

# Resilience Breeding for Food Security



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# Agriculture, Pakistan and Food Security

GDP Share 25%

Labour 48%

Export Earnings >80%

## Agriculture Dynamics

### Food security

- Self sufficient in cereals, sugar, fruits, vegetables
- Importer of edible oil, pulses, cotton, tea, dry milk

### Nutritional security

- Around 50% population suffers nutritional deficiency
- Nearly 44% children are stunted

## Challenges

- Population
- Water
- Climate change
- Land
- Pest and diseases
- Salinity and water logging

## Population Growth

|      |   |             |
|------|---|-------------|
| 1951 | → | 41 million  |
| 2012 | → | 185 million |
| 2030 | → | 261 million |

# How to Enhance Agriculture Productivity in climate change scenario?

- **Management**

May involve huge cost

- **Genetic gain**

Less cost but needs very strong knowledge base

Major increase in yield came through improved genetics, often termed as '**genetic gain**'

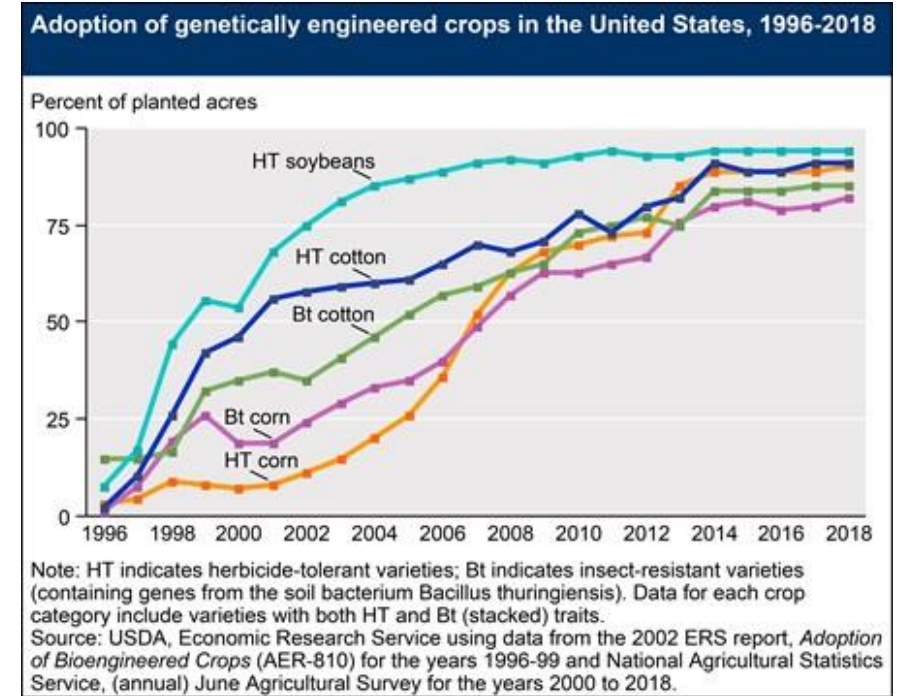
**Historically major increase in wheat, corn, milk, meat and eggs through genetic gain**

The rate of genetic gain is slow;  
requires new technologies

# Challenges in GM crops; Only cotton is commercialized

- Regulatory frame work
  - Trade issues
  - Public acceptance
- Perception from Europe vs North America/South America

Cotton, oilseed crops, sugar and low lignin trees accepted



## The way out for developing world

- Education of policy makers/masses
- Genomics assisted breeding/speed breeding
- New breeding technologies



# New breeding technologies

New breeding technologies (NBTs) include

Genome editing/engineering technologies

- a) zinc finger nucleases
- b) transcriptional activator-like nucleases
- c) clustered regularly interspaced short palindromic repeats (CRISPR)/CRISPR-associated Cas9 systems
- d) Modified CRISPR/Cas9 for nucleotide change without DNA cutting



## Applications in food crops

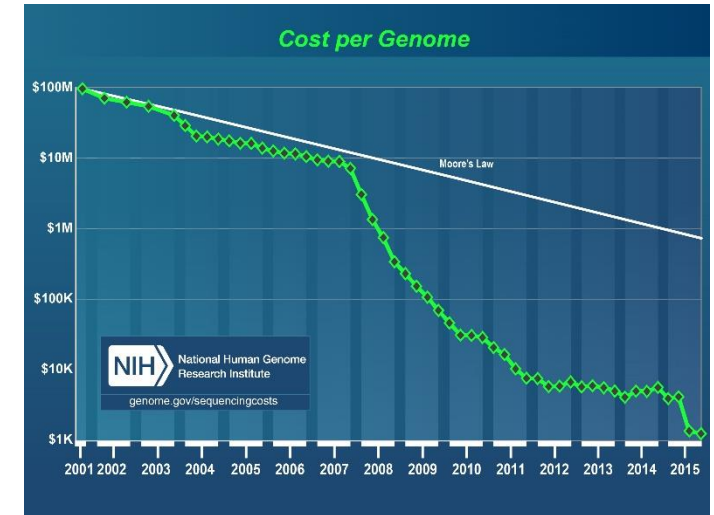
- Rice; yield, better grain, nutritional value, herbicide tolerance
- Potato; virus resistance, sweetening control, stress tolerance
- Wheat; yield, disease resistance, nutritional value
- Cotton; disease resistance, better quality, nutritional value
- Oilseed crops; higher yield, better quality, nutritional value

# Genomics; a silent revolution

Major improvements in sequencing technologies

All major crops/livestock have been sequenced

- DNA based marker technologies
- Genotyping by sequencing (GBS);
- Automated phenotyping
- Bioinformatics; Our ability to handle genomic data



## Applications

- Genome sequencing of important livestock breeds and crops
- Platform for genomic selection in livestock and plants
- Genome sequencing of microbes selected as microbial factories

# **New initiatives food security; applications of biotechnology in livestock and poultry sector**

- Constitute around 60% of agriculture
- 2.3 trillion Dollar Halal market
- 10-15% growth rate
- Potential to control nutritional deficiency and stunting
- Application of genomics can double milk yield in ten years



**Innovative breeding practices for fast-track development of crops and livestock by establishing technological platforms**

**Investment**

**Centralized "Speed Breeding" facilities**

- \* NIBGE, Faisalabad

**Improving selection accuracy during crop and livestock breeding**

- \* Centralized "SNPLine facility"
- \* Multispecies crop and animal SNP chips

**Genome editing capacity**

- \* NIBGE

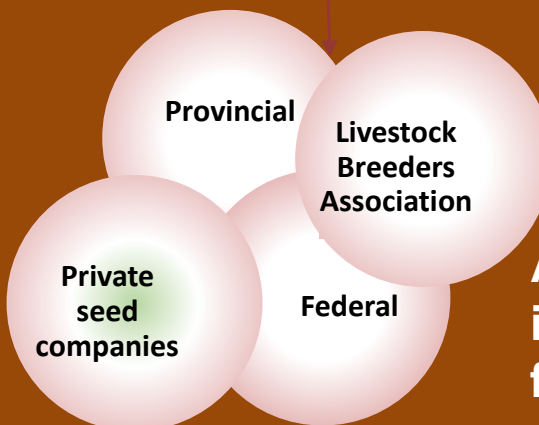
**Platforms**

- \* Reduce the time span of varietal development by 4-6 years
- \* Integrated with provincial and national breeding programs

- \* Improving selection accuracy in crops and livestock
- \* Achieving steady genetic gain by practicing 'Genomic Selection'

- \* Fine-tune the expression of major genes associated with traits of breeding interest

**Benefits**



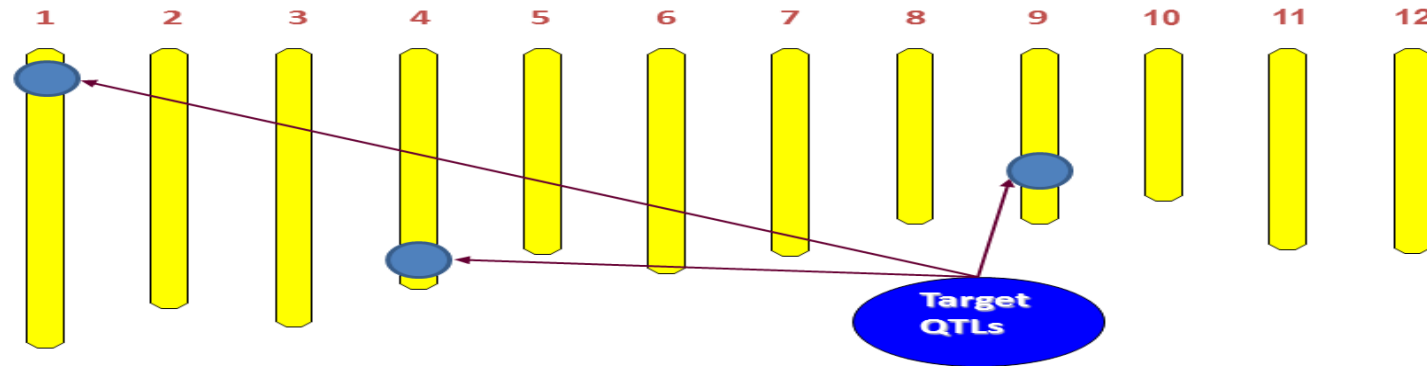
**Stakeholders**

**Approval and provision of improved seeds/livestock to farmers**

Success stories

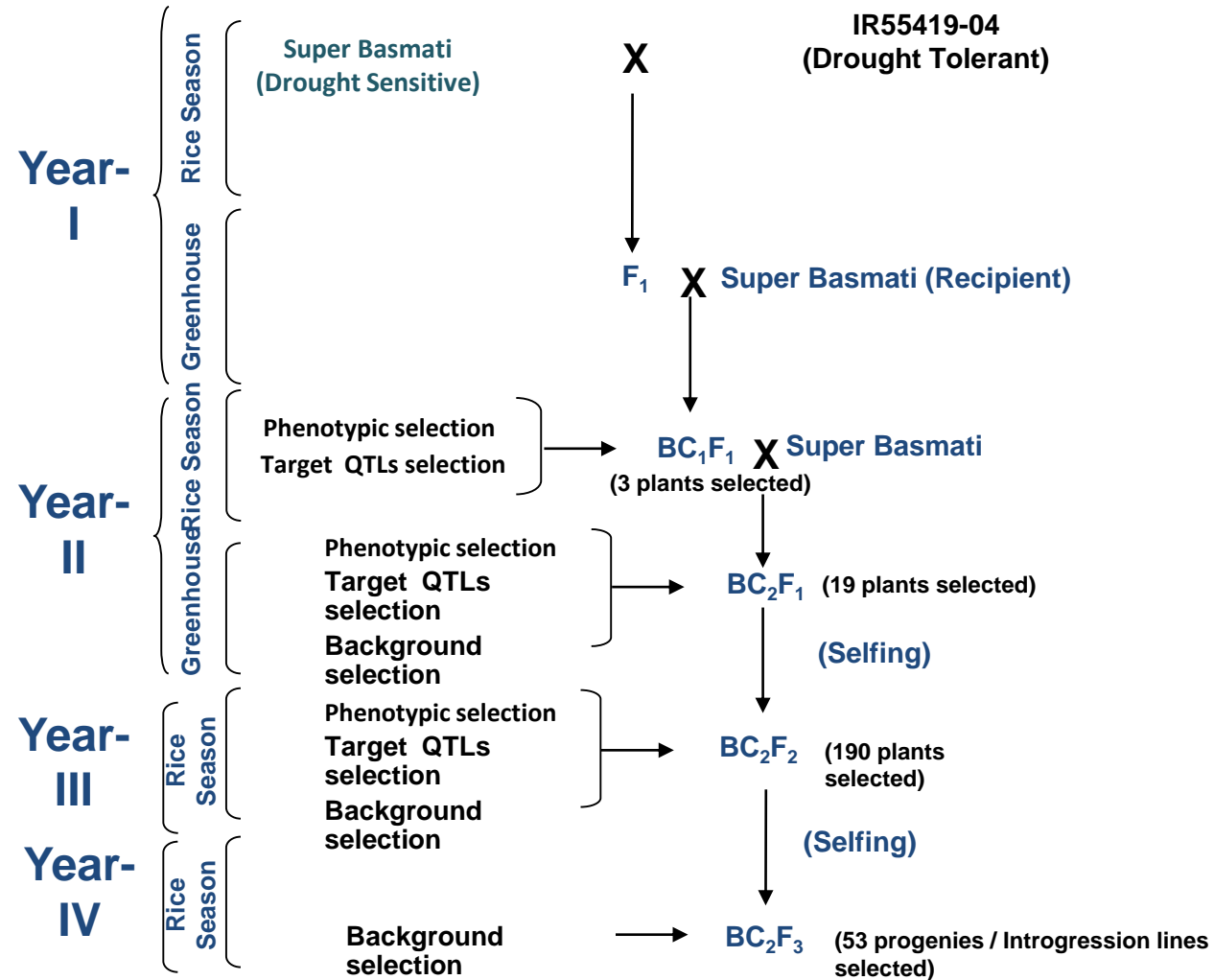
**Introgression of Drought Tolerant QTLs/genes in  
Basmati Rice:  
*A Molecular Breeding Approach***

# Drought tolerant QTLs used for Introgression through Marker Assisted Backcrossing



| Chromosome | Linked / selectable marker(s) | QTL  | Reference   |
|------------|-------------------------------|--|---|
| 1          | RM84<br>RM220                 | Osmotic adjustment   | Robin et al., 2003<br>(IR622/IR600 QTL 2003)  |
| 4          | RM559                         | deep root dry weight;<br>drought tolerance                         | IGCN ZYQ18/JX17 DH QTL 1998;<br>TTU IR58821/IR52561 QTL 2002  |
| 9          | RM201,<br>RM242               | Maximum Root length;<br>Root thickness , Relative<br>water content | IR64/Azucena DH<br>Courtius et al, 2000<br>Price et al, 2002<br>Steele et al., 2006<br>Chaitra et al., 2006 |

# Schematic summary of procedures for introgression of drought tolerant QTLs in Basmati background

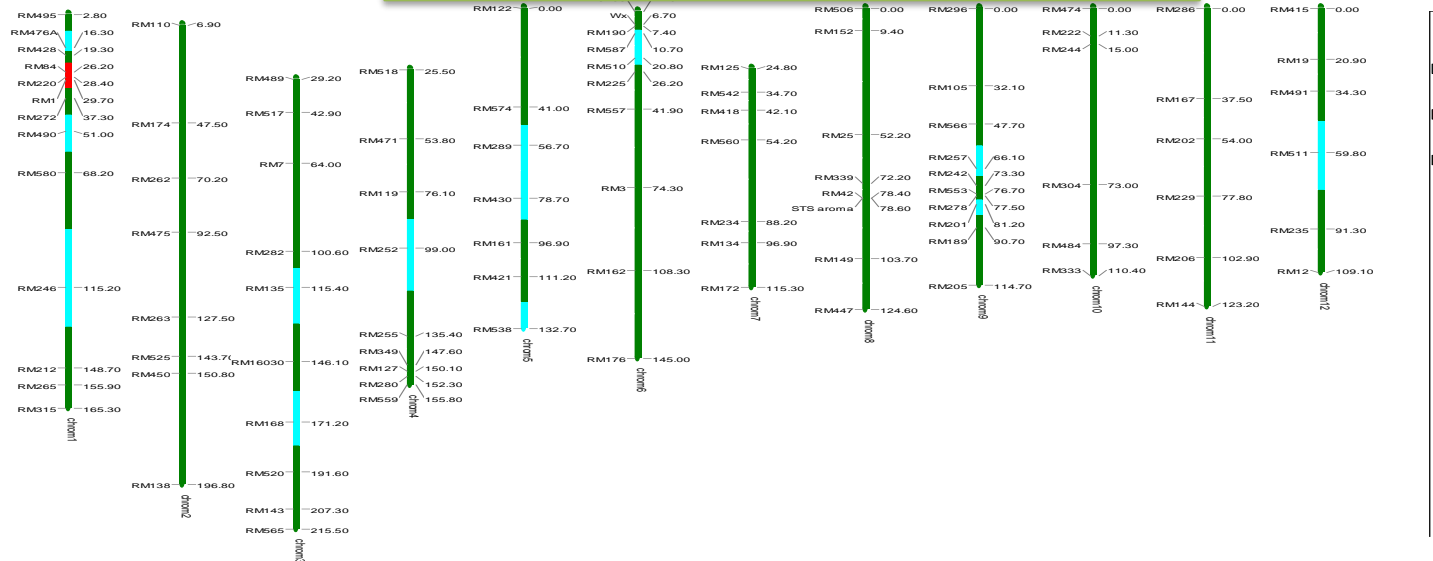


# Field Performance of DT Introgression Line – DTIL60

| Traits                |              | DTIL60<br>(Q1) | Super Basmati |            |
|-----------------------|--------------|----------------|---------------|------------|
| Plant height (cm)     |              | 111.3          | 118.8         |            |
| Maturity Days         |              | 138.3          | 143.9         |            |
| 1000 grain wt (g)     |              | 20.7           | 21.5          |            |
| Paddy Yield           | Well Watered | 3466           | 3735          |            |
|                       | Stress       | 1628*          | 1186          |            |
| % Reduction           |              | 55.1           | 68.2          | IR55419-04 |
| Grain dimensions (mm) | Length       | 7.0            | 7.5           | 6.4        |
|                       | Width        | 1.7            | 1.8           | 2.3        |
|                       | Thickness    | 1.5            | 1.5           | 1.67       |

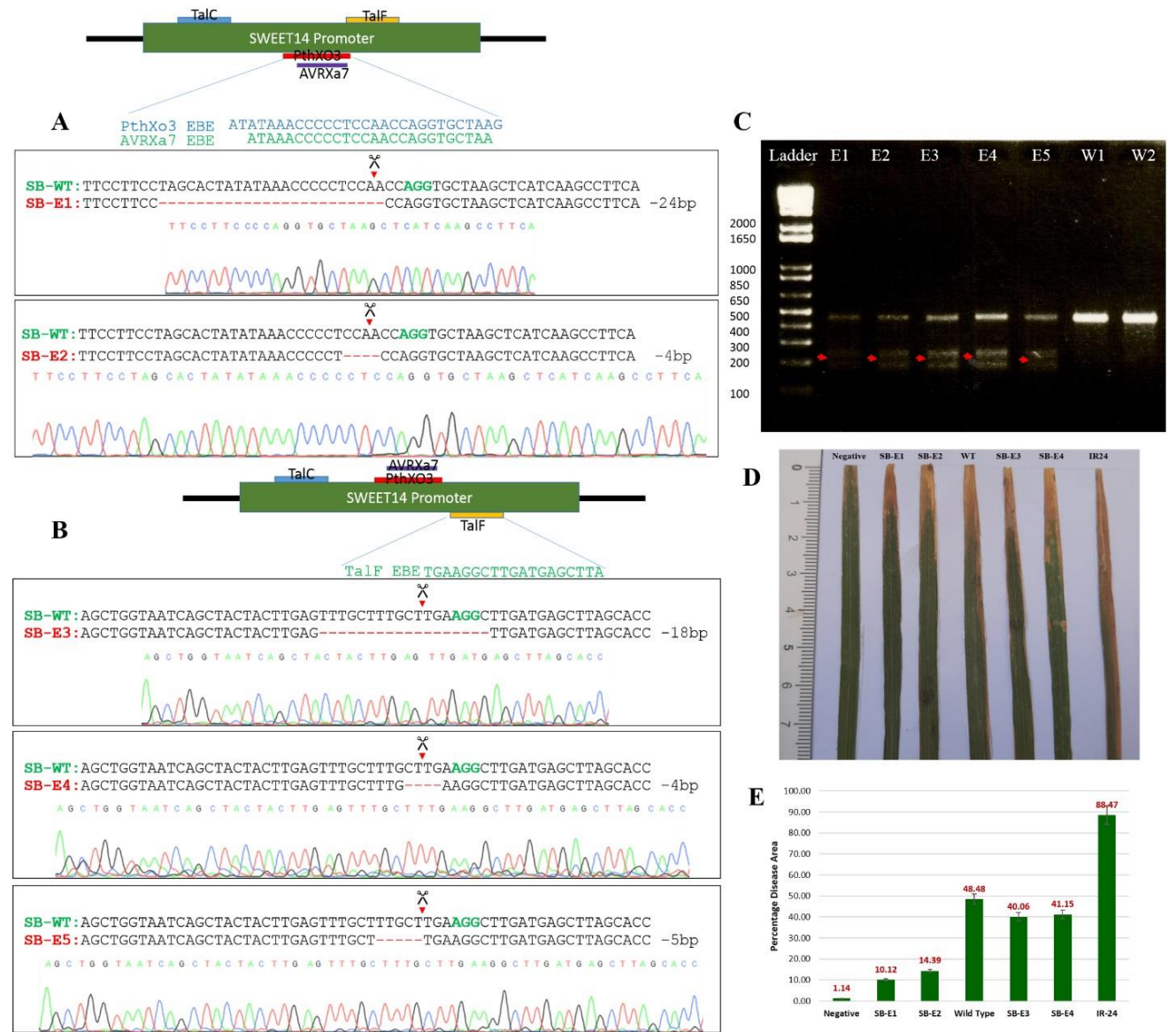


**Genome Recovery = 92.9%**



# Success story; genome editing in rice for bacterial blight resistance in Super basmati

- SWEET14 is a major susceptibility gene because majority of geographically distant Xoo strains target SWEET 14
- The effector binding elements of 4 TALENs are present in the promoter of this gene
- These TALENs are AVRxa7, PthXo3, TalC and Tal 5
- We designed 3 gRNAs targeting these 4 TALENS (AVRxa7 and PthXo3 are overlapping)
- We were successful in getting



# Biofertilizers; reduce chemical fertilizers by 25%

## Next generation Biofertilizers

Value additions, e.g., biopesticides, growth promoting hormones, P- solubilization, insecticide and herbicide degradation, new carrier material

## Way forward

Coating on chemical fertilizers



Field testing of BioPower

## **New technologies for future investment**

- New breeding technologies (genome editing, CRISPR), speed breeding
- Genomics in improvement of crops, livestock and human health
- Minimize use of chemical fertilizers/pesticides
- Applications of automated phenotyping/satellite imaging
- Microbial technologies
- Synthetic biology

## **Collaboration**

- Rothamsted
- John Innes Centre
- NIAB, Cambridge
- University of Bangor
- Pirbright



Thanks