

AN AUTOMATED WHOLE-CANOPY MULTI CHAMBER SYSTEM FOR LIVE MONITORING WATER USE EFFICIENCY IN SORGHUM

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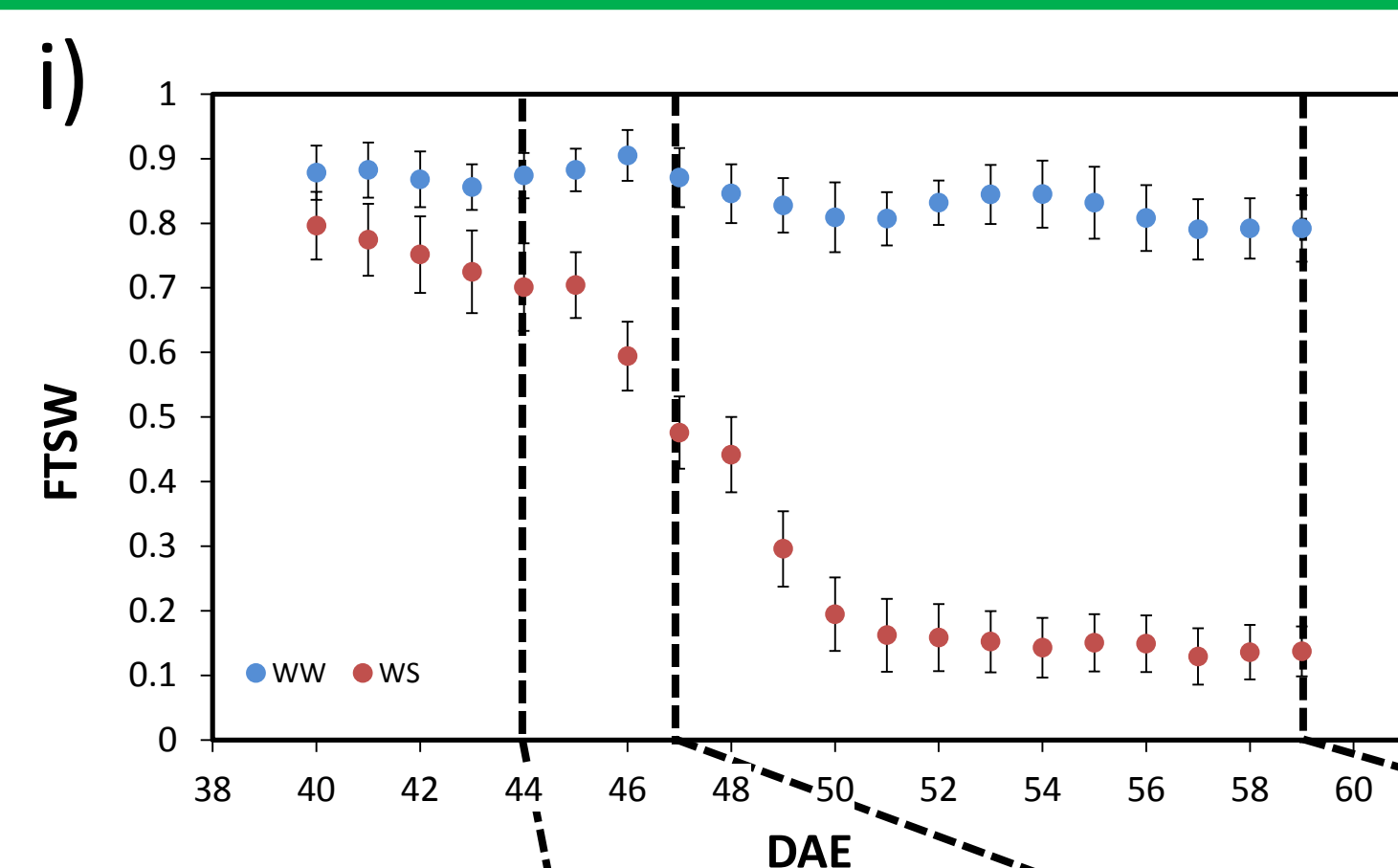
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Introduction

Plant growth and productivity are strongly affected by limited water availability in drought prone environments. The current climate change scenario, characterized by long periods without precipitations followed by short but intense rainfall, force plants to implement different strategies to cope with drought stress.

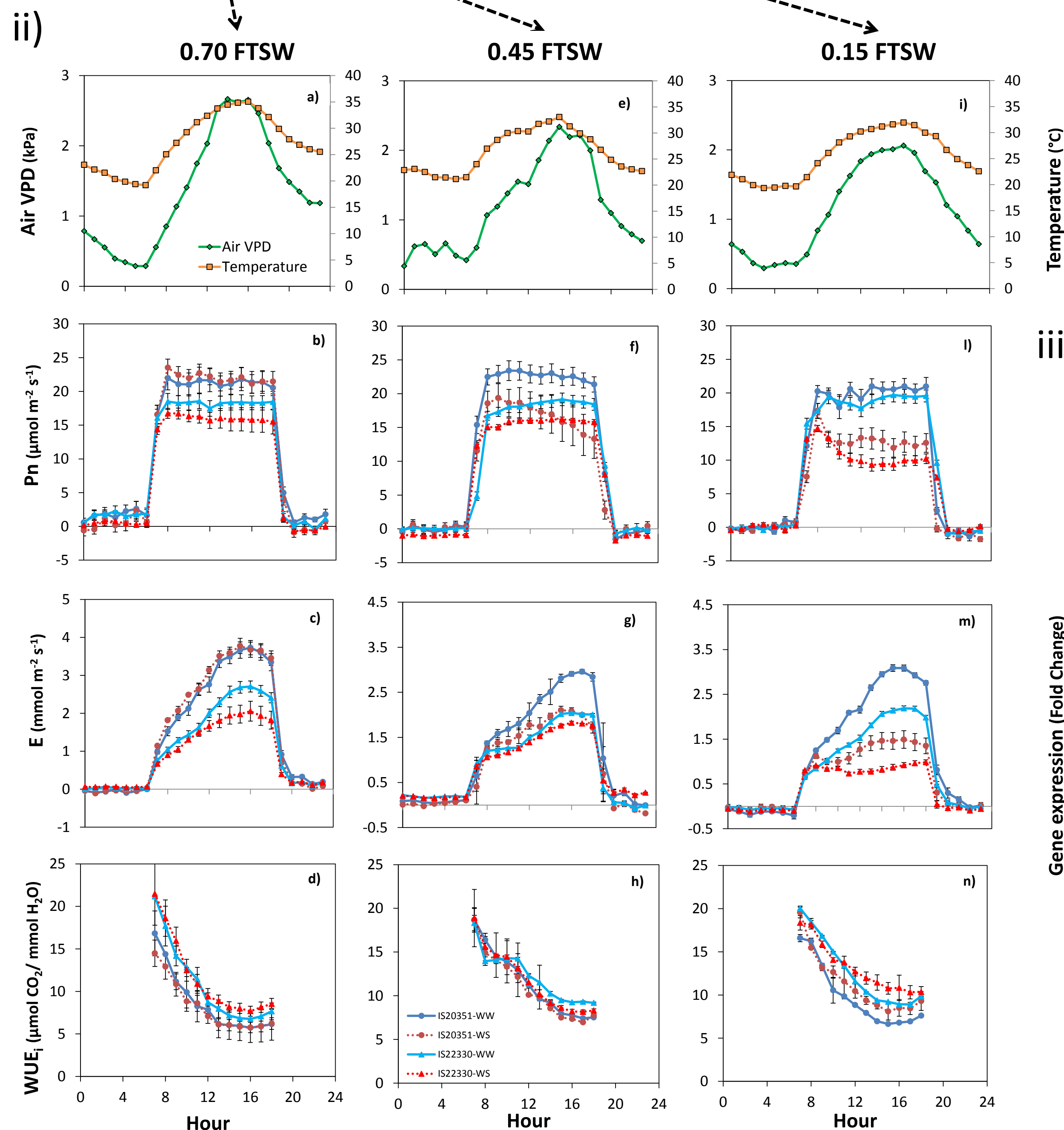
Objective

Understanding how plants use water during these period of limited water availability, and which are the molecular mechanisms at the basis, is of primary importance to identify and select the best adapted genotypes to a certain environment.



Material and methods

An automated whole-canopy multi chamber system was used to monitor net photosynthetic rate (Pn), transpiration rate (E) and water use efficiency of two sorghum genotypes (IS20351 and IS22330) during a dry-down experiment (Fig.1 and 2i) with a daily trend (Fig. 2ii). Gene expression dynamics of five drought related genes was evaluated every four hours at 0,15 FTSW (Figure 2iii).



Results

- Continuous and non-destructive measurements of the **whole-canopy Pn, E and WUE_i** were discriminated in response to progressive drought stress;
- The drought tolerant IS22330 adopts a “pessimistic” strategy, having low Pn and E under WS conditions and achieving overall a WUE_i higher than the drought sensitive IS20351 (Fig. 2B)

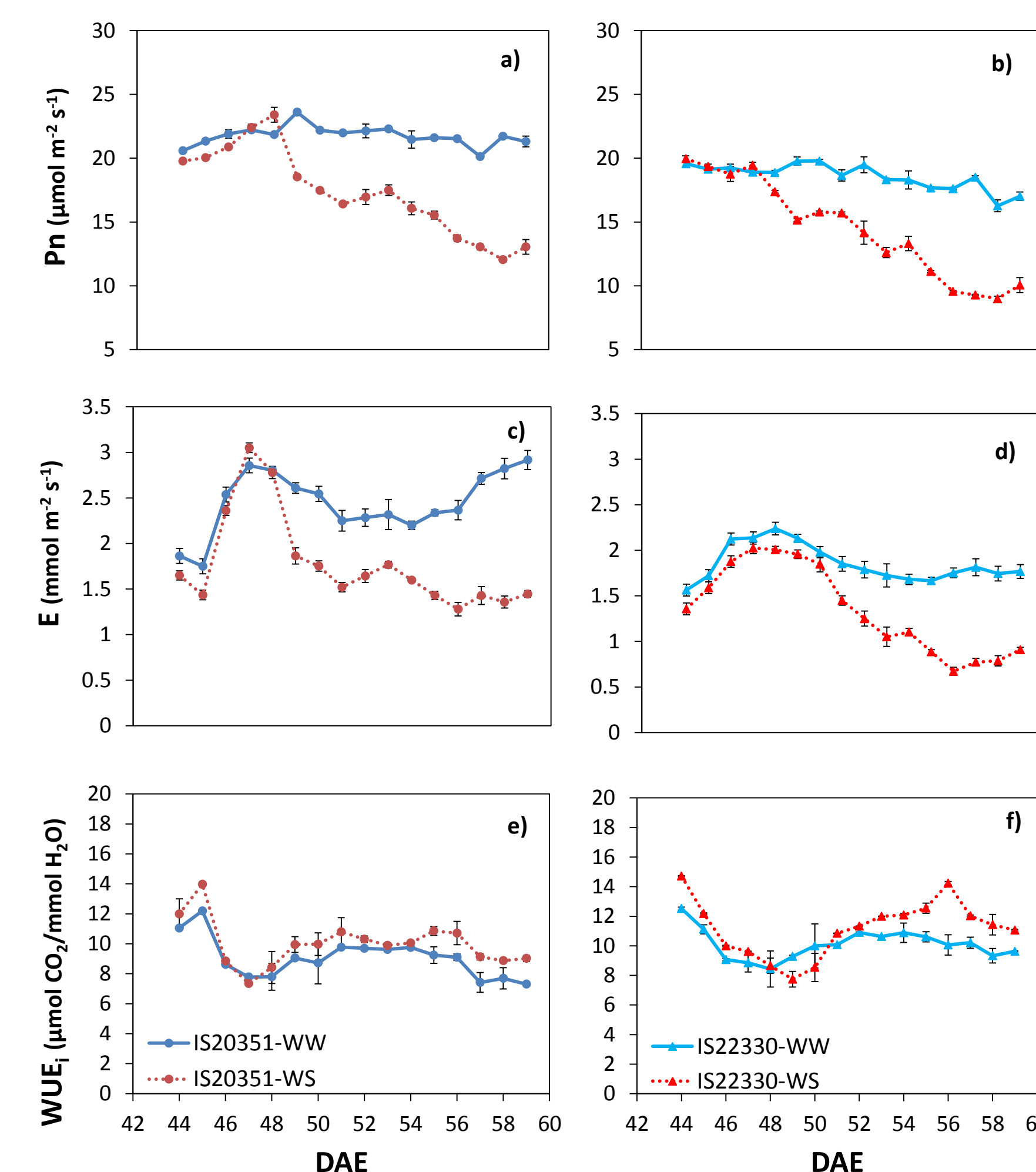
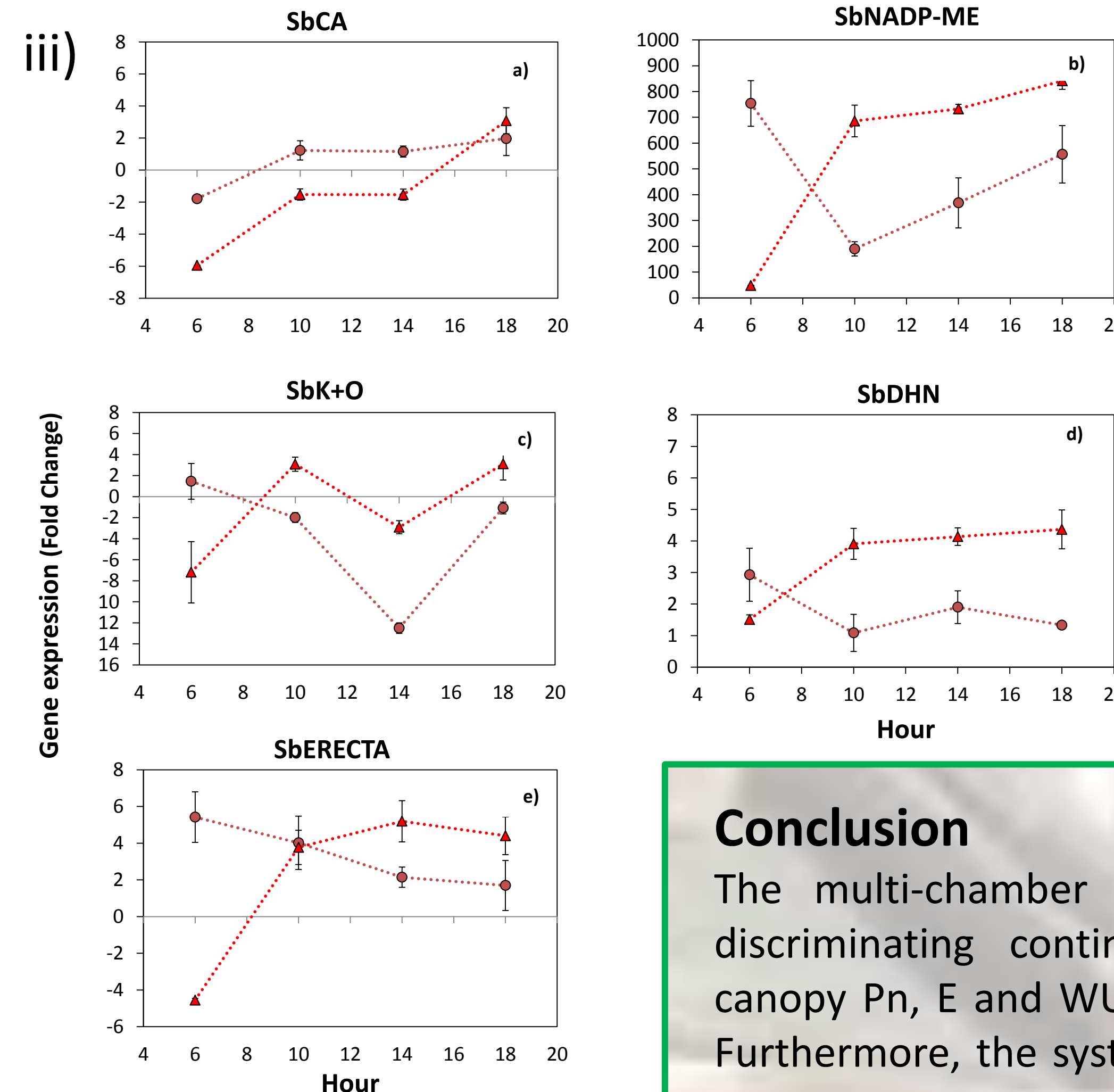


Figure 1: Net photosynthetic rate, transpiration rate and WUE_i recorded at whole-canopy level during the experiment for the two genotype IS20351 (circles) and IS22330 (triangles) under well-watered (WW, blue) and water-stressed (WS, red) conditions.

- The drought tolerance strategy of IS22330 is pursued through a fine control of stomata apertures involving regulation of K⁺ channels expression levels (*SbK+O*) and regulation of C4 cycle gene expression levels (*SbNADP-ME* and *SbCA*).
- *SbERECTA* could be used as a proxy for WUE_i determination while *SbDHN* was confirmed as candidate gene to assess drought tolerance in sorghum.

Conclusion

The multi-chamber system proved to be a valuable tool in discriminating continuously and non-destructively the whole-canopy Pn, E and WUE_i in response to progressive drought stress. Furthermore, the system allowed to unravel strategies adopted by genotypes to cope with drought stress and to identify the right moment to perform destructive sampling for transcriptomic analysis.