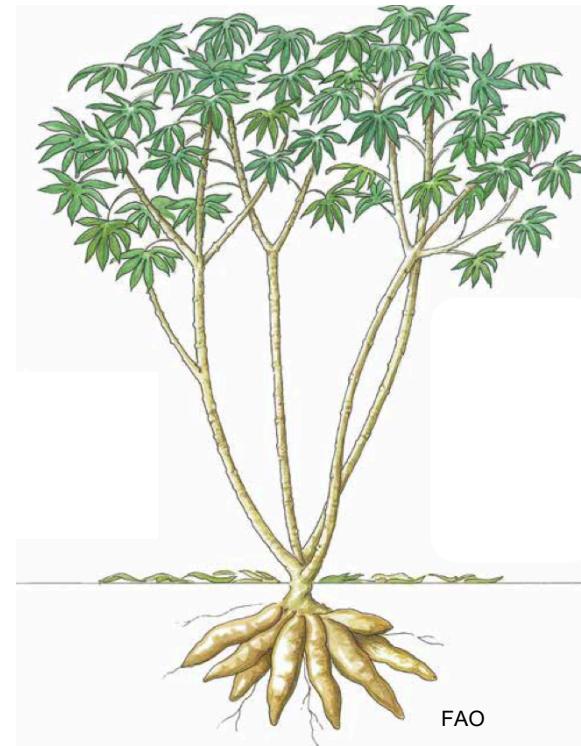


# **Development and integration of handheld NIRS-based phenotyping for cassava breeding**

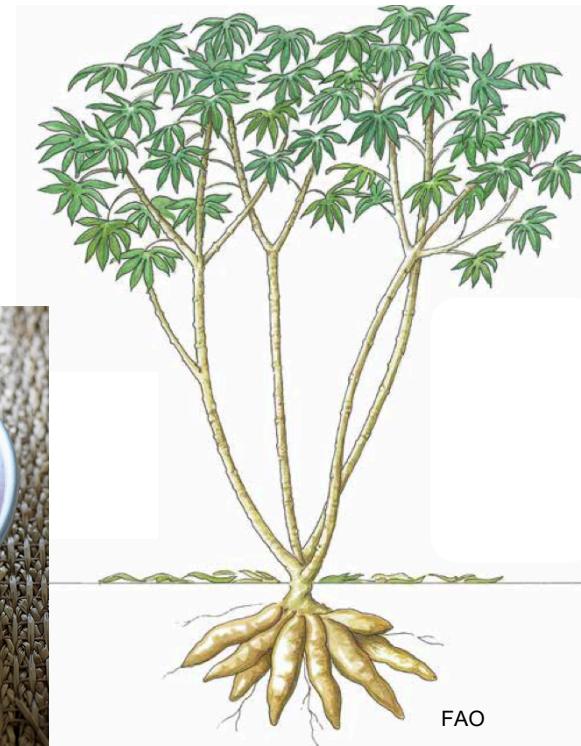
*Jenna Hershberger*

# Cassava: plant of many uses



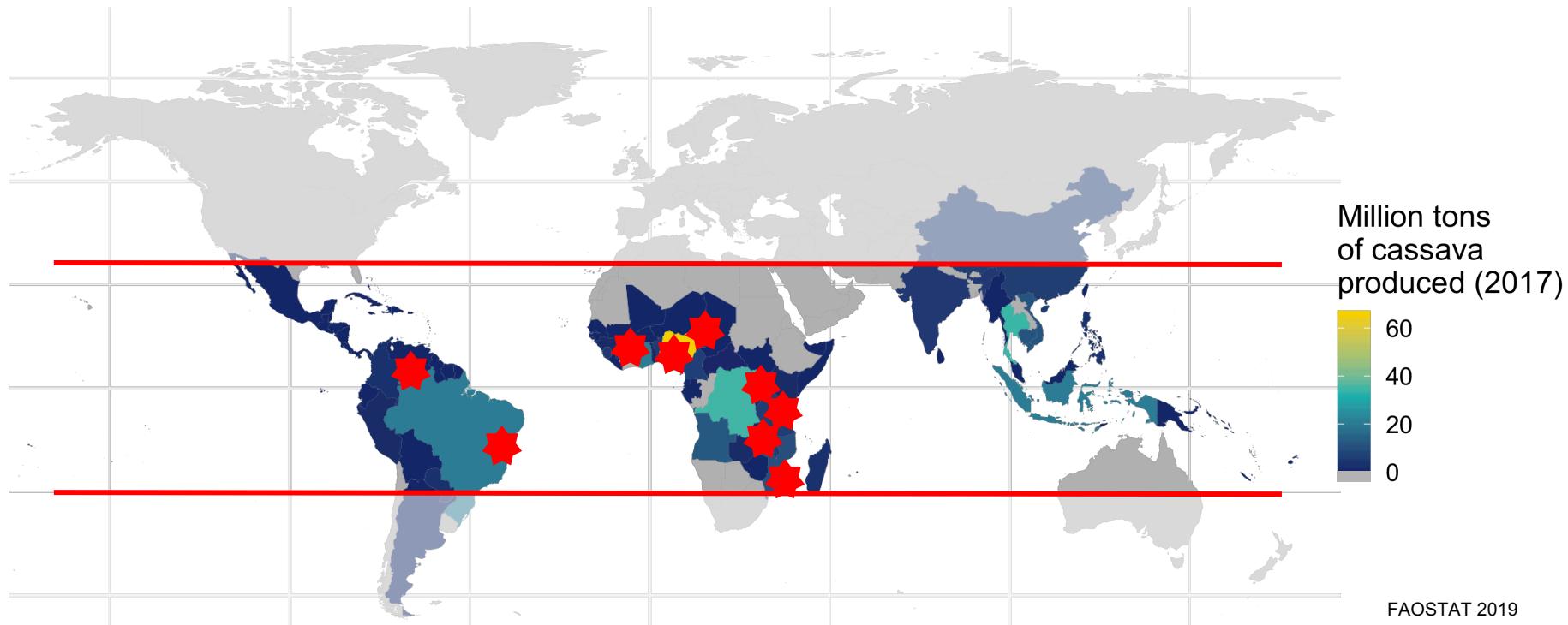
# Cassava: a tropical staple

- Roots consumed by over 800 million people
- Drought-tolerant subsistence crop



FAO

# Cassava: a tropical staple



# Cassava breeding targets

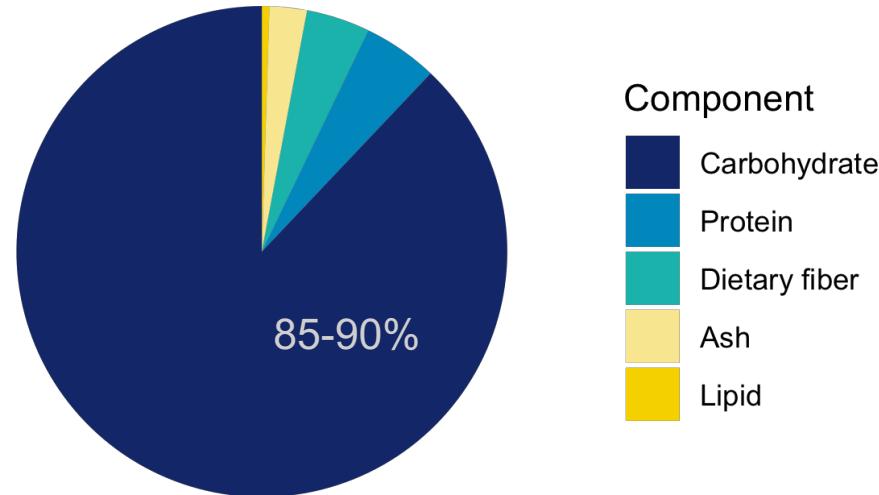
- Pest and disease resistance
- Yield and performance
- Quality traits



# Root dry matter content (DMC)

- $\frac{\text{dry weight}}{\text{fresh weight}} \times 100\%$
- Major component of dry yield and food quality
- Typically ~20-45%
- Not uniform within the root

DMC composition

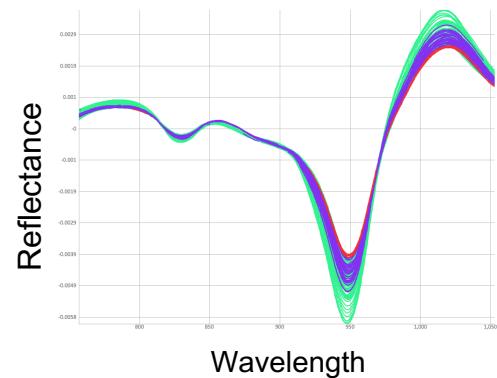
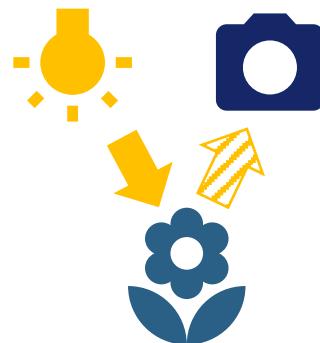


# DMC is time-consuming and laborious to measure



# Near-infrared spectroscopy (NIRS)

- Reflectance from near-infrared region of the electromagnetic spectrum
- Complex signal from biophysical and biochemical properties of sample
- Multivariate statistical techniques required for calibration
- Allows for rapid, non-destructive analyses



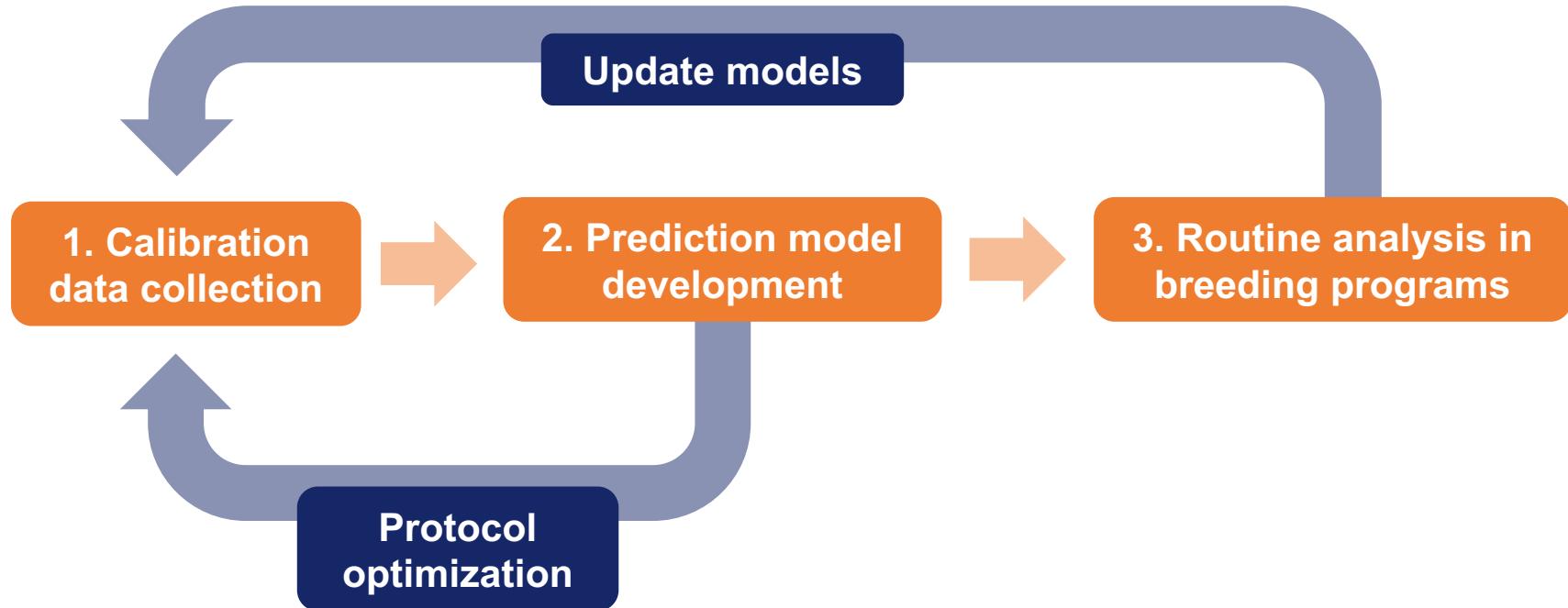
# NIRS for cassava phenotyping

- Highly predictive of dry matter in cassava roots
- Validated spectrometers are prohibitively expensive

Foss 6500	
Spectral range	400-2500nm
Predictive ability	> 0.94



# NIRS validation workflow



# 1. Calibration data collection

**IITA** (Nigeria)



**NaCRRI** (Uganda)



**Embrapa** (Brazil)



# 1. Calibration data collection

Sample three roots/plot



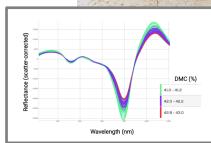
Slice root in three regions



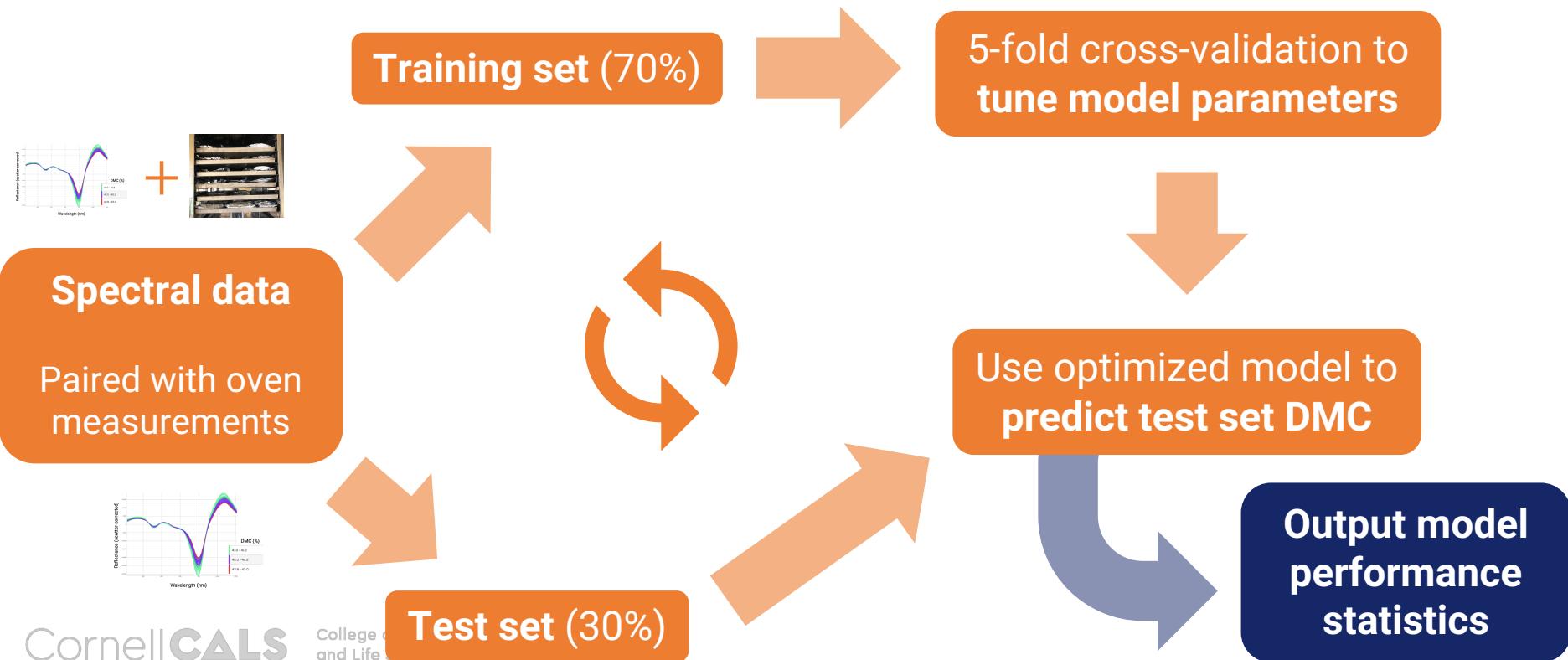
Scan the cut surface from each region



Measure oven DMC with a representative sample from the entire root



## 2. Prediction model development



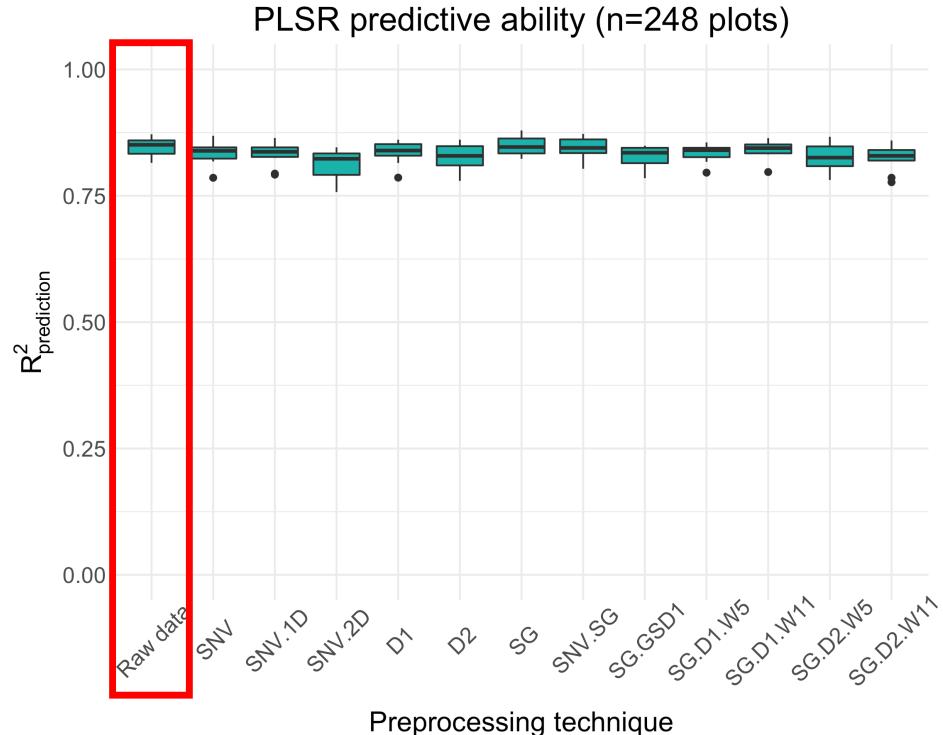
# Optimization: PLSR outperforms other model types

Model type
Partial Least Squares Regression (PLSR)
Support Vector Machine (SVM)
Random Forest (RF)

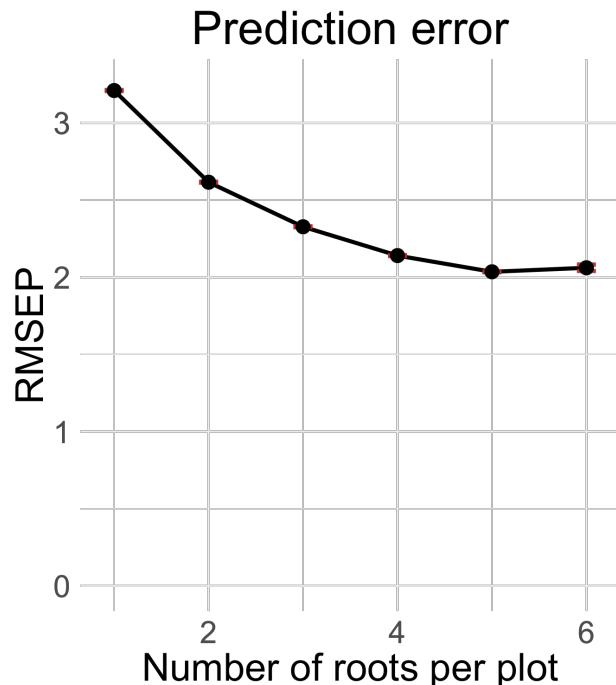
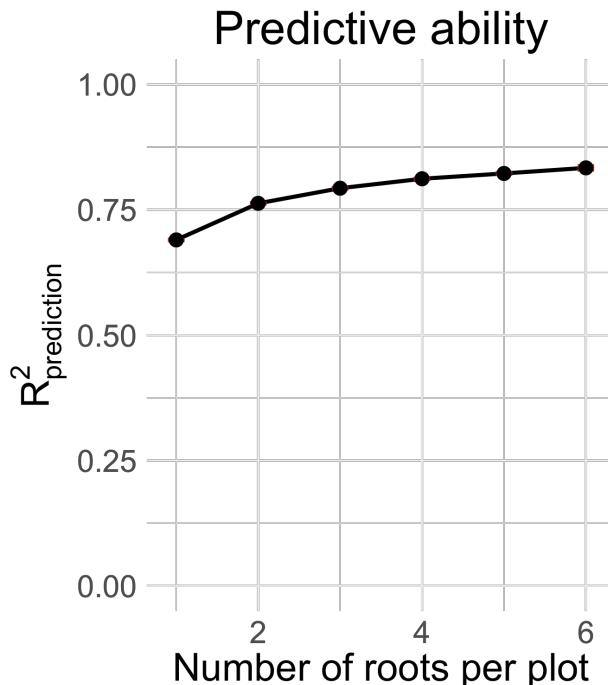
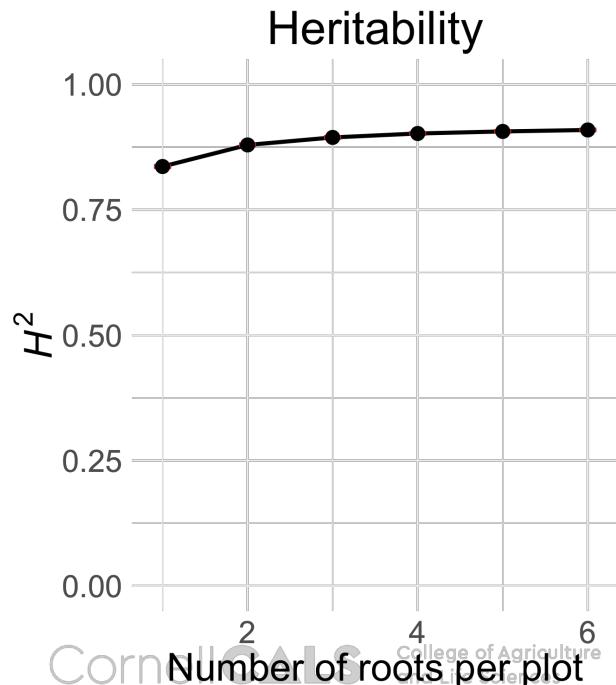
- Models developed and evaluated using IITA dataset
- Ten iterations of model development pipeline

# Optimization: No significant change with preprocessing

- Scatter correction
  - Standard normal variate (SNV)
- Derivatives
  - First and second derivatives (1D, 2D)
  - Window size 11 unless noted by “W”
  - Gap segment derivative (GSD)
  - Savitzky-Golay (SG)



# Optimization: Diminishing returns after three roots





## 1. Calibration data collection

## 2. Prediction model development

## 3. Routine analysis in breeding programs



### Protocol optimization

- PLSR
- No preprocessing
- 3 roots per plot

Spectral range

Predictive ability

Price per unit

SCiO

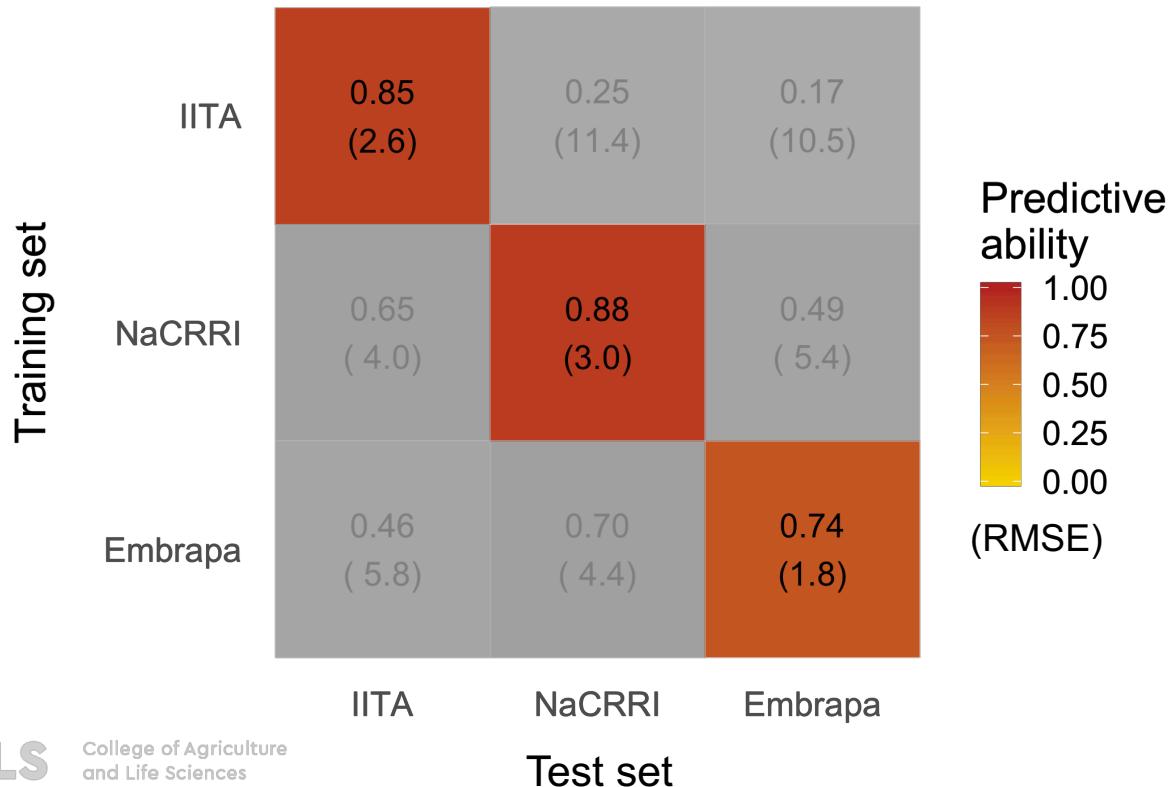
740-1070nm

??

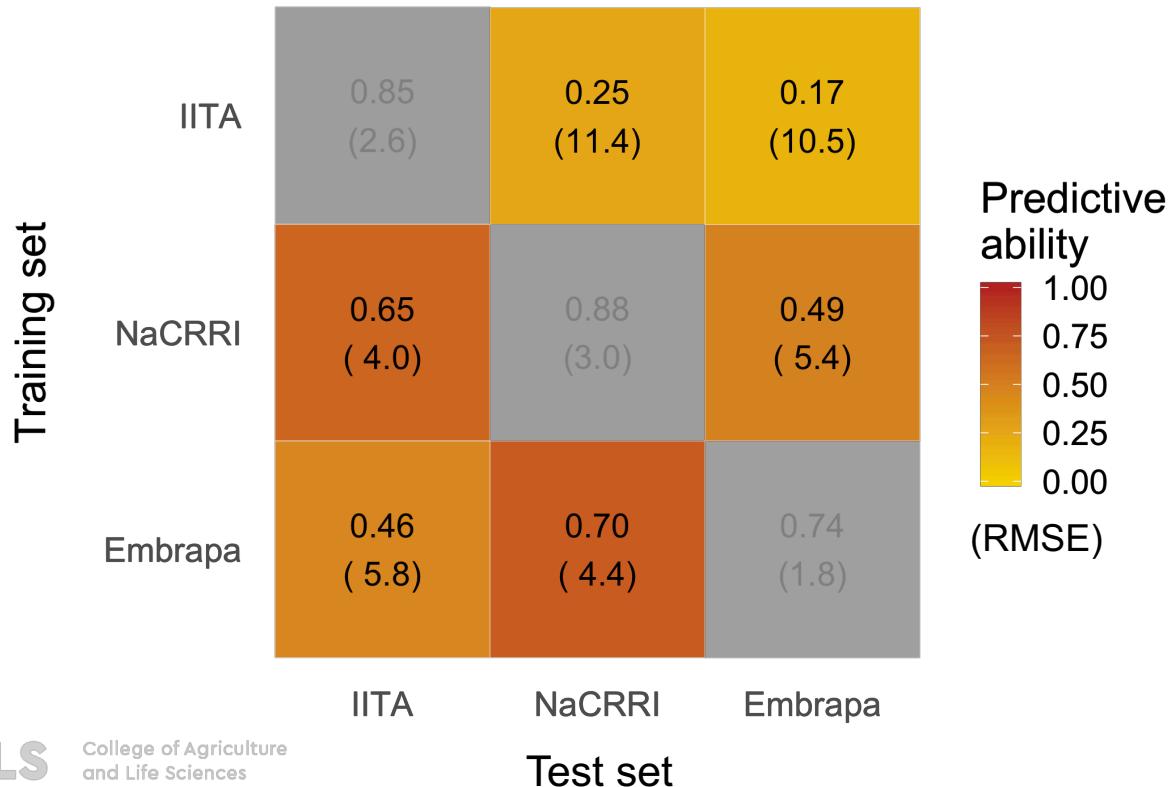


\$299

# SCiO is highly predictive of DMC within breeding programs

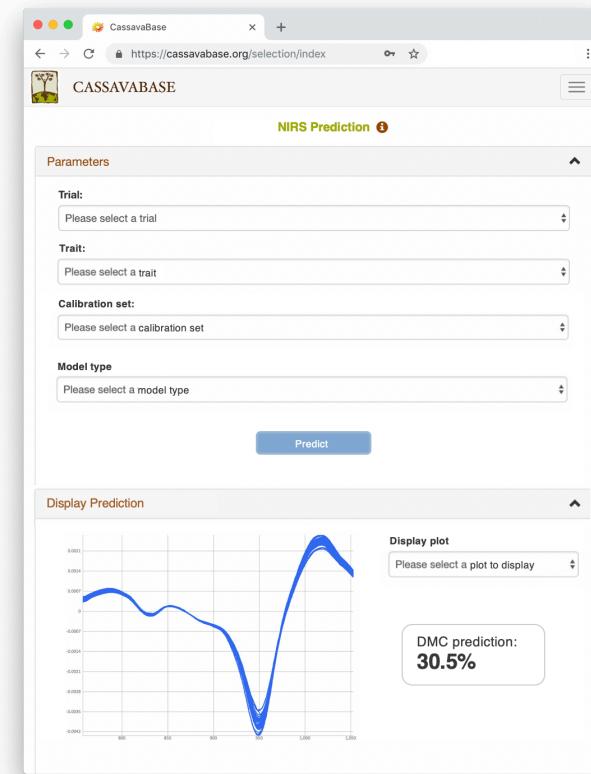


# SCiO is moderately predictive of DMC across breeding programs



# 3. Routine use in breeding programs

- Integrate NIRS data and models into Cassavabase
- Implement model update strategy



# Acknowledgements

## Gore Lab at Cornell University

## NextGen Cassava

- **IITA** – Ismail Rabbi and team
- **NaCRRI** – Robert Kawuki and team
- **Embrapa** – Eder de Oliveira and team
- **Cassavabase** – Lukas Mueller and team

## PhenoApps team

- Kansas State University
- Makerere University

## Committee Members

- Michael Gore
- Rebecca Nelson
- Tim Setter

## Funding

- NSF BREAD IOS-1543958

College of Agriculture  
and Life Sciences

