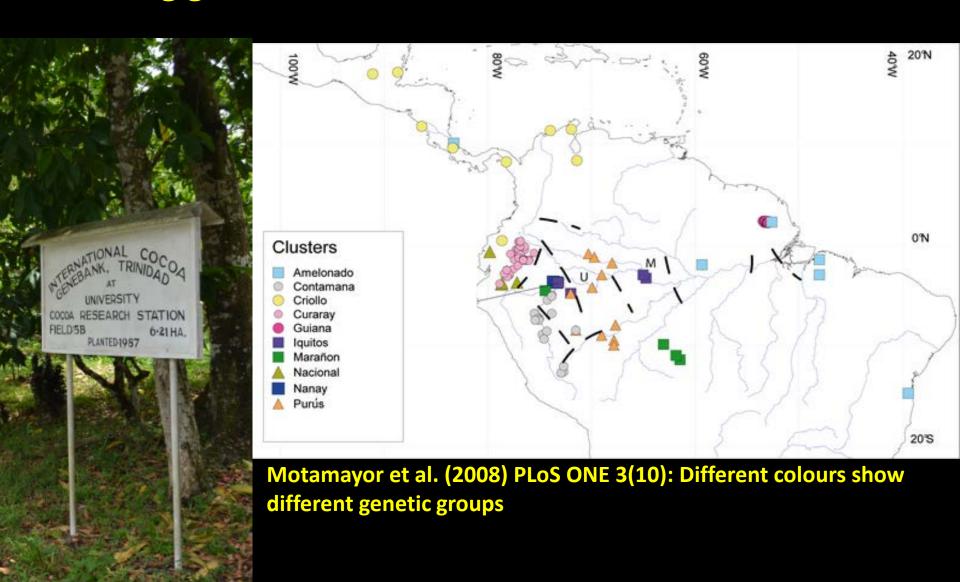


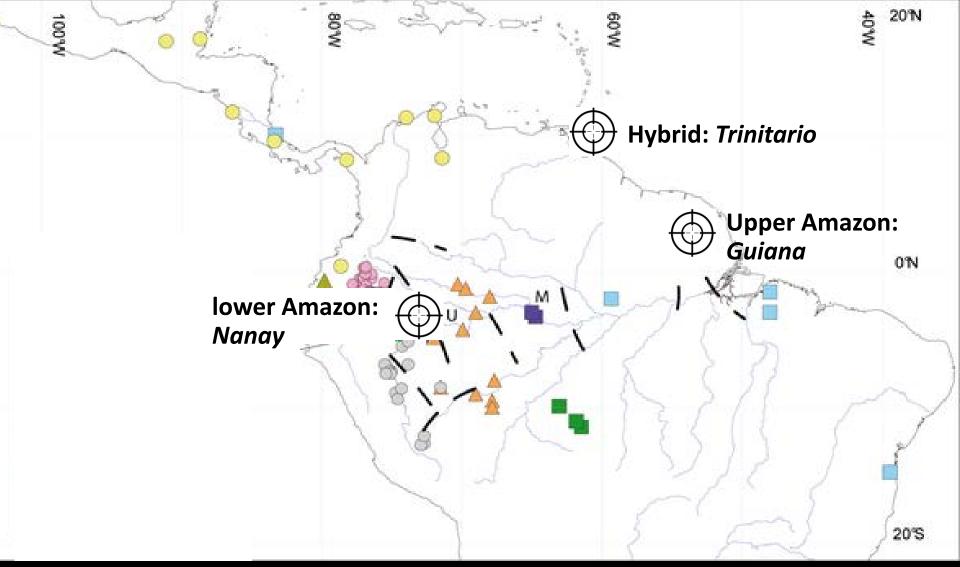
### Harnessing genetic differences to increase cocoa resilience?



#### Aims of workpackage 1: Climate Proof Screen

- Assess available methods for identifying differences in resilience to drought and heat stress
- Establish protocols for testing resilience and use these to compare three genetic groups
- Assess the potential impact of climate resilience on current and future cocoa crops (via workpackage 2 & 3)

## Three groups: testing three accessions from each



Motamayor et al. (2008) PLoS ONE 3(10):





## Assessing 'performance' in the field and greenhouse



Leaf temperature: Infrared camera (FLUKE 410)



**Transpiration**: Porometer (SC-1 Decagon Devices)



Photosynthesis: Chlorophyll fluorescence (Walz MINI PAM)

# Visual assessment of stress symptoms



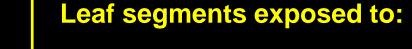
# Membrane stability test: to assess tolerance to 'damage' using leaves from field and greenhouse







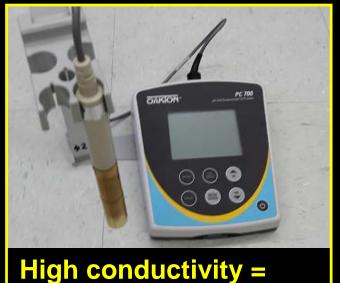




**30**; 35; 40; 45; **55** Celsius or **0**; -1.0; 2.0; **-3.0** Megapascals

Then incubated overnight



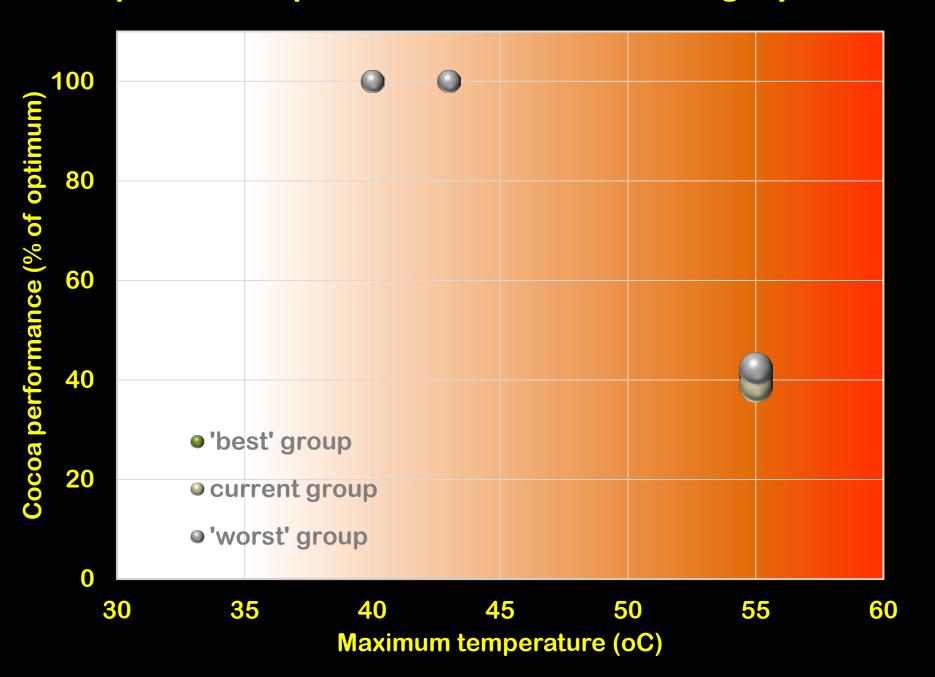


low membrane stability

#### Best method for assessing differences in heat stress

- In the greenhouse
  - Max air temperatures were approx. 5 C higher
  - Transpiration was consistently elevated
  - Photosynthesis unaffected
  - Visual assessment unaffected
  - Membrane stability test was consistent...

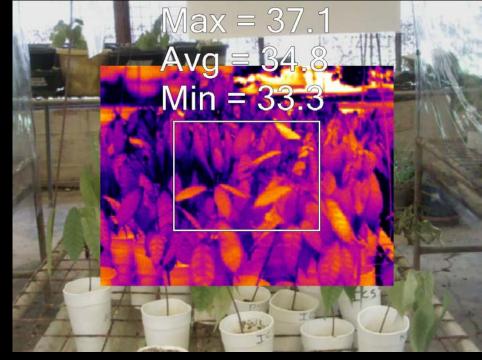
#### The impact of temperature on membrane integrity



#### Best method for assessing differences in water stress

- In the greenhouse
  - Stomatal conductance reduced
  - Leaf temperature elevated
  - Membrane stability test- inconsistent
  - Visual index was reduced
  - Photosynthesis was reduced...





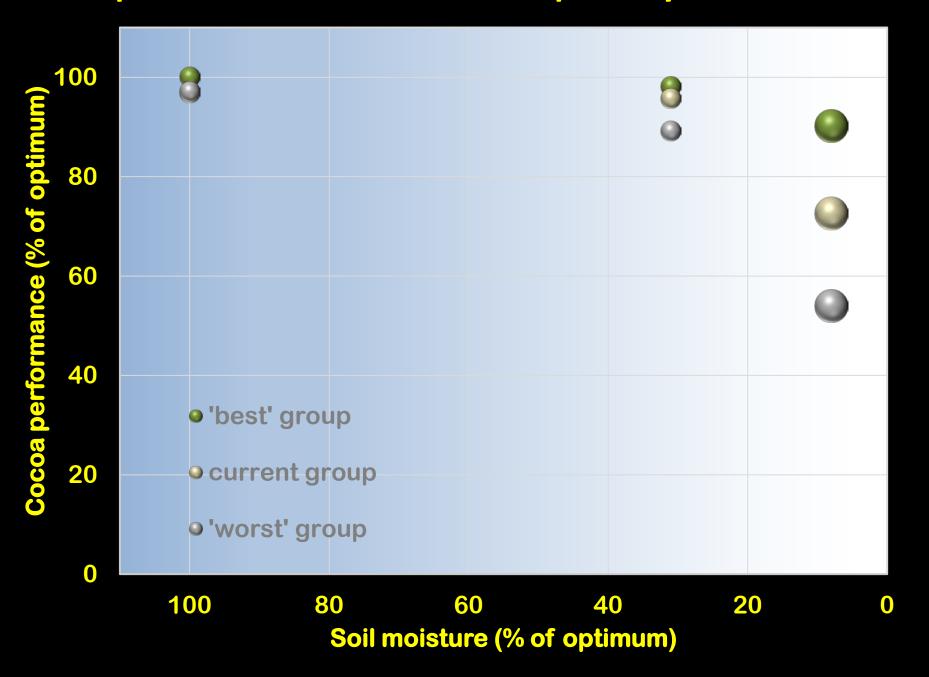
**Control Bench** 



**Dry Bench** 



#### The impact of moisture content on photosythesis



**Conclusions of workpackage 1 : Climate Proof Screen** 

Selection for resilience to drought stress has great potential

Several methods available, including low-tech methods that could be implemented through participatory selection

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Figure 1. Localization of the origin of individuals analyzed; colors indicate the inferred genetic cluster to which they belong.

