# Utilizing Genomics and Phenomics in CIMMYT wheat breeding

#### **Suchismita Mondal**





#### **Outline**

Genomics & phenomics in breeding pipeline

Key learnings

New directions





#### Collaboration



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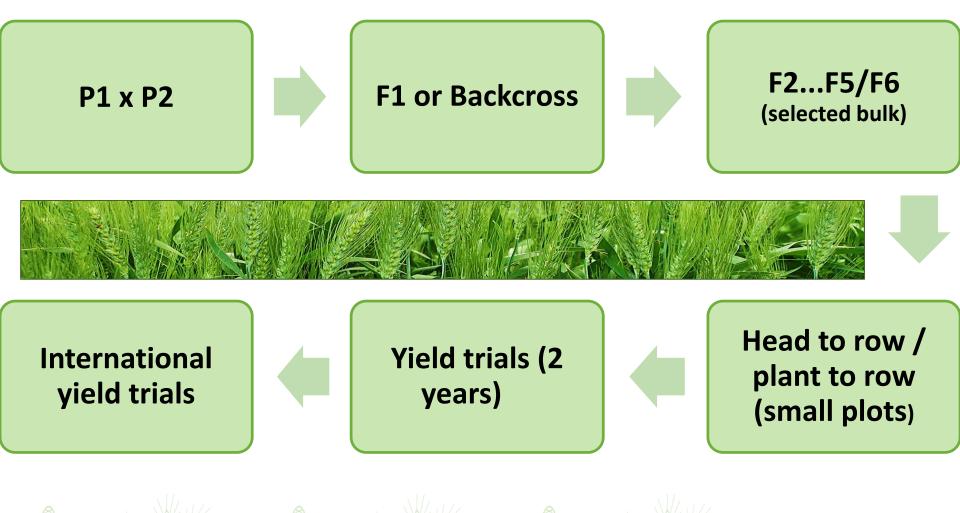






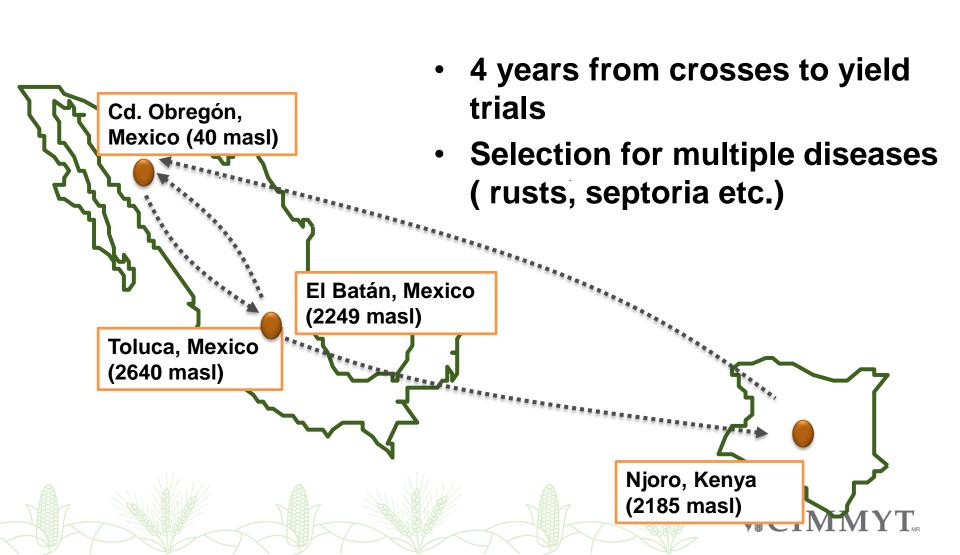
## **Breeding cycle at CIMMYT**

**CIMMYT** 



# Rapid cycling of breeding materials (Field based selection)

Mexico (Cd. Obregon-Toluca/El Batan)- Kenya International Shuttle Breeding



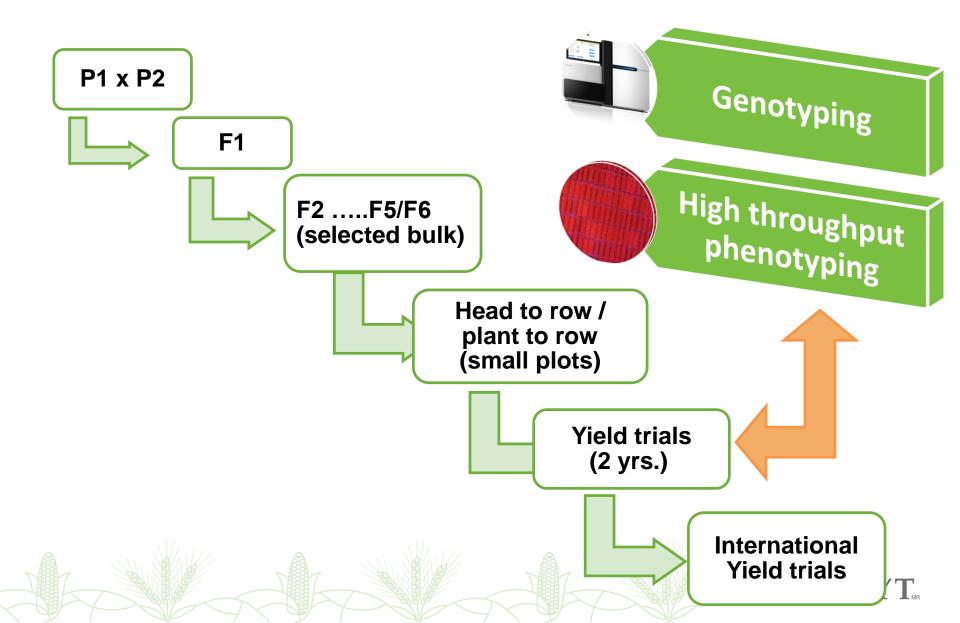
### **Objectives**

#### Improving genetic gains:

- Reducing the breeding time
- Improving selecting efficiency
- Phenotyping under diverse environment



## Integrating technologies



#### **Strategy**

1st year yield trial(~10,000 lines)

2<sup>nd</sup> year Multi-environment yield trial (~1000 lines)

**Irrigated** 



Irrigated (beds & flats)

Drought (severe & moderate)

Heat



1st year yield trial(~10,000 lines)

2<sup>nd</sup> year Multienvironment yield trial (~1000 Lines)

#### **Pedigree Information**

- GBS markers (KSU)
- 10,000 lines each year for 4 years

Genotyping



- Ariel phenotyping
- Thermal and hyperspectral camera

High throughput phenotyping





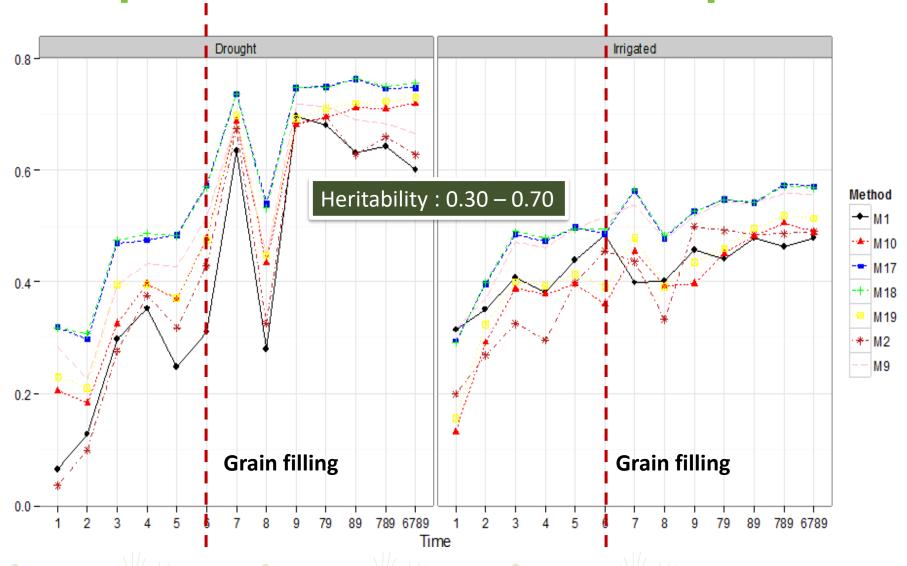
## **Phenotyping**

- Key questions
  - Association of traits measured through high throughput phenotyping (HTP)
    - All bands vs indices (R-NDVI, G-NDVI, Canopy temperatures)
  - Timing of data collection





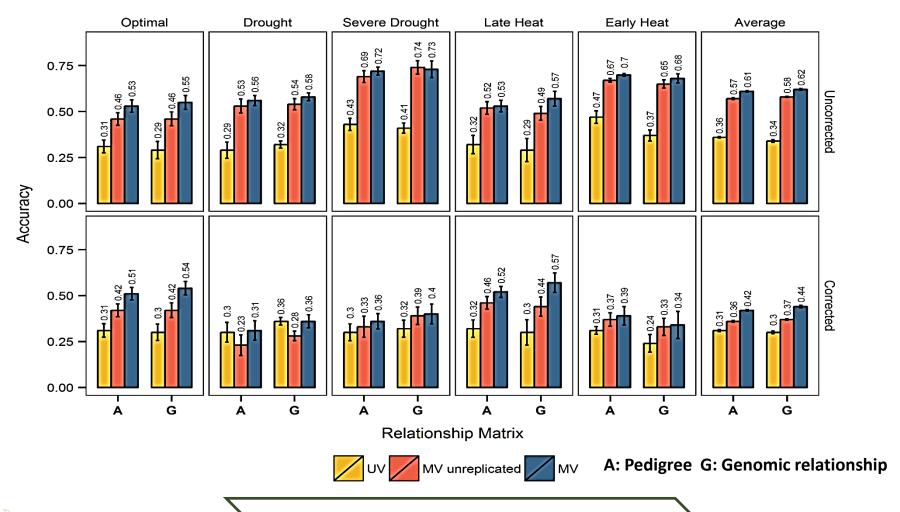
#### Comparison of models & time points



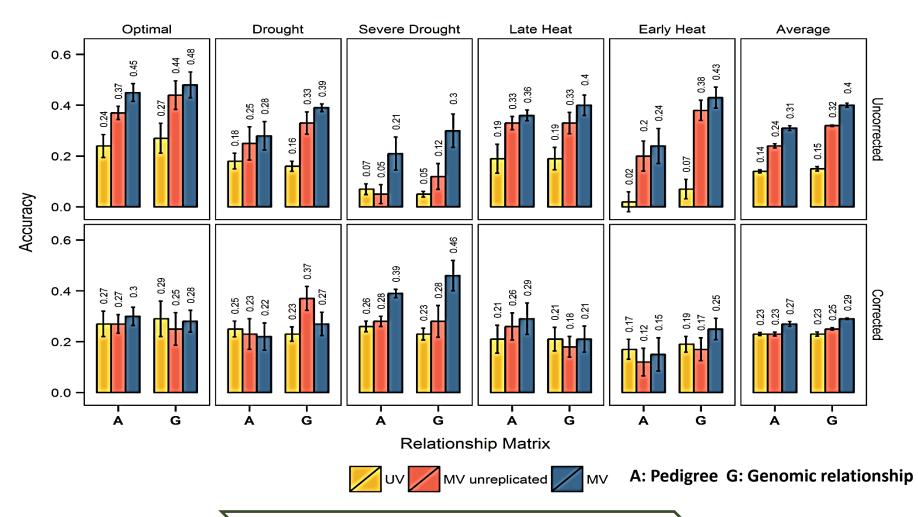
2<sup>nd</sup> year Multi-environment yield trial

CIMMYT

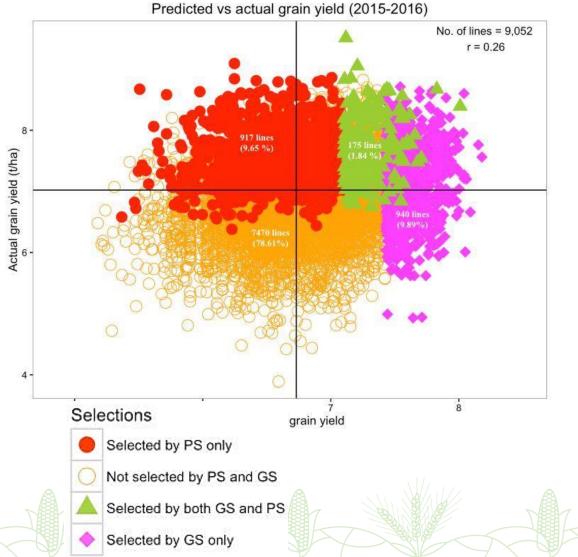
# Within environment predictions using multivariate (HTP indices) model



# Across environment using multivariate (HTP indices) model



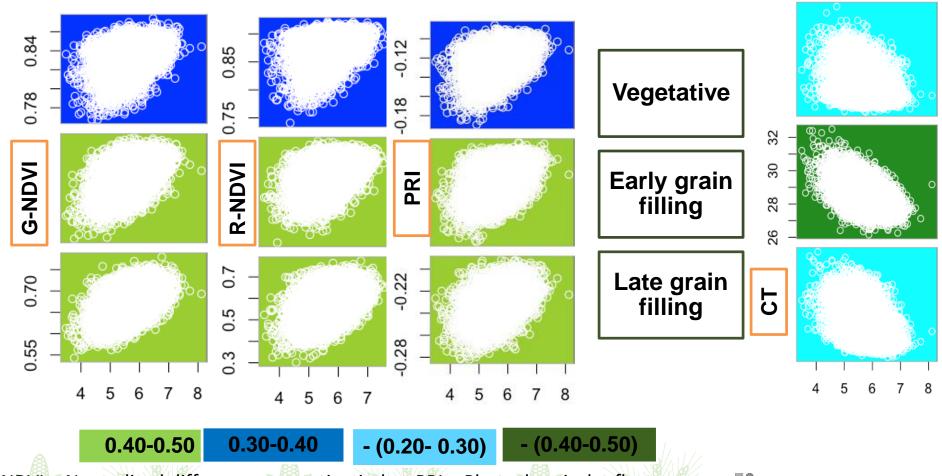
#### 1st year yield trial(~10,000 lines)



- Grain Yield : Predicted vs Balance
- Rejected:
  - Sib with better performance
  - Seed color, height
- Evaluations underway in 2016/ 2017



## Grain yield association with...



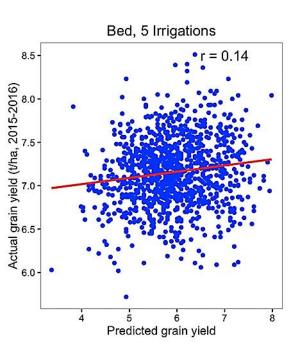
NDVI – Normalized difference vegetation index, PRI – Photochemical reflectance index, CT – canopy temperatures

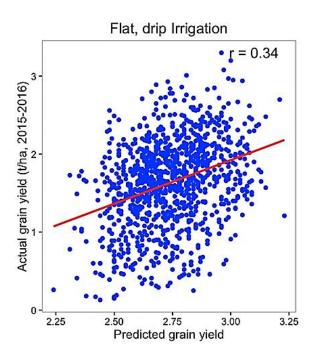


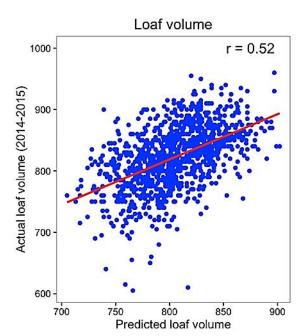
#### Forward predictions vs Balance

1st year yield trial(2014)

2<sup>nd</sup> year Multi-environment yield trial (2015)

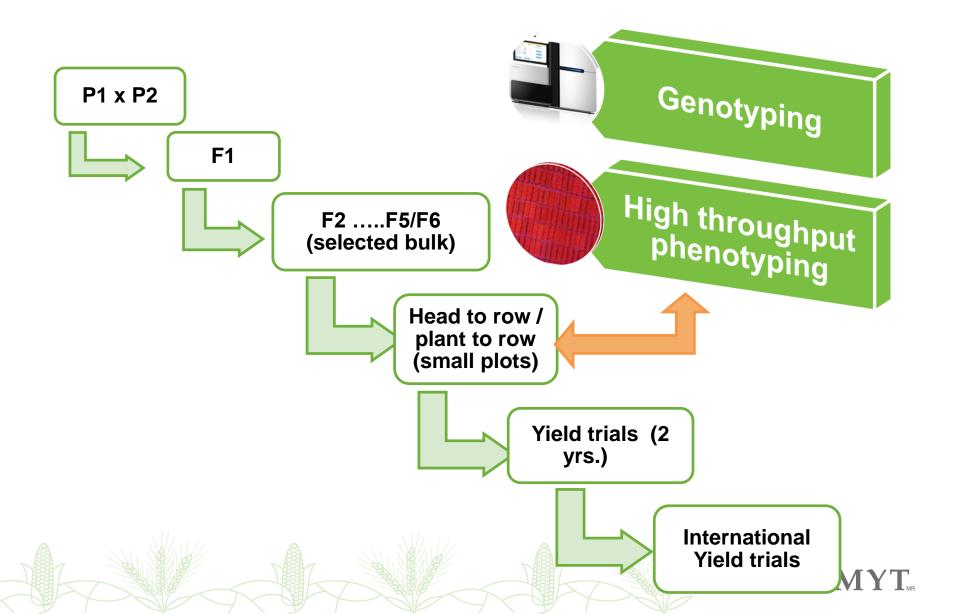








#### **New Directions**



#### Head rows/plant to rows



- 50,000-70,000 head rows of 1m for visual selection
- GS or HTP could aid the selection process
- Data collection on way





#### **Predictions across locations**



- Selected Yield trials evaluated
  - Bangladesh (Jamalpur)
  - India (Ludhiana, Jabalpur, Pusa)
  - Faisalabad
- Phenotyping and performance prediction ongoing

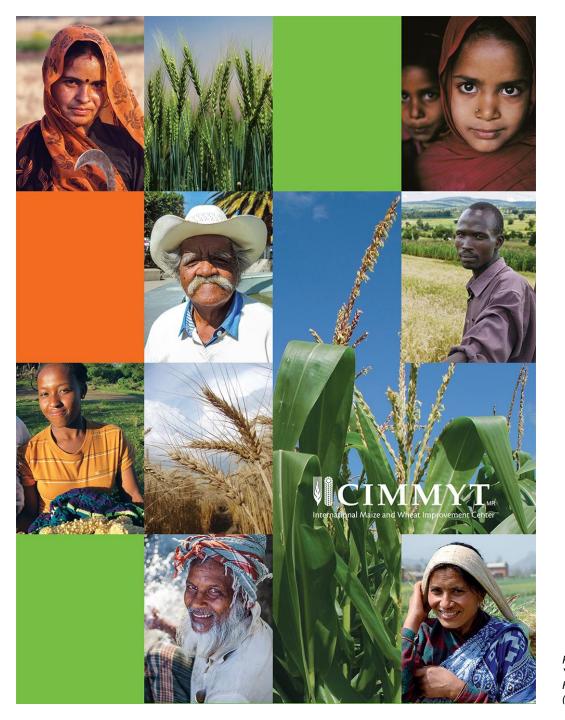


#### **Summary**

- Potential of genomics and phenomics in improving selection efficiencies
- Further work in including GxE to improve prediction
- GWAS is also being used to standardize GS models (Sehgal et al. 2016, Poster No. 179)







# Thank you for your interest!

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