



# Breeding informatics

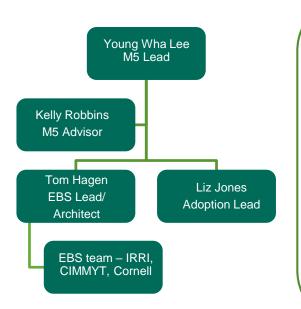
## Young Wha Lee

Breeding informatics lead



## Module 5 Breeding Informatics

M5 focuses on activities that increase the incorporation of modern informatics capabilities into best practice breeding processes.



#### **Deliver software**

Delivery of Enterprise Breeding System

### Deliver integrated and centralized analytic capability

- Develop a generalized analytics framework for EBS
- Drive towards a centralized analytics resource for 1CG

#### Support breeding programs in adoption, including but not limited to software

- Adoption of EBS
- Adoption of/fluency with BMS
- Transition of BMS users to EBS
- Digitization of data collection

Coordinate long term strategy on data management systems for public breeding, with the EBS leadership team, external stakeholders, and funders



## Please connect with us!

EBS website: <a href="https://ebs.excellenceinbreeding.org/">https://ebs.excellenceinbreeding.org/</a>

EBS Leadership Team: Jan DaBaene (chair), Young Wha Lee, Tom Hagen, Liz Jones, Eng Hwa Ng, Eduardo

\*the EBS LT is responsible for internal strategic governance of Module 5 activities

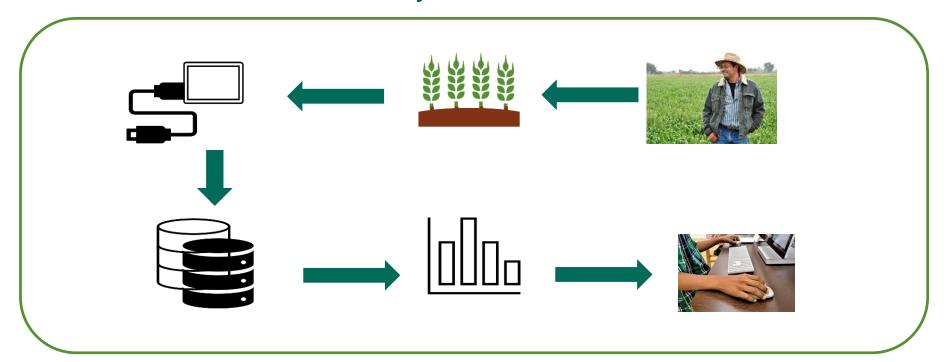




# **Enterprise Breeding System**



# Modern breeding requires data management systems





## Key Features of the Enterprise Breeding System

- □ Software-as-a-Service with centralized development and deployment
- Supports a diverse array of breeding programs under OneCGIAR
- □ Supports core breeding activities
- Includes a genotype database
- ☐ Includes an analytical framework capable of advanced biometrics







## **EBS Version Roadmap**

Goal: Enable a complete execution of a breeding cycle from Version 4

Version	WF Name	
2	Breeding Trials	
2	Nurseries	Core
2	Germplasm Management	breeding Integration of
2	Phenotyping	operations — genotype data
2, 3	Field Operations	management,
3, 4	Analytical Service Submission	sample tracking, and analytics
3, 4	Breeding Analytics	

Projected delivery dates

**V2: Nov 2020** 

V3: June 2021

V4: Dec 2021



## EBS 2.0 components and functionalities

Name	Abbreviation	Supported activities
Breeding4Results	B4R	Core Breeding
Analytics Framework (*New*)	AF	Trial Design
KDXplore (*New*)	KDX	Field Mapping (plot layout)
Service Gateway (*New*)	SG	Login and System Connectivity

- Create trials and nurseries
- Generate alpha lattice, row-column, RCBD, and augmented RCB designs
- Make crosses
- Create and capture plot layouts
- Generate electronic fieldbooks for data capture apps
- Upload phenotypic data and perform basic phenotypic data QC
- Search for germplasm, seeds, traits, and experiments



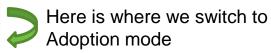


# **EBS Adoption**



# Understanding stages of testing: example of CIMMYT maize/wheat and IITA maize

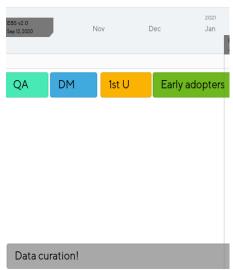
Environment	Testing stage/ process	Version EBS	Who is organizing?	Who is testing/using?
Develop	Develop	2.0, 2.1, 3.0	EBS	EBS developers
QA & UAT &	QA & UAT	2.0, 2.1, 3.0	EBS	EBS requirements analysts and QA/QC team
CIMMYT* Test	Test	2.0, 2.1, 3.0	CIMMYT DM team	Data managers
CIMMYT Staging	Product owner	2.0, 2.1, 3.0	CIMMYT DM team	Product Owners and Data Managers
CIMMYT Staging	Early adopter	2.0, 2.1, 3.0	CIMMYT DM team	Early adopters
CIMMYT Production	Production	3.0	CIMMYT DM team	All





## EBS 2.0 Rollout: Where we are now

## **V2.0**



A data manager embedded in the Centers is critical to coordinate and facilitate adoption activities

Adoption activities include:

- EBS training
- ☐ Feedback to development team
- System adoption KPIs
- Data curation
- Digitization training



# A broader focus: Module 5 supports adoption of breeding data management systems in general

- □ Not just EBS we have ongoing collaborations to support BMS adoption
- We believe the adoption of a data management system (whether BMS, Breedbase, or EBS) is more urgent
- With IBP, we will build tools for BMS-EBS data migration to facilitate eventual convergence into EBS





## **Breeding Informatics Network**



# Biometric/bioinformatic resources are Center specific and unevenly distributed across the CGIAR

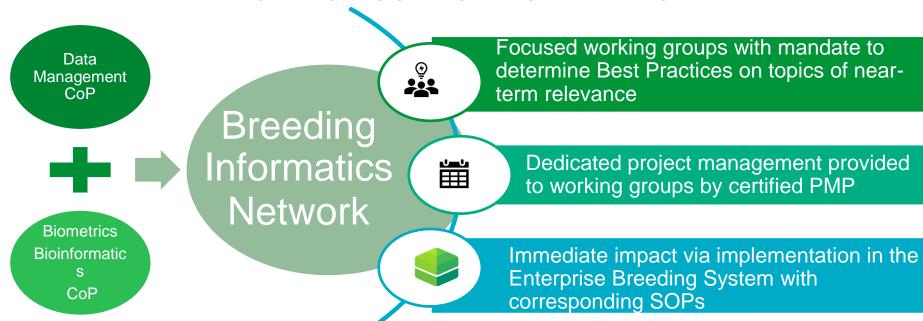
	CIMMYT	IRRI	IITA	ICRISAT	CIP	CIAT	ICARDA	AfricaRice
biometrics			1 FTE		1 FTE			
Bio-informatics	Molecular bree but no bioinform		1.5 FTE					

EiB's support of data analytics efforts up till 2020 has been mainly in the form of:

- supporting Communities of Practice in data curation and biometrics/bioinformatics
- organizing workshops & trainings
- funding sabbaticals
- providing best practices consulting & manuals



# With the existing CoPs as a foundation, we will initiate a cross-Center Breeding Informatics Network in 2021



Excellence in Breeding Platform

# We have begun the organizational and alignment process for this cross-Center effort

### M5 resources

- ☐ Funding partial salaries of Biometric leads, so they can participate and lead in the Breeding Informatics Network
- Building up biometric capacity in IITA
- ☐ Hiring of EiB Bioinformatics Lead to bridge M5, M3, and coordinate CG bioinformaticians
- Access to expert contractors via CIMMYT for one-off projects
- □ Project Manager support

### **Watchouts**

- □ We are building the plane as we fly it – breeding modernization within programs, EBS development and adoption, and operational capacity building are all being pushed simultaneously
- ☐ Coordination across centers is hard
- ☐ Bandwidth can we get people to fully participate in the working groups?

## **Mitigations**

- ☐ Your buy-in and support
- □ Develop, communicate, and stick to a process for determining, approving, and modifying best practices for implementation – enable full transparency
- ☐ Project manage for defined time limited outcomes
- Make sure focus is on near term needs that are 100% aligned with implementation of existing continuous improvement plans

