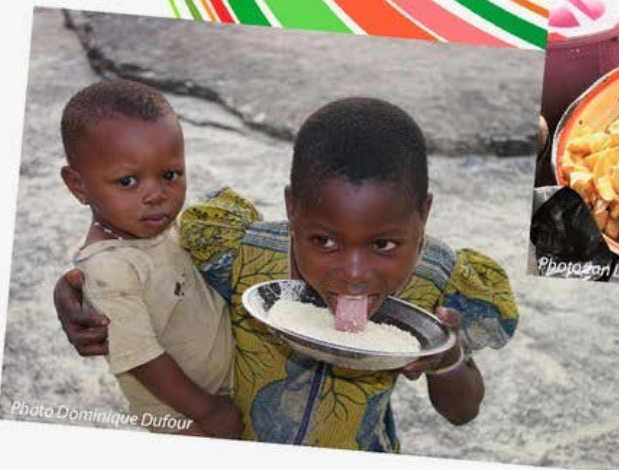


# RTB foods



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# Hyperspectral imaging in Breeding Program

## *Characterisation of cooking ability of boiled cassava*

24ème Rencontres HélioSPIR  
13/06/2023

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### Contributors

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CIAT : Luis Fernando Delgado, Maria Alejandra Ospina,  
Jhon Larry Moreno, Jorge Luna, Luis Londoño.



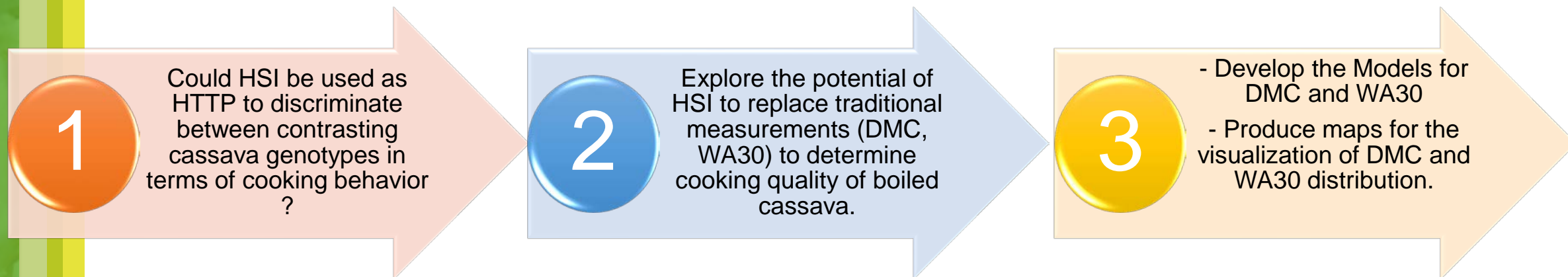
# Outline

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- Introduction
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  - What hyperspectral imaging provides ?
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  - Protocol of measurements.
  - Hyperspectral spectral images processing.
- Results
  - Prediction of DMC in fresh cassava.
  - Prediction of WA30 in boiled cassava.
- Conclusions and perspectives

# Challenge and objectives

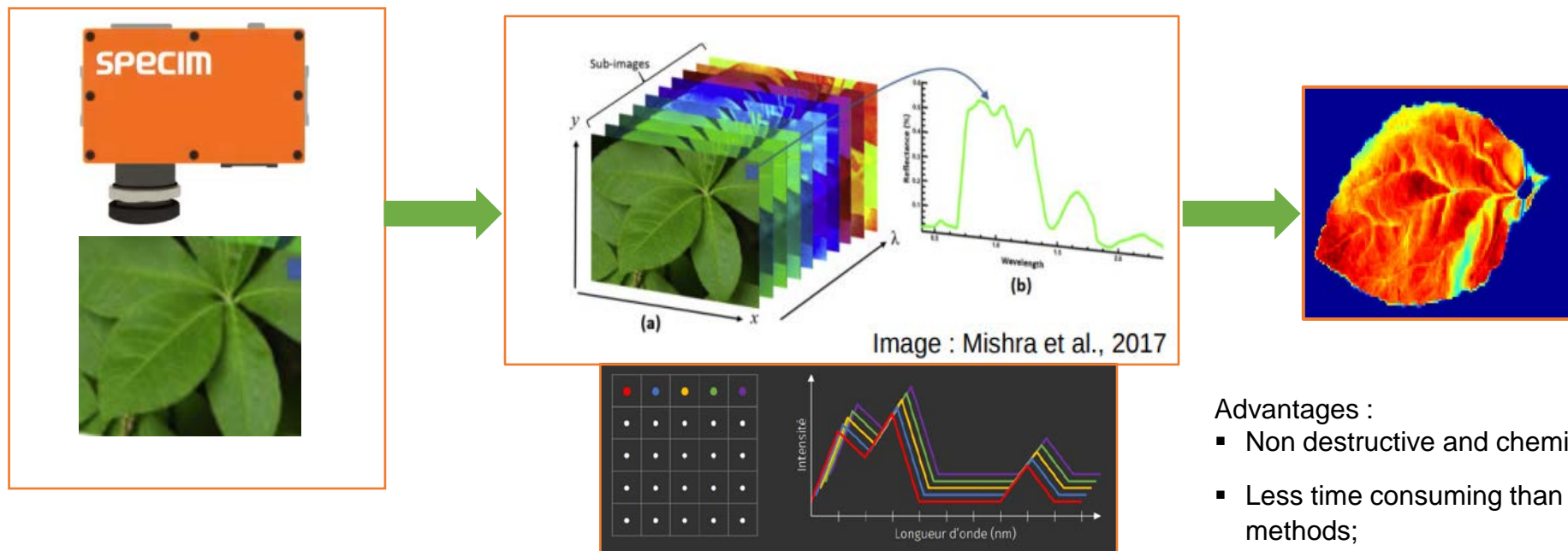
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# What hyperspectral imaging provides ?

- Hyperspectral imaging combines spectroscopic and imaging techniques to enable direct identification of different components and their spatial distribution within the sample.

- Each plan is one image acquired for one wavelength.
- Each pixel in hyperspectral image represents one spectrum.



## Advantages :

- Non destructive and chemical free;
- Less time consuming than conventional methods;
- Spatial visualization of different biochemical constituents of the sample.

# Plant material and Samples preparation



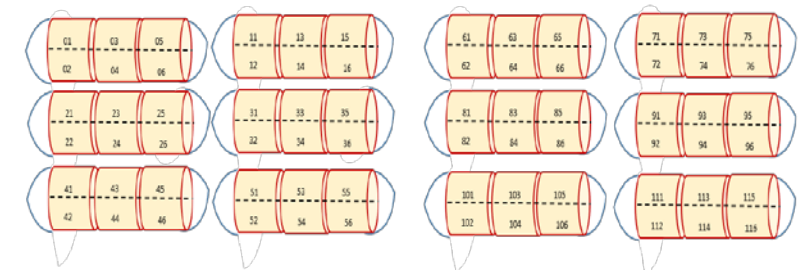
Cassava roots from 31 genotypes (RTBfoods progenitors) harvested in March 2022 at CIAT (Colombia)

12 roots selected per genotype, washed and peeled.

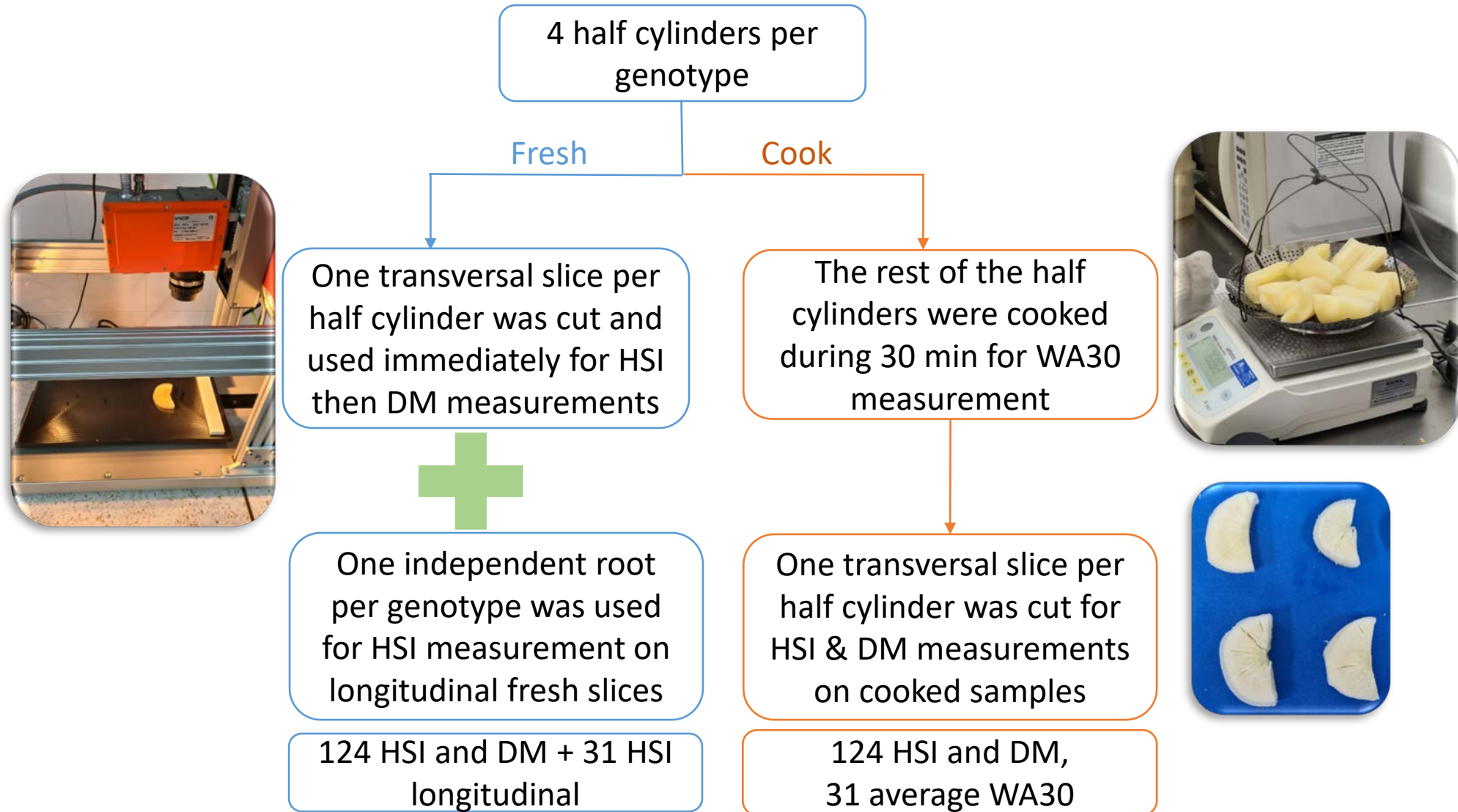


Each root was divided into 6 half-cylinders,  $6 \times 12 = 72$  half-cylinders per genotype

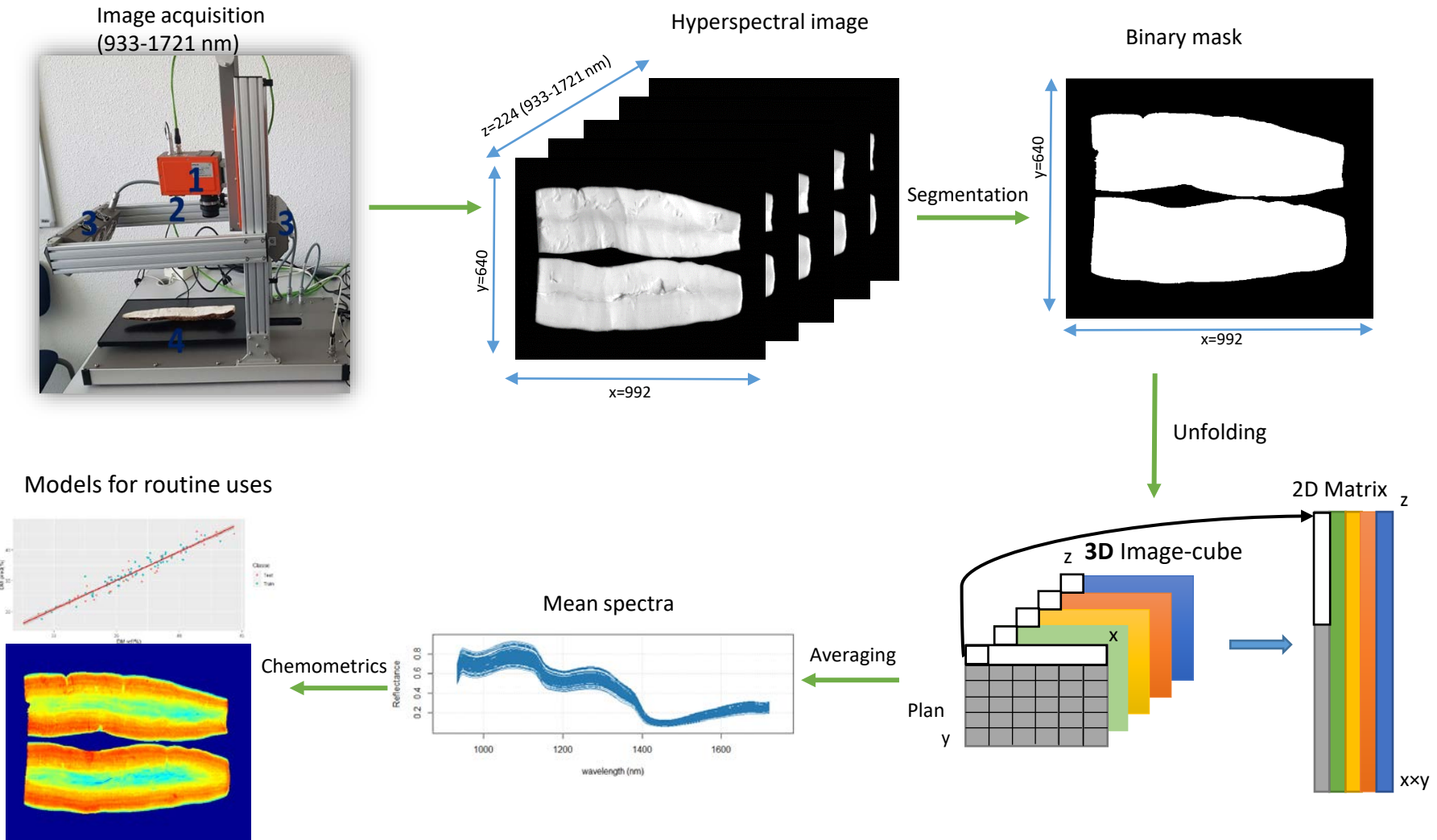
central semi-cylinders number 03,43,103, 114 were selected for HSI, DM and WA30



# Protocol of measurements



# Hyperspectral images processing

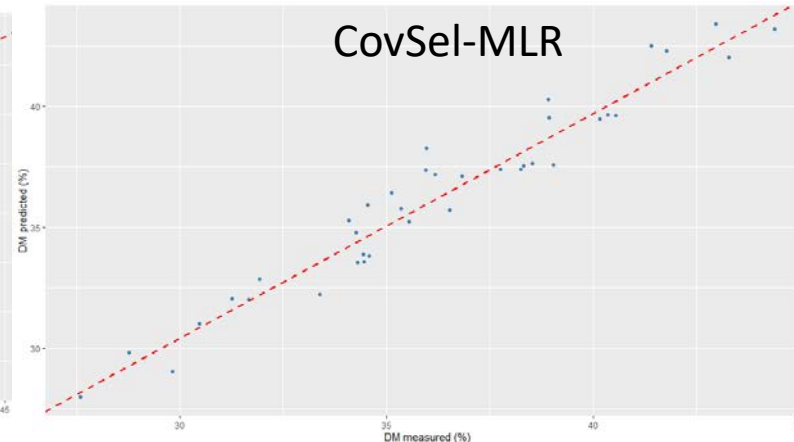
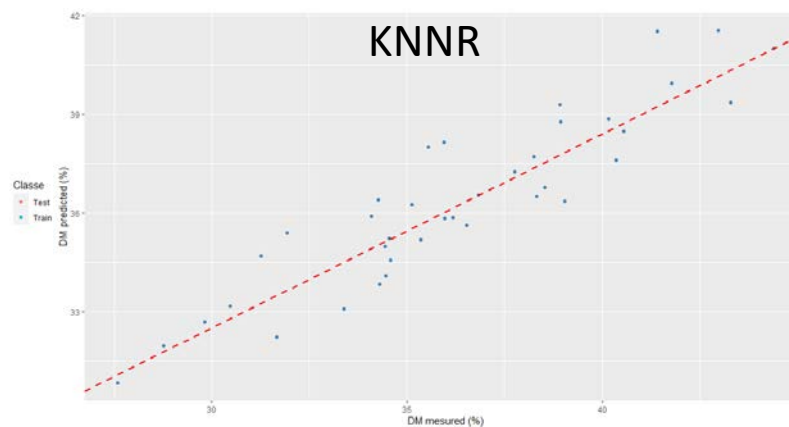
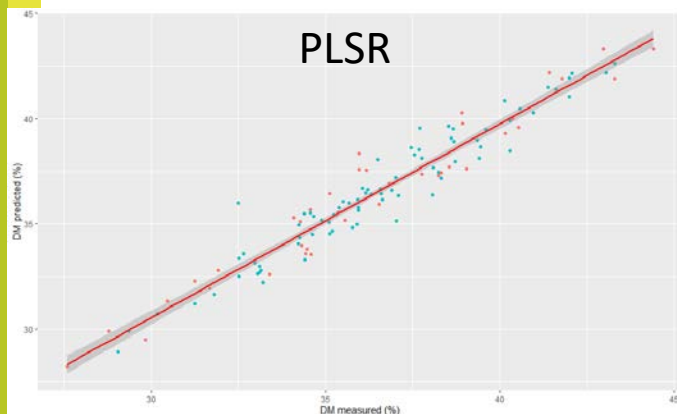


# Results : Prediction of DMC in fresh cassava



- Performances of different models developed

| Model      | Ncal | Parameters   | Calibration |       |        |        |           |                    | Validation |                  |           |      |
|------------|------|--|-------------|-------|--------|--------|-----------|--------------------|------------|------------------|-----------|------|
|            |      |  | Mean(%)     | SD(%) | Min(%) | Max(%) | RMSEC (%) | R <sup>2</sup> cal | Nval       | R <sup>2</sup> p | RMSEP (%) | RPD  |
| PLSR       | 78   | LV=7   | 36.40       | 3.46  | 27.59  | 44.40  | 0.78      | 0.94               | 38         | 0.93             | 1.06      | 3.26 |
| KNNR       | 78   | nIvdis=5, h=.2, k=40                               | 36.40       | 3.46  | 27.59  | 44.40  | 0.16      | 0.99               | 38         | 0.77             | 1.94      | 1.78 |
| SVMR       | 78   | cost=5, epsilon=0.2, gamma=3                       | 36.40       | 3.46  | 27.59  | 44.40  | 2.85      | 0.17               | 38         | 0.46             | 3.63      | 0.95 |
| Covsel_MLR | 78   | (932,1021,1130,1169,1222,1385,1410,1444,1484,1717) | 36.40       | 3.46  | 27.59  | 44.40  | 0.88      | 0.92               | 38         | 0.94             | 0.96      | 3.60 |

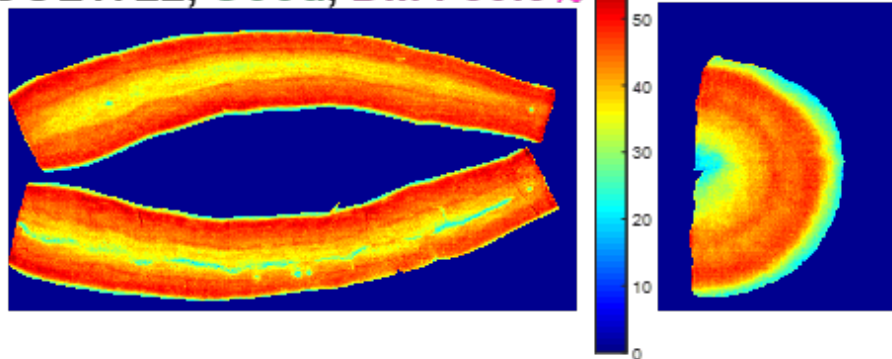




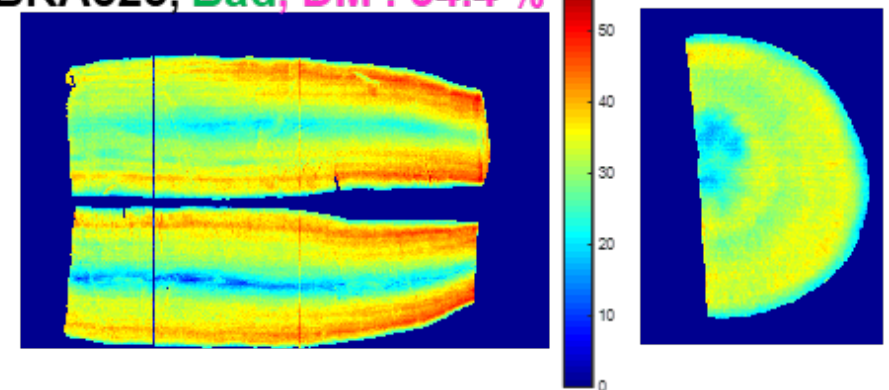
# Results : Prediction of DMC in fresh cassava

- Visualization of DMC distribution in contrasting cooking ability of cassava genotypes

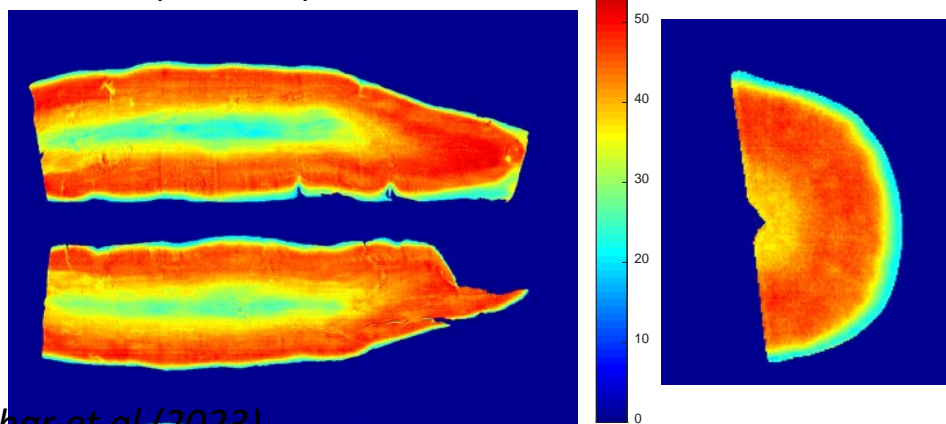
COL1722, Good, DM : 39.0%



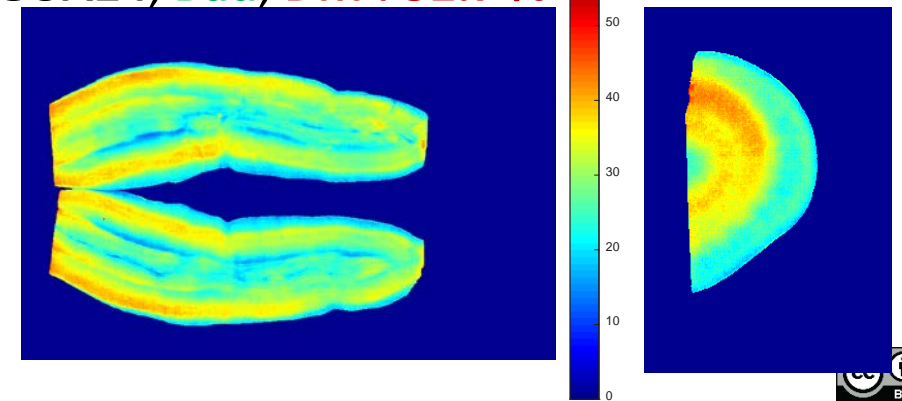
BRA325, Bad, DM : 34.4 %



IND135, Good, DM : 40.5 %

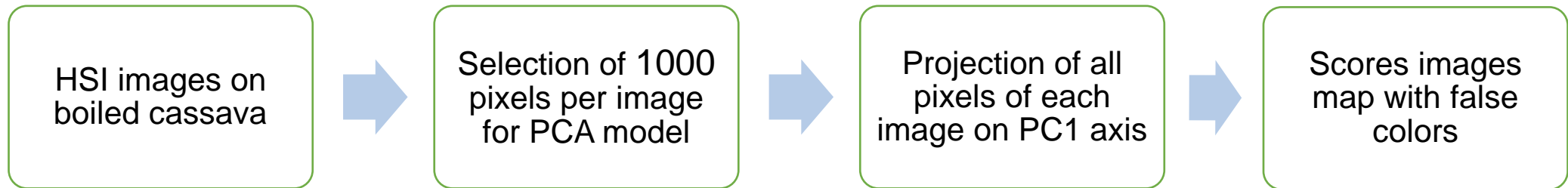


GUA24, Bad, DM : 32.7 %

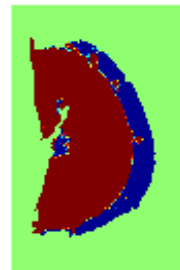


# Results : Prediction of WA30 in boiled cassava

- Visualization of Water absorption in contrasting cooking ability of boiled cassava



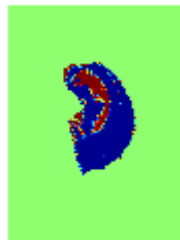
Good (CR63) : WAB=18.31 %, OCT = 20 min



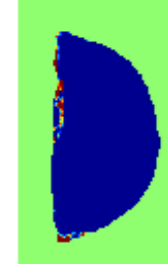
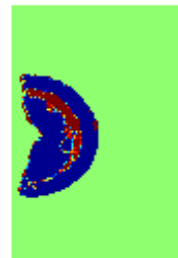
Bad (VEN25) : WAB =0.39 %, OCT=60 min



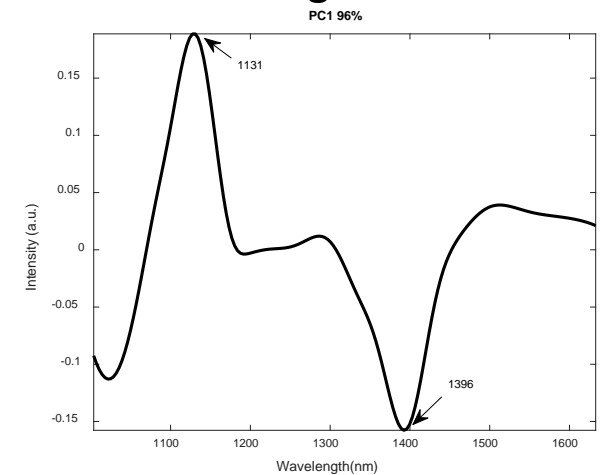
Good (CUB46) : WAB=13.34 %, OCT = 25 min



Bad (BRA512) : WAB=3.05 %, OCT = 60 min



Loading PC1



Absorption region from 1191 to 1385 nm which is associated with the first overtone of OH band in water ([Ignat et al., 2014](#));

# Conclusions and perspectives

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- High efficient model for visualization of DMC in fresh cassava genotypes,  $R^2_p=0.94$ ,  $RMSEP=0.96\%$ ,  $RPD=3.60$
- Visualisation of WAB30 using PC1 scores of PCA model.

## Perspectives

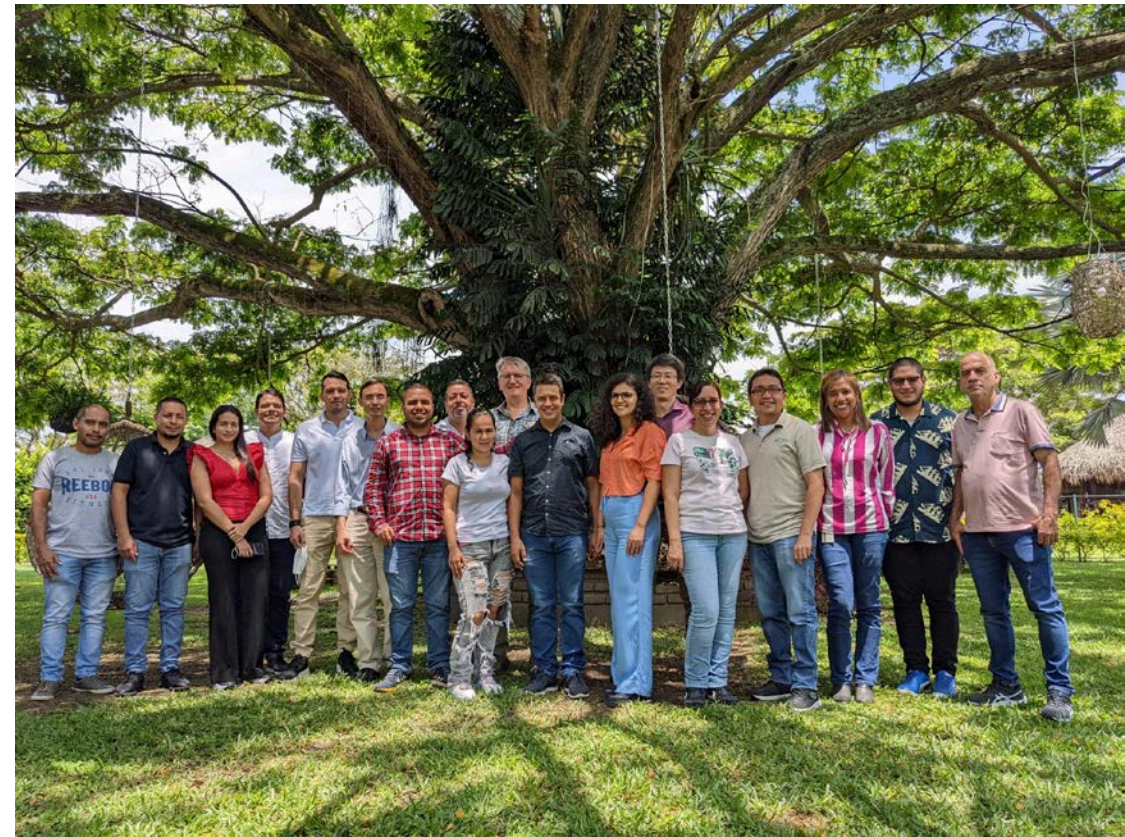
- Enhance the DMC model accuracy and robustness using more genotypes.
- Implementation of the HSI procedure within CIAT breeding program.
- Define acceptability threshold for WA30 classification model.

# Acknowledgements

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BILL & MELINDA  
GATES *foundation*



# Thank You For Your Attention !

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