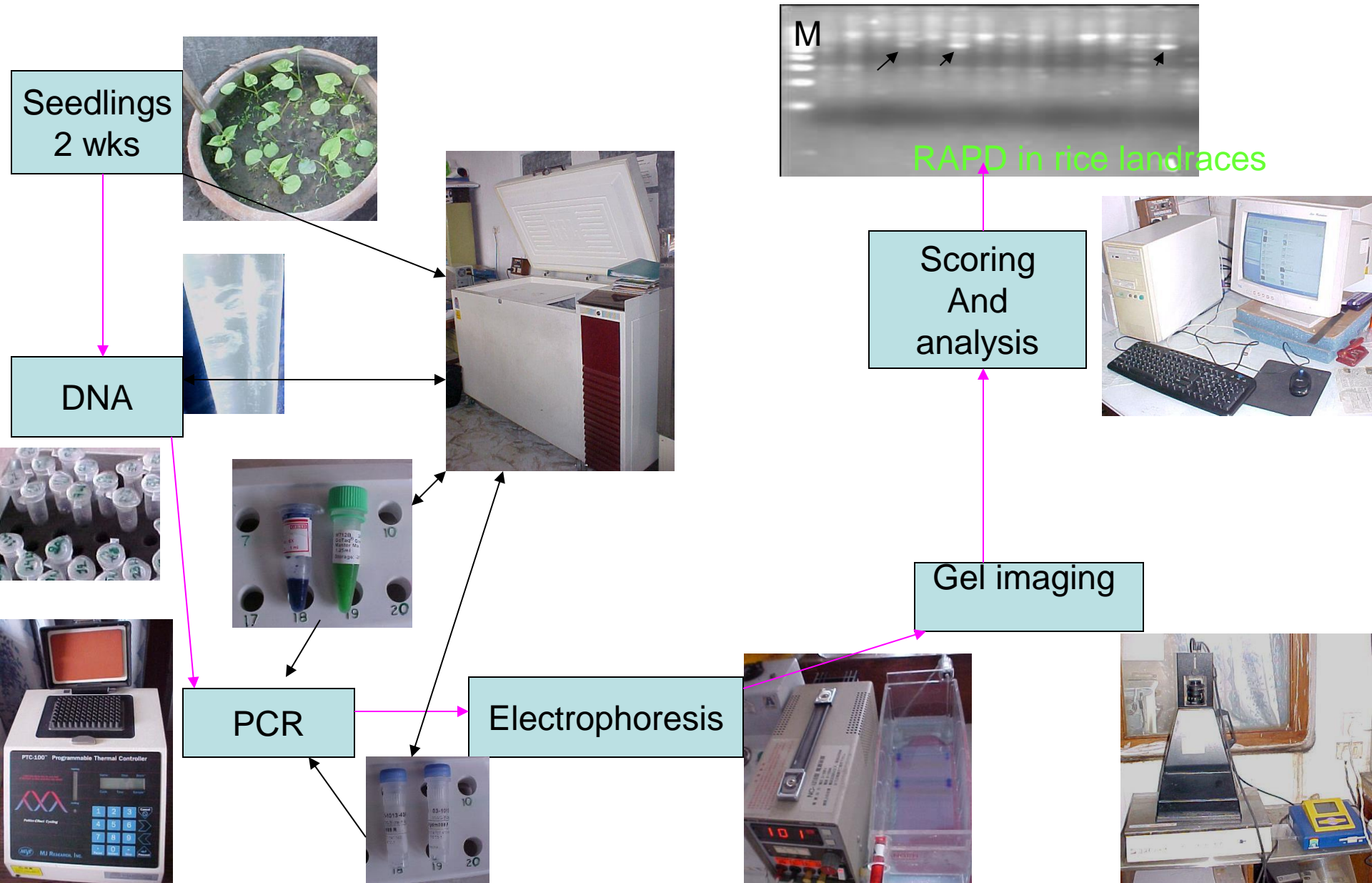
Dominant marker: RAPD, Morphological...
 Co-dominant markers: SSR, EST, Isozymes ...

- Phenotypic neutral
- Codominant manner
- Free of epistatic effect

Which one is the best in term of yield &/or disease resistance?



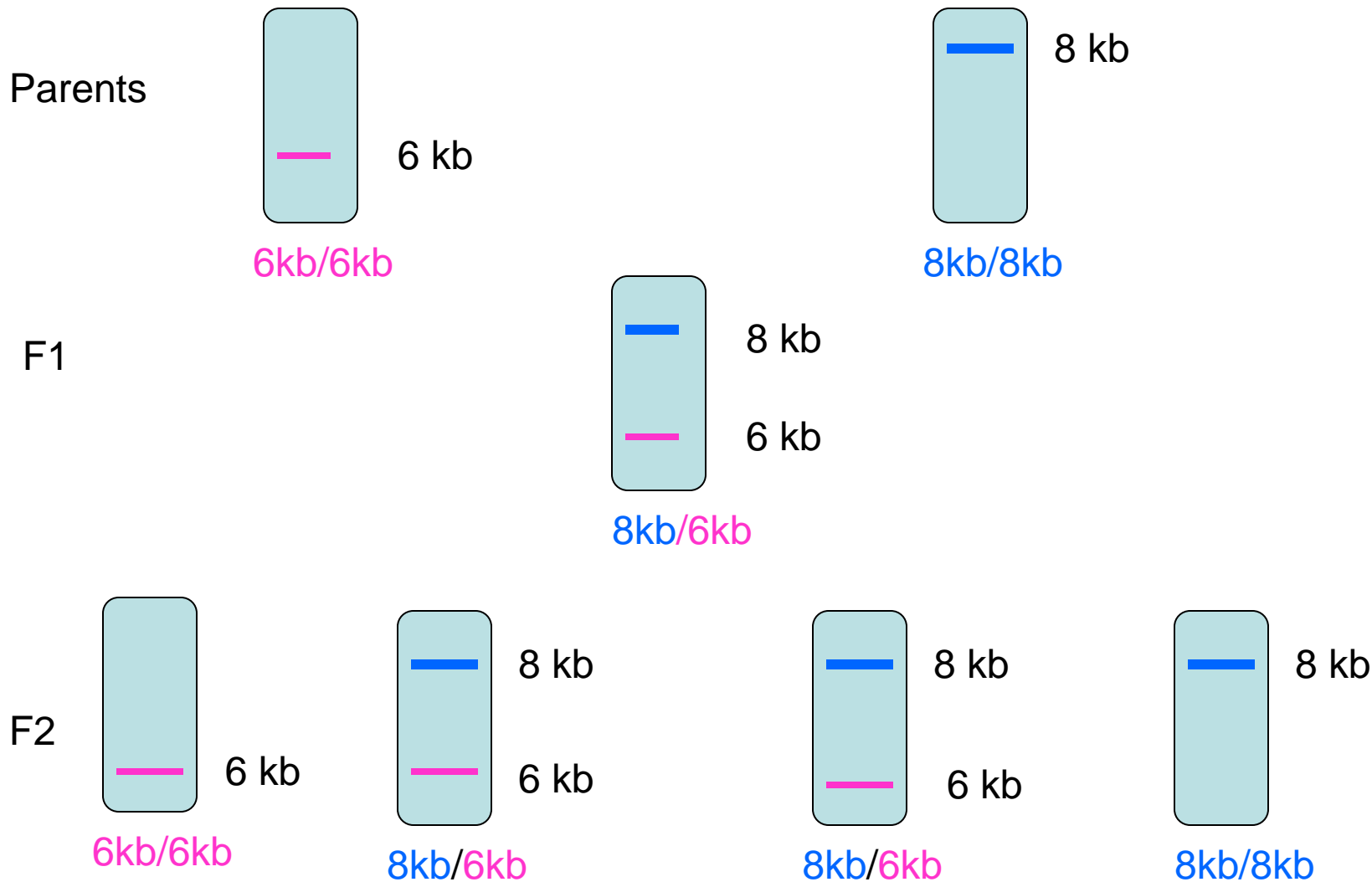
Experimental steps in molecular based research



Inheritance of SSR marker

- dominance, recessive
- allelism test, number of alleles

QTL also in Mendelian principle



Achievements

Diversity study

- **SSR**: rice, barley
- **RAPD**: wild and cultivated buckwheat, rice
- **Isozyme**: wild and cultivated rice, barley, mango, pigeon pea, buckwheat, swertia, taro, citrus

Running activities in rice and buckwheat

- Molecular tagging of blast resistance gene
- Marker Aided Selection (MAS) and Marker Aided Backcrossing (MAB)
- Developing NILs
- Linkage map construction
- QTL mapping

Software for analysis of biochemical and molecular data

- NTSYSpc
 - PHYLIP
 - Tree View
 - Free Tree
 - WinAMOVA Arelquin
 - GGT
 - PhotoCapMw
- | | |
|----------|------------------|
| GDA | QTL Cartographer |
| GenePop | MapMaker/EXP |
| PopGene | MapMaker/QTL |
| GeneStat | Scion Image |

Prospects

- Information based on the DNA is considered the most reliable
- Opportunity to select desirable lines based on the genotypes rather than phenotypic- extremely attractive to plant breeders
- Must identify marker linked to target traits

1. Characterization
2. Identification
3. Screening
4. Diversity and biosystematics study
5. Population structure
6. Linkage map construction and gene tagging
7. MAS and MAB
8. Comparative mapping

A. Characterization

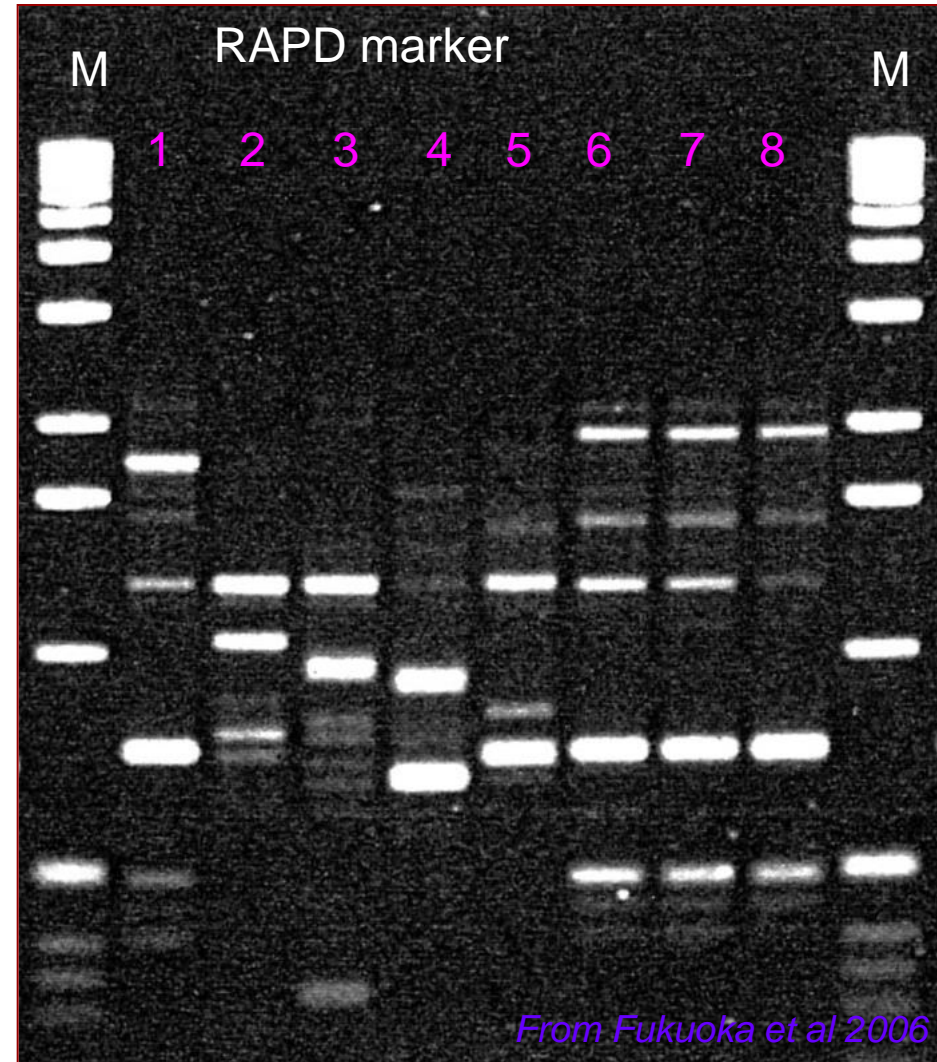
Descriptive statistics

- No. of bands, Size, Homo/heterozygous, Private allele

Diversity indices

- Observed/ Expected heterozygosity
- Effective/ Number of alleles/locus
- Percent polymorphic loci
- Shannon's gene diversity
- Gene/genotype frequency
- PIC

- A. Characterization of accessions
- B. Characterization of markers

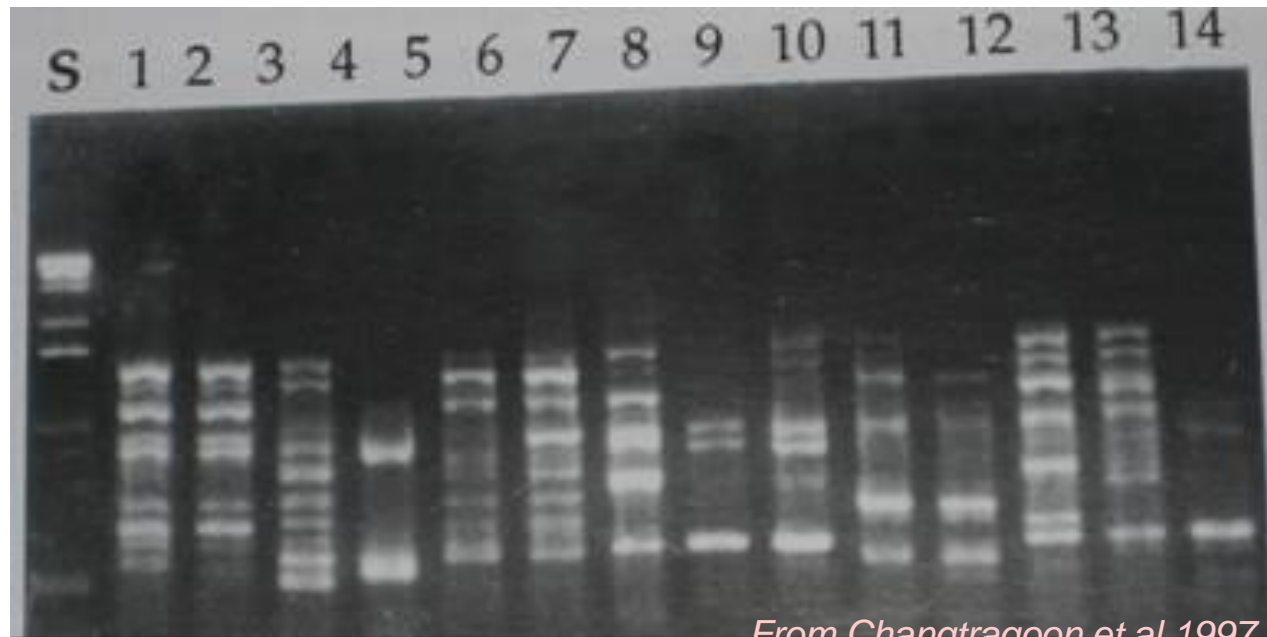


B. Identification

- **Uses:**
- Plant variety/ cultivar protection, plant patenting
- Hybrid testing
- PGR management (duplicate identification)
- Breeding works (parent selection, introgression etc)

- Similarity/ dissimilarity in bands
- Private allele/s
- Fingerprinting
- Standard

RAPD fragments amplified by OPA 13 primer in 10 spp of rattan showing species specific polymorphism (1=2, 3, 4, 5=6, 7, 8=9, 10=11, 12=13, 14, S= 1kb ladder



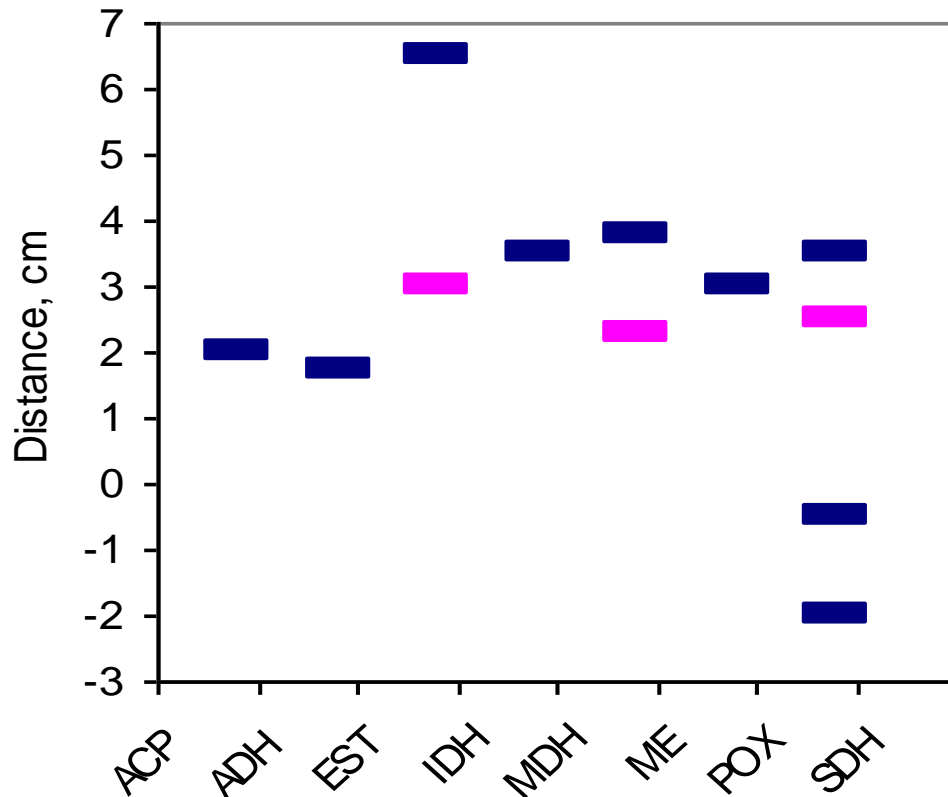
From Changtragoon et al 1997

Fingerprinting (Genotyping)

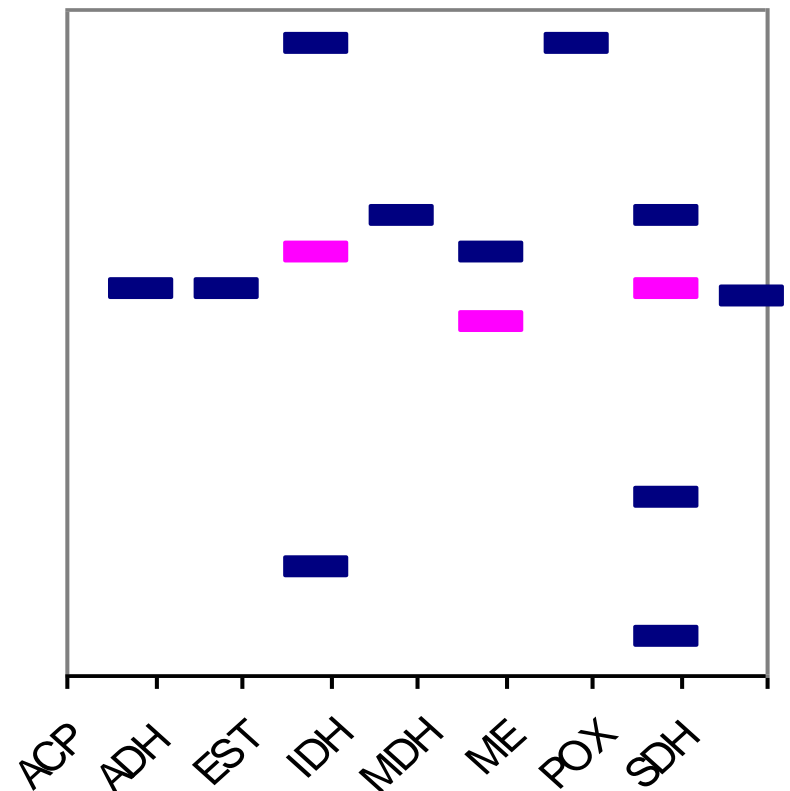
- 15 individual samples per variety in cross pollinated crop species
- 1 individual sample per variety in self pollinated crop species

Isozyme profiles of rice cultivars (Multiple markers band)

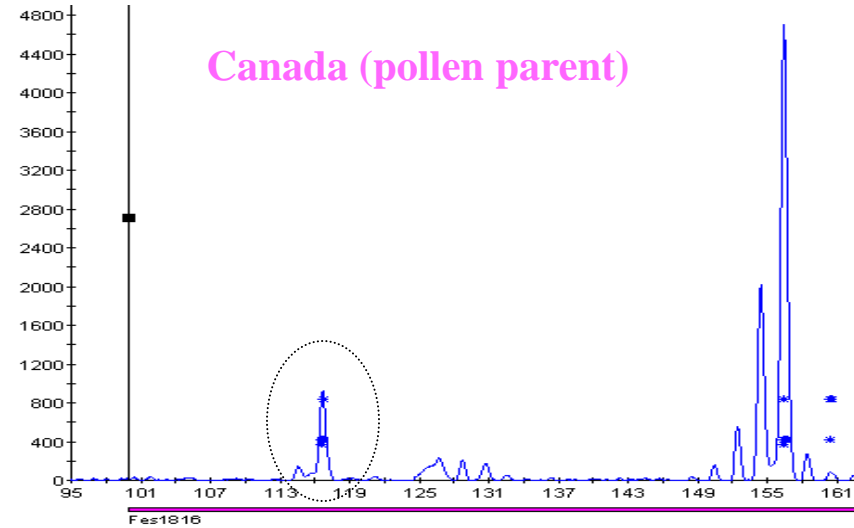
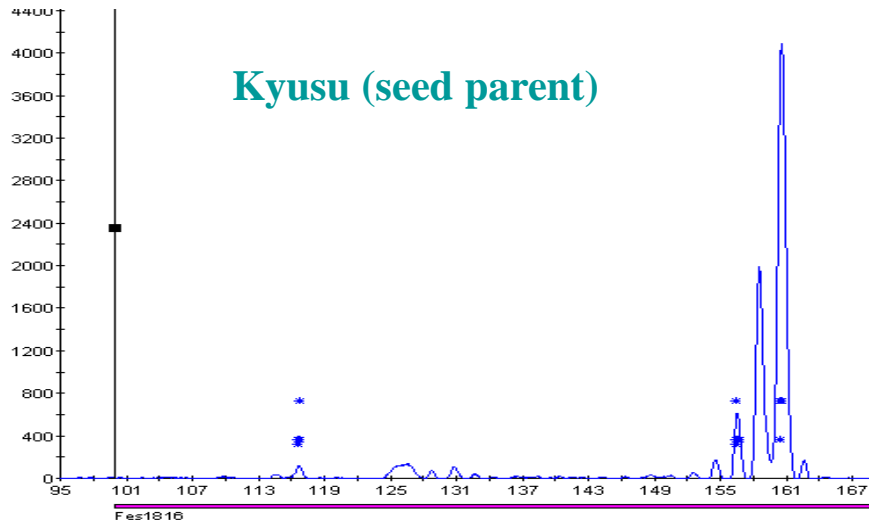
A. Chhomriong



B. Khumal 4

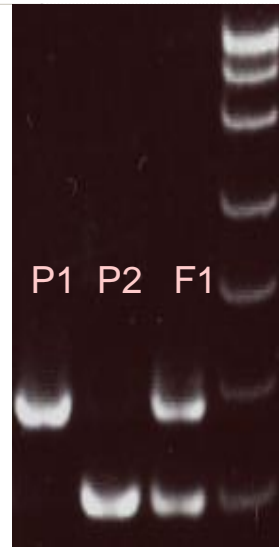
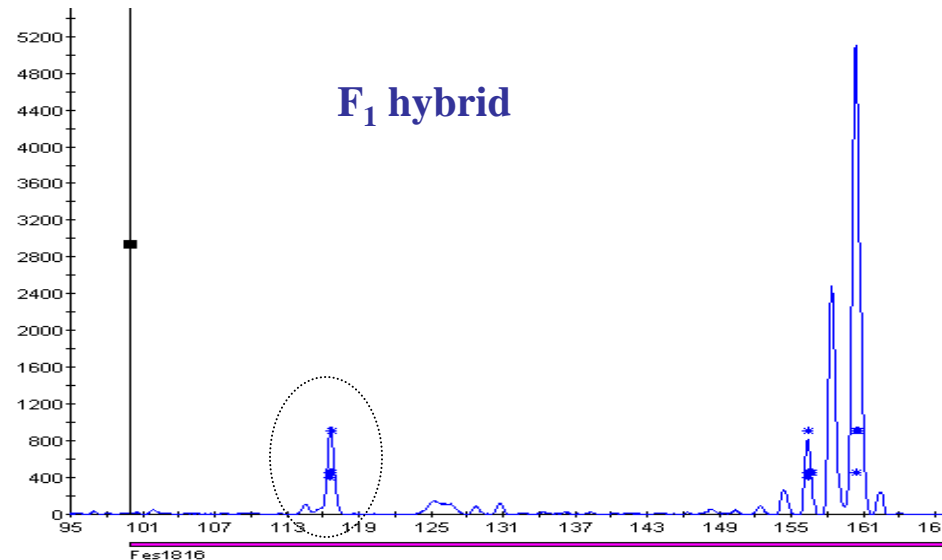


DNA sequence based F_1 testing



Primer: Fes 1816

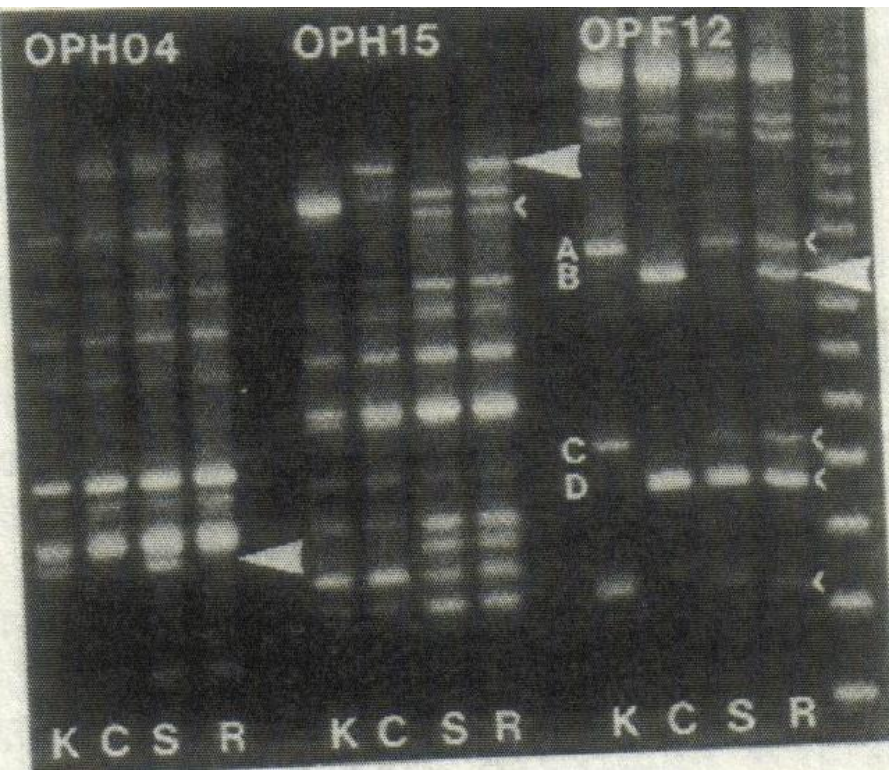
**SSR
electropherogram
clearly distinguished
 F_1 hybrid**



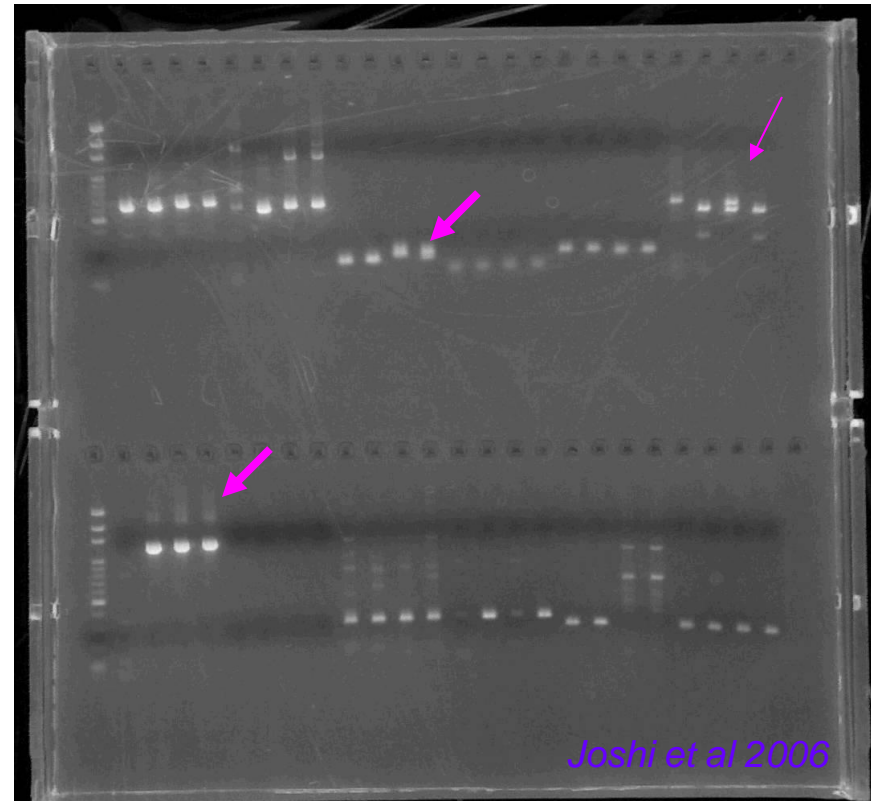
C. Screening

- Resistance and susceptible
- Cold tolerance
- Drought tolerance
- High protein content

RAPD markers detecting downy mildew resistance gene in lettuce parent and progeny bulk



Different EST markers in 4 lines of common buckwheat



D. Diversity and biosystematics study

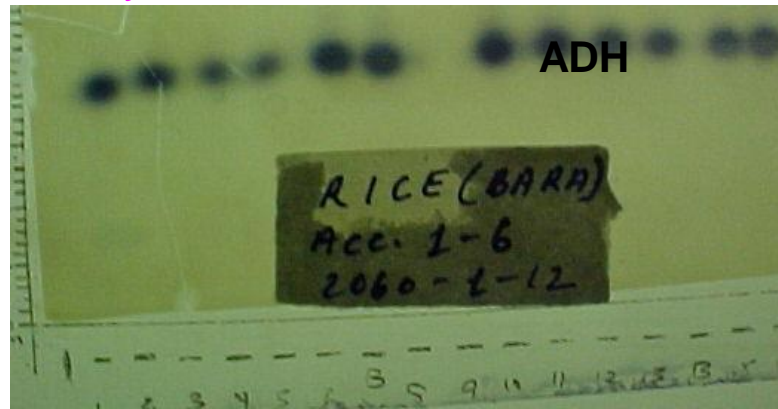
Diversity indices
Genetic distance

- Nei's
- Roger's

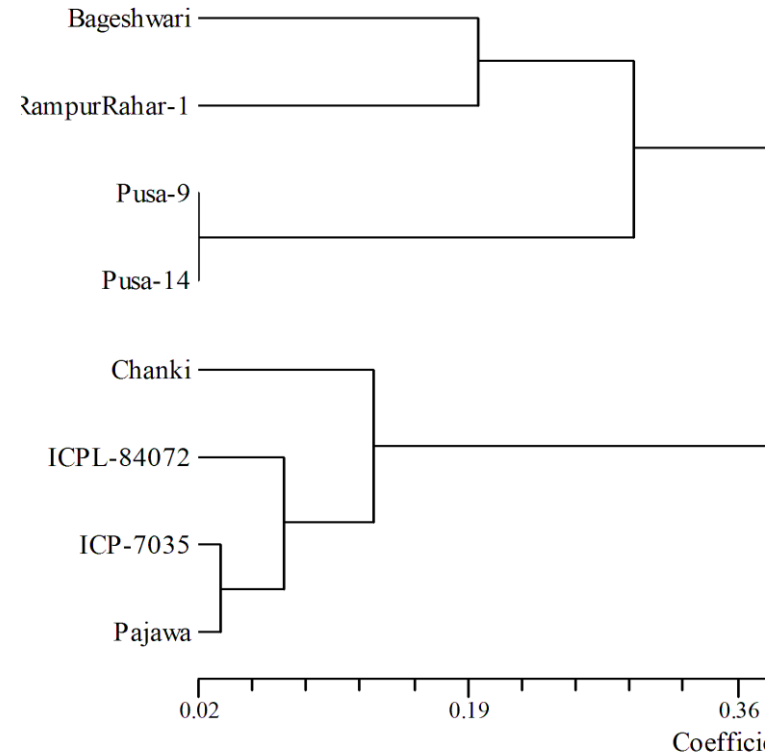
Multivariate analysis

- Cluster
- PCA

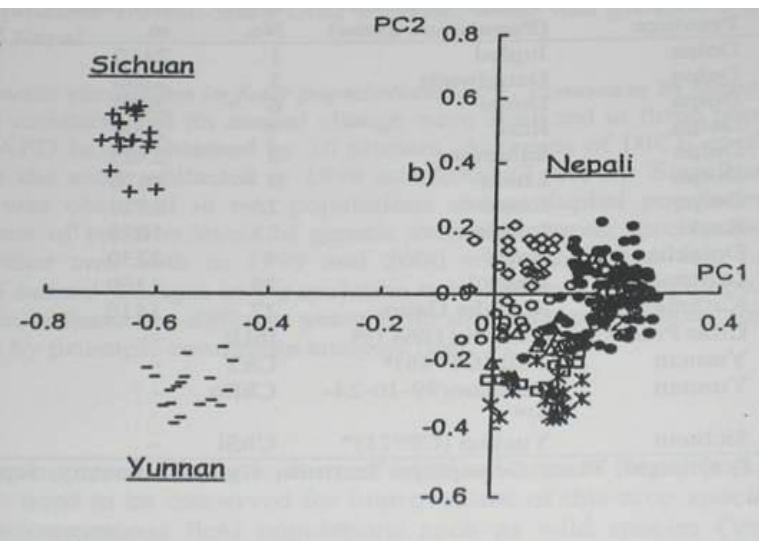
Isozyme bands in Bara rice landraces



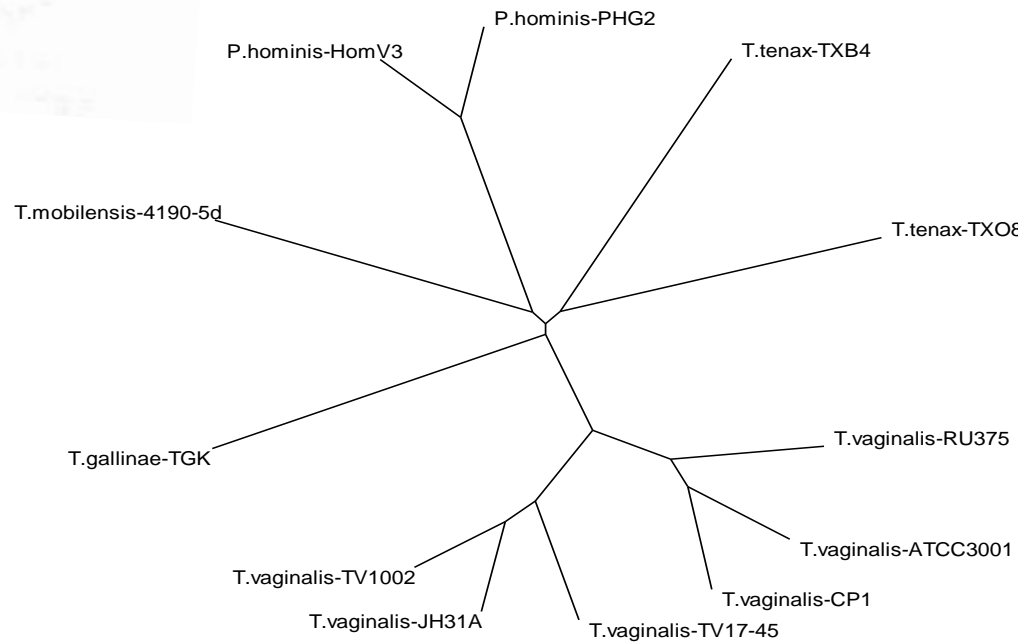
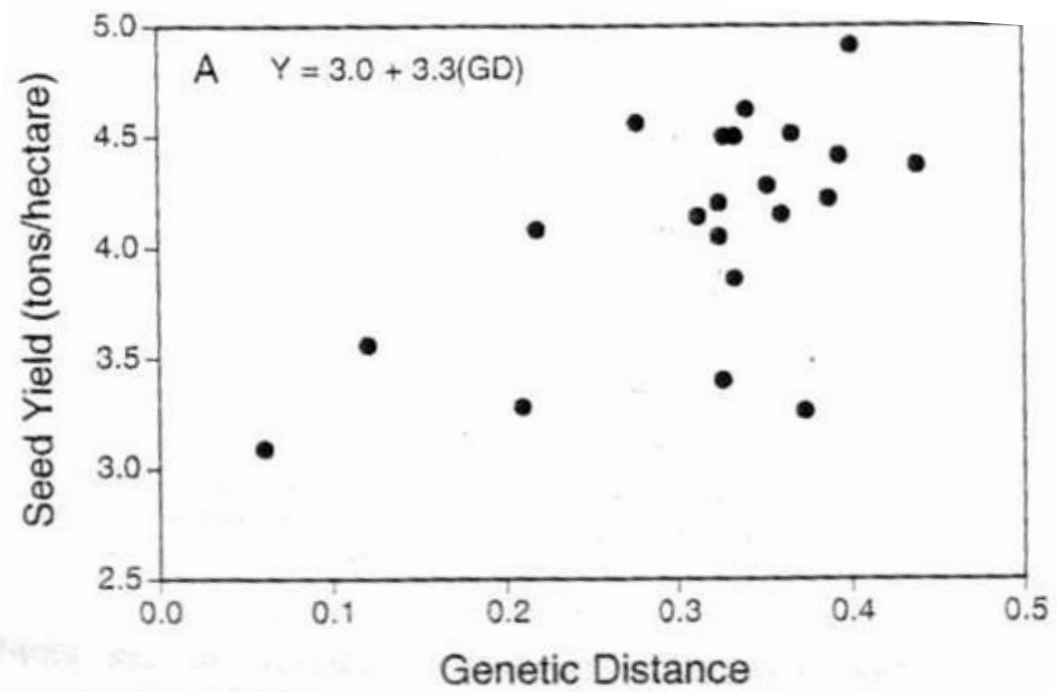
Dendrogram of 8 pigeon pea populations based on isozymes with Nei's genetic distance



Principal component analysis based on 46 RAPD primers in *Fagopyrum cymosum*



Divergence and Heterosis study



Phylogenetic (radial) tree of trichomonads based on 731 RAPD traits: Phylogeny and evolution

E. Population structure

Parameters

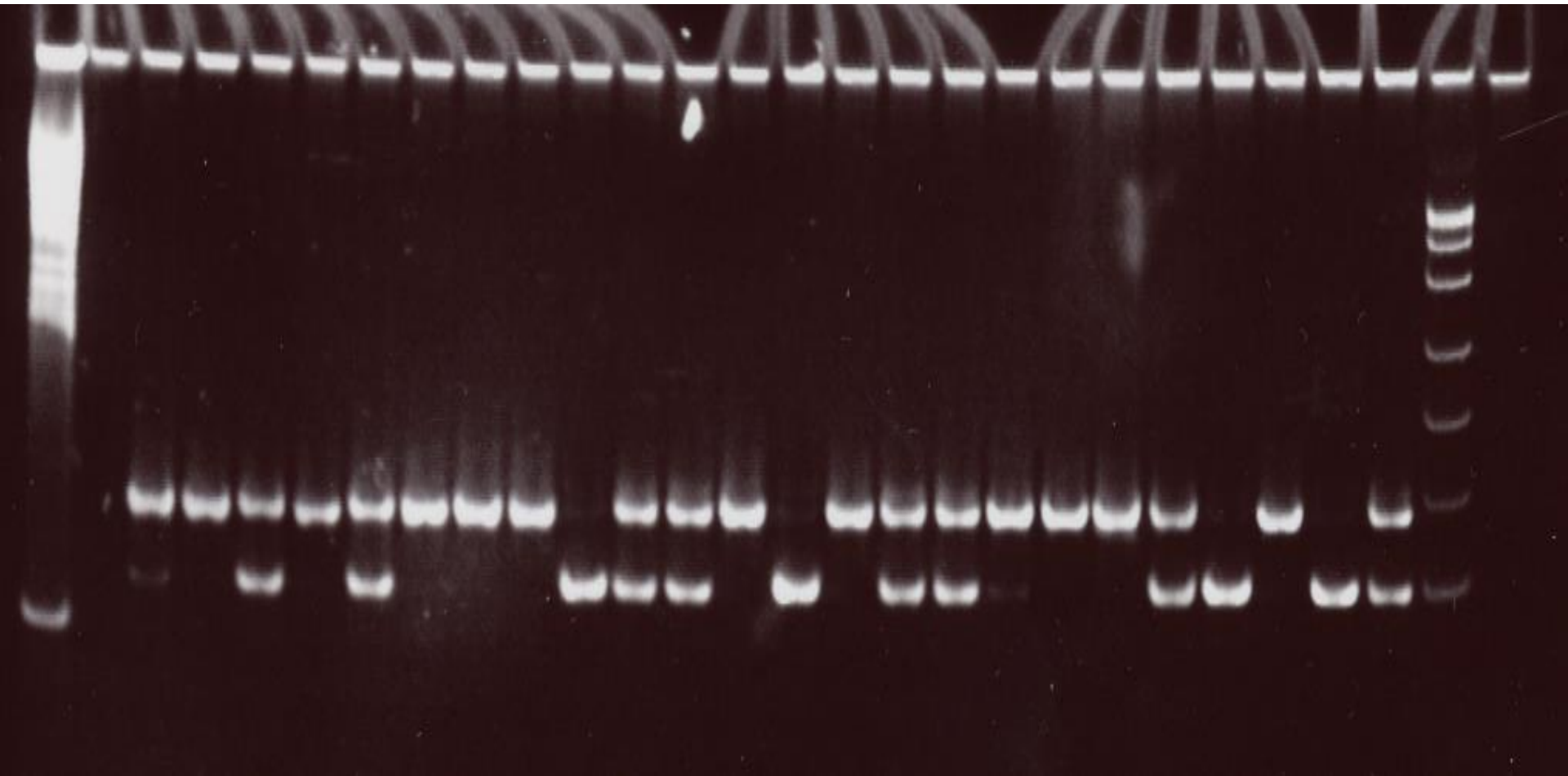
- Wright' F statistics
- Nei's G statistic
- Rho statistics

AMOVA, partitions of variations

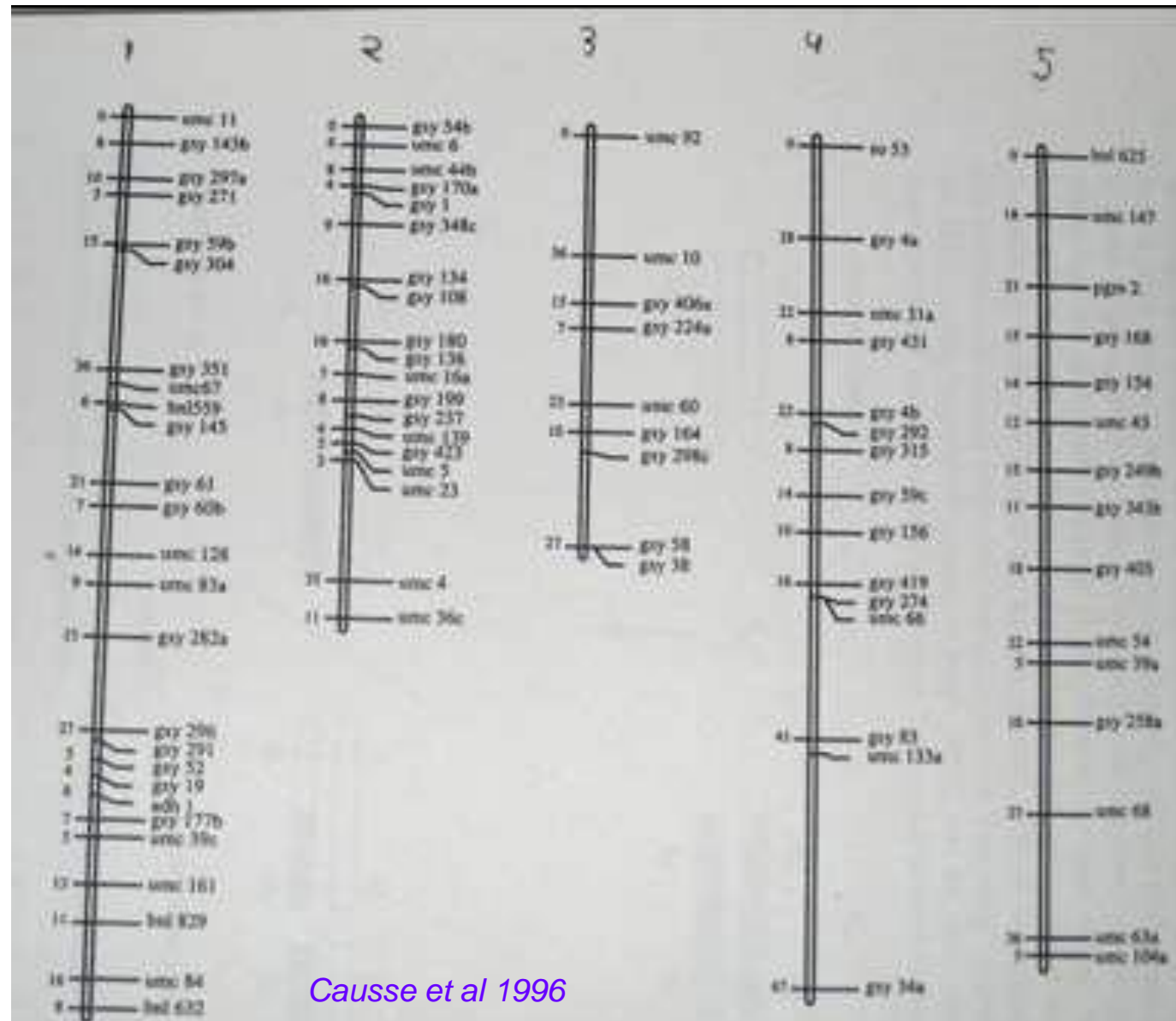
Source of variance	df	SS	EMS
Among groups, species	G-1	SSD (AG)	$n''\sigma_a^2 + n'\sigma_b^2 + 2\sigma_c^2 + \sigma_d^2$
Among populations/ within groups	P-G	SSD (AP/WG)	$n\sigma_b^2 + 2\sigma_c^2 + \sigma_d^2$
Among individuals/ within populations	N-P	SSD (AI/WP)	$2\sigma_c^2 + \sigma_d^2$
Within individuals	N	SSD (WI)	σ_d^2

F. Linkage map construction, gene tagging and QTL mapping

CAPS marker in parents and F2 population of common buckwheat



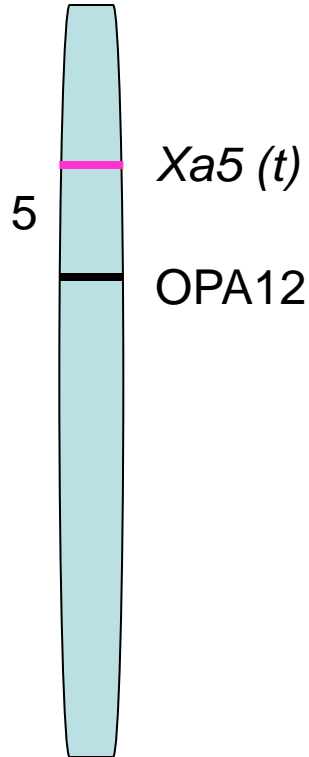
A linkage map of maize using cDNA markers in 5 chromosomes



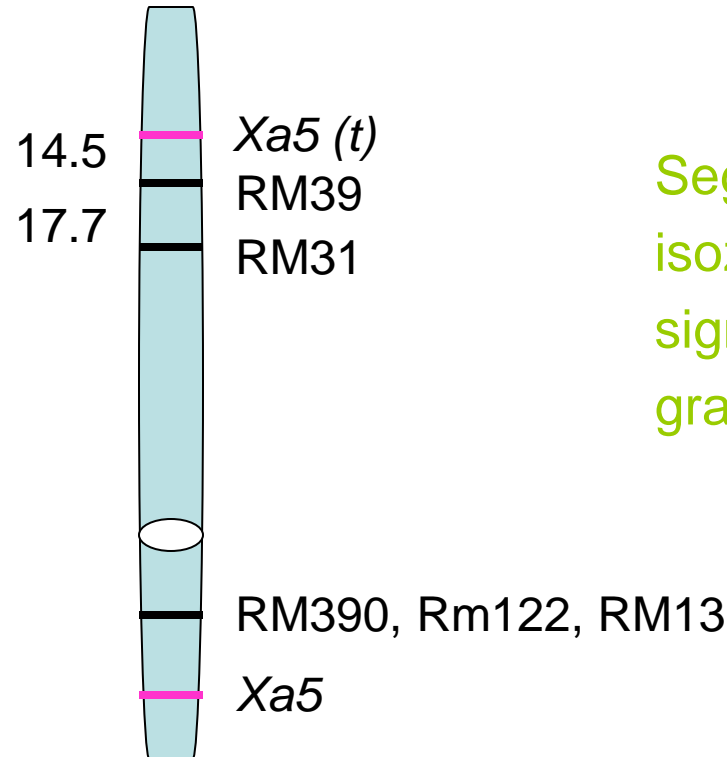
Causse et al 1996

RAPD and SSR markers linked to bacterial blight resistance gene of rice

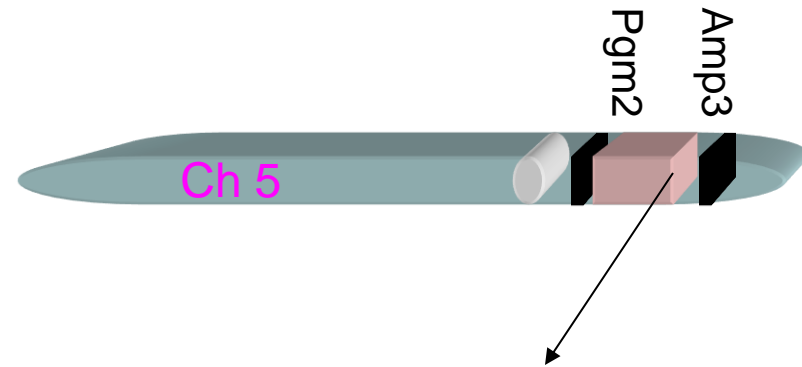
RAPD maker



SSR markers

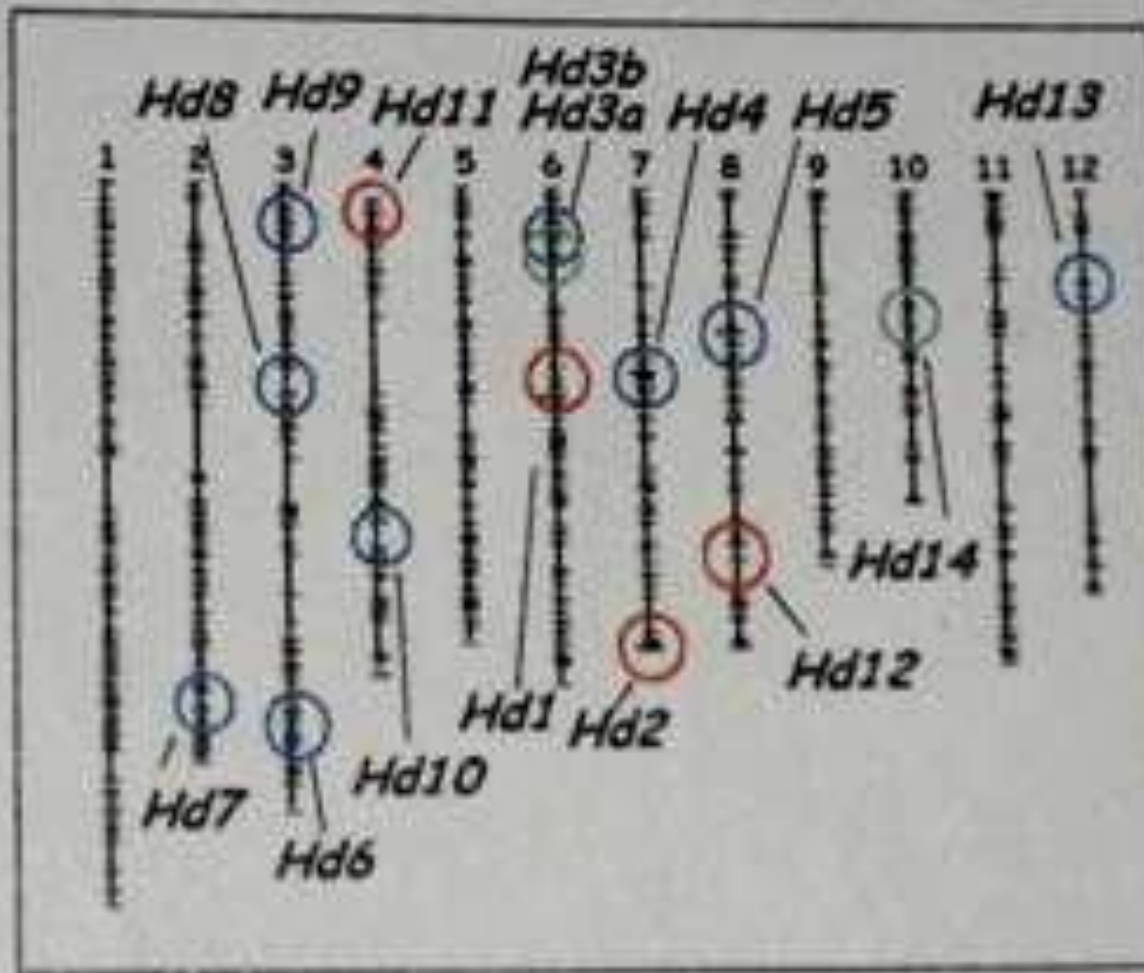
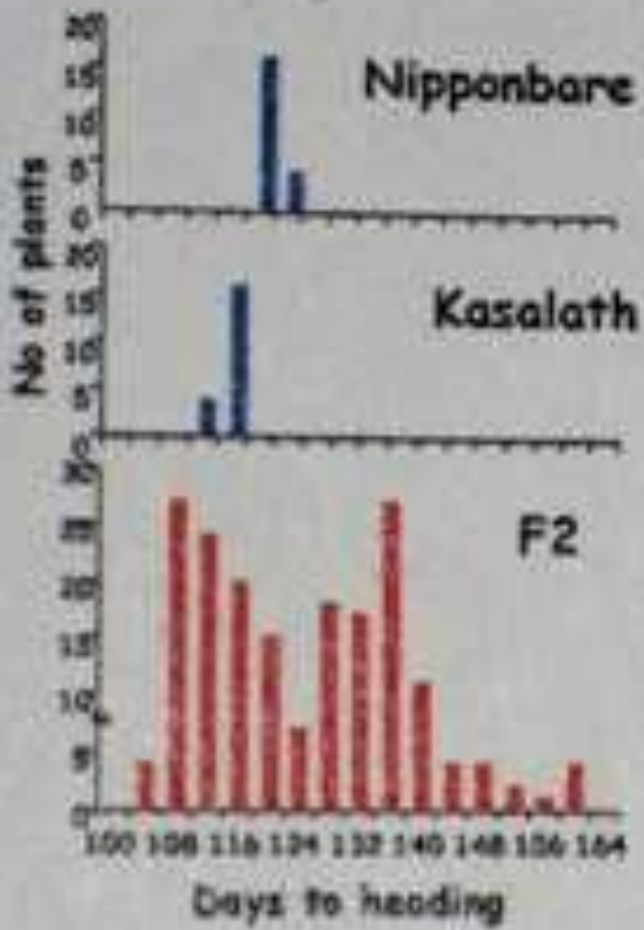


Ch 5

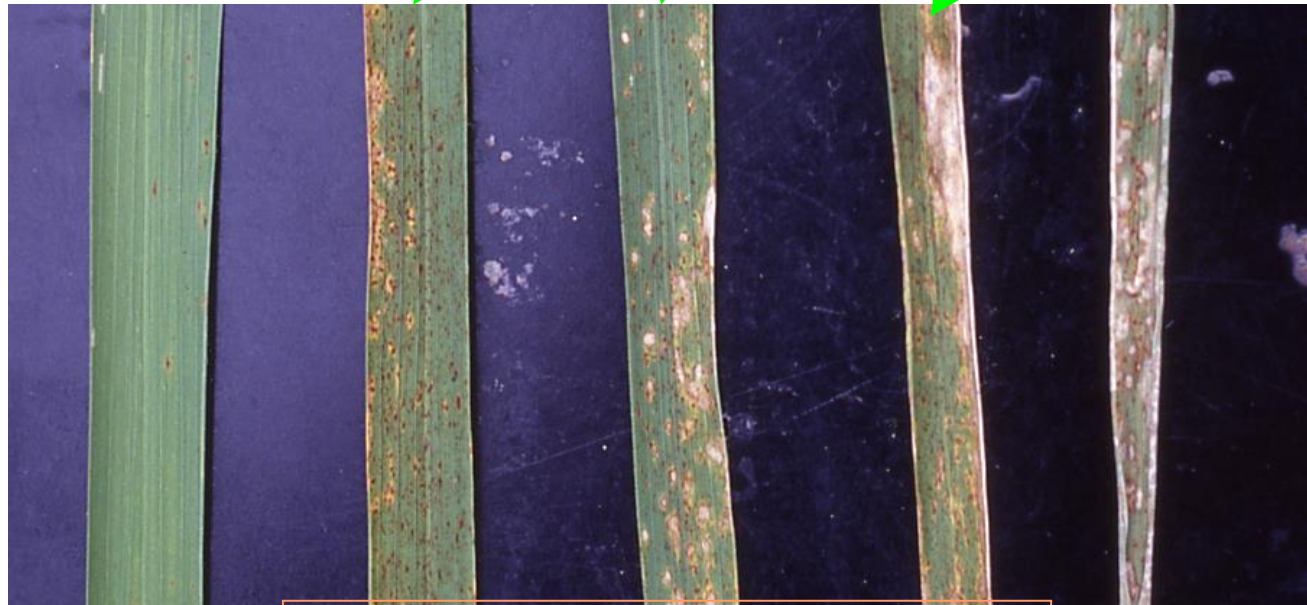
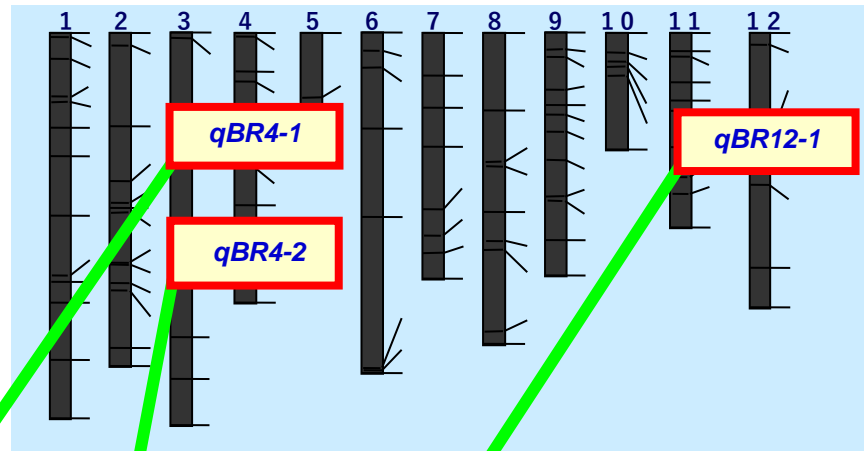


Segment bracketed by isozyme markers: very significant effect on grain yield in maize

QTL mapping for heading date in rice = ETL



Effect of each QTL for rice blast



pi21
(*qBR4-1*) *qBR4-2* *qBR12-1*

upland

lowland

From Fukuoka et al 2006

G. Marker assisted selection (MAS) and marker assisted backcrossing (MAB)

Phenotypic selection

Genotypic selection

Combined selection

- Parents selection/ tagging desired traits
- Crossing
- Marker assisted selection in segregating population
- Graphical genotyping (Foreground and background selection)

- At any stage of crop development
- Many round of selection
- No need of gene expression
- No need of sample destruction
- Large portions of recurrent parent
- No need of selfing in case of recessiveness of gene

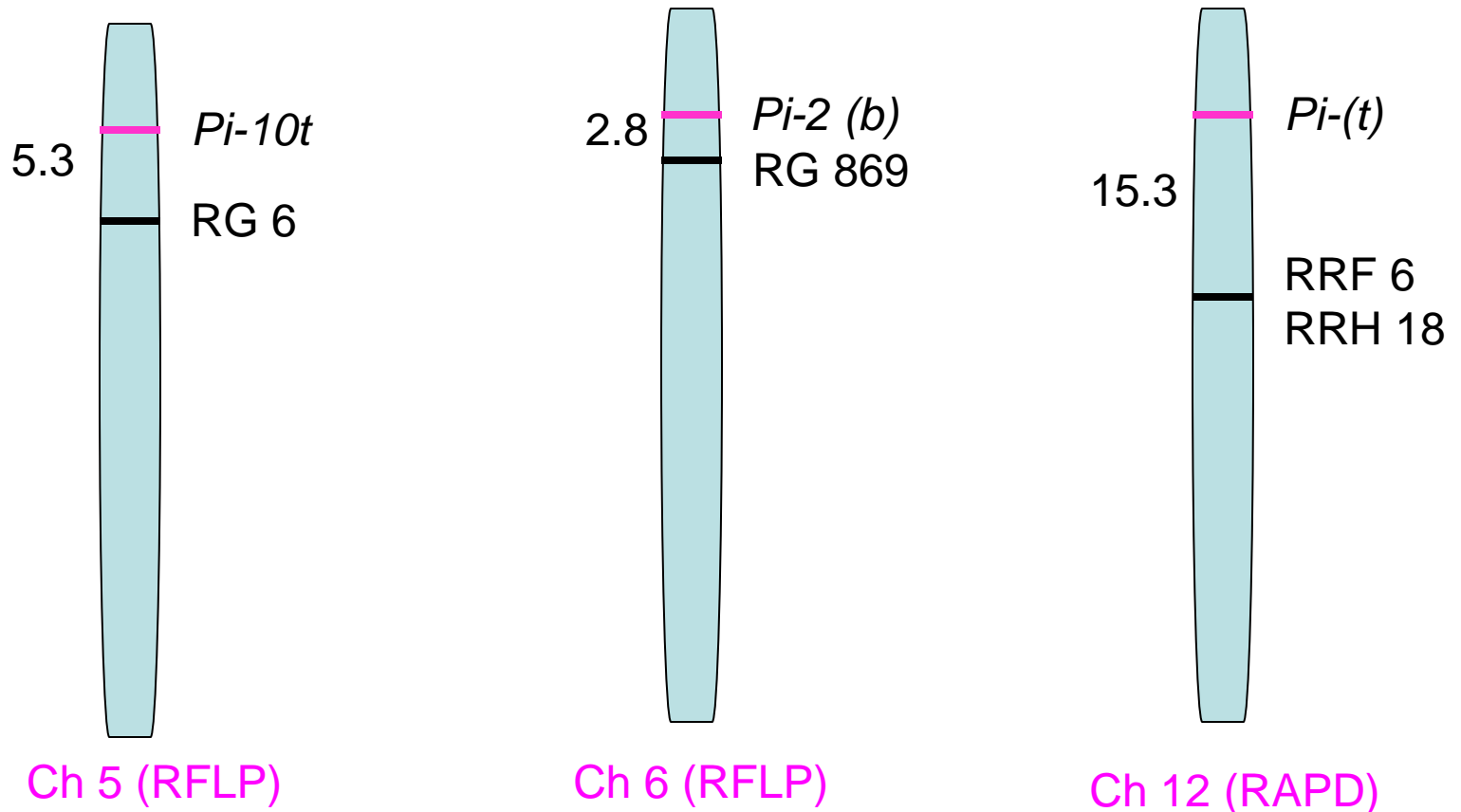
Co segregation of *Pi-g(t)* and SSR marker RM166 in the BC1 population.



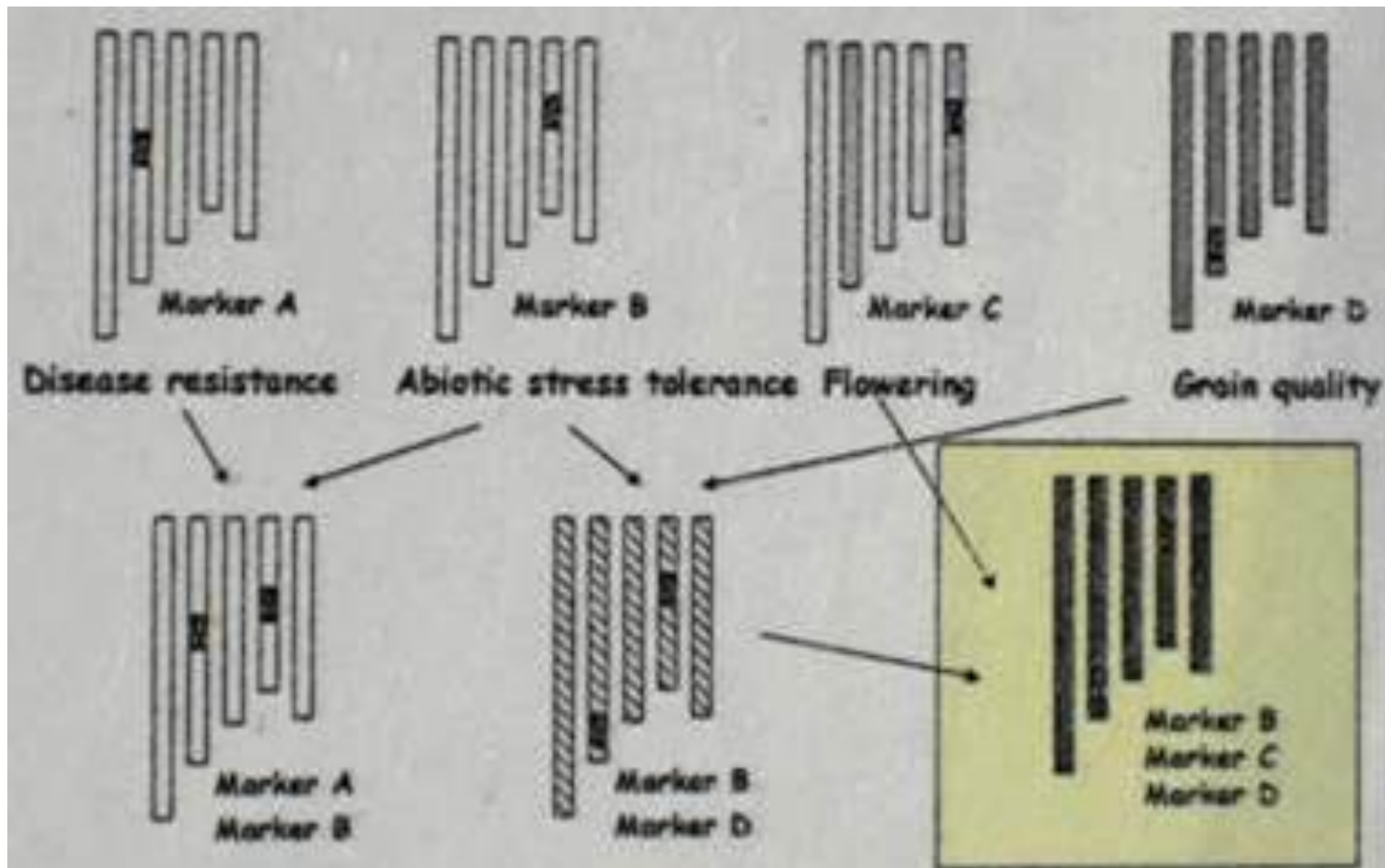
M P₁ P₂ 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 M



Different markers linked to blast resistance gene in rice

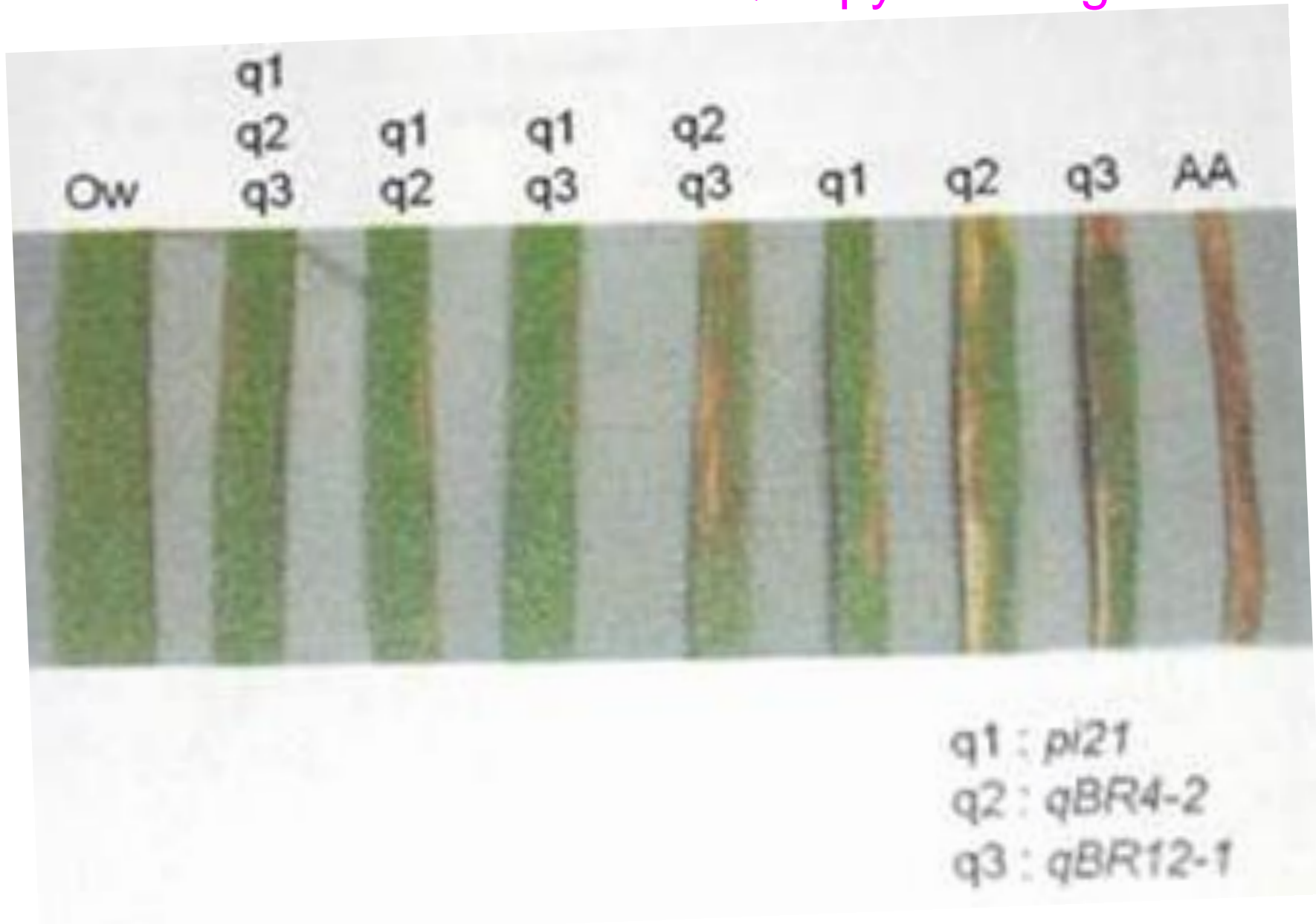


Gene pyramiding



Selection can be performed based on the genotypes instead of the phenotypes

Durable rice blast resistance QTL pyramiding



Comparative study

- Vavilov's law of homologous series in variation
- DNA markers can be utilized among different crop species, not possible in case of phenotypic markers
- Use of complete DNA sequences of Rice, Bacteria, Fungi, Human, Drosophila, Arabidopsis
- Evolutionary study



Scope

- Any areas, if your material content DNA

Issues

Technical

- Biochemical or DNA base
- PCR or non PCR base
- Arbitrary (single primer) or site specific (pair primer)
- Dominance or codominance
- Exon or intron base
- Single or multi locus

Operational

- Cost
- Time
- Protocol
- Skill etc

Molecular marker technology

- Should be integrated in our plant breeding program
- Never replace the conventional plant breeding
- Complementary to conventional plant breeding (improve the efficiency and precision)

Plant breeder

- Fingerprinting
- Linkage map construction and QTL mapping
- Graphical genotyping
- Gene tagging
- MAS/ MAB (Parents selection, desired genotypes selection, NIL development, screening)
- Gene pyramiding