

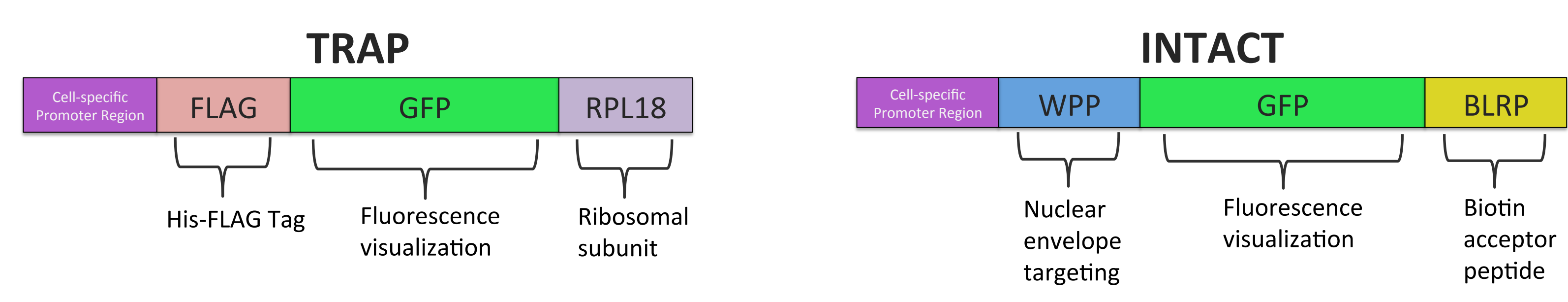


How do cells respond to stress and developmental cues?

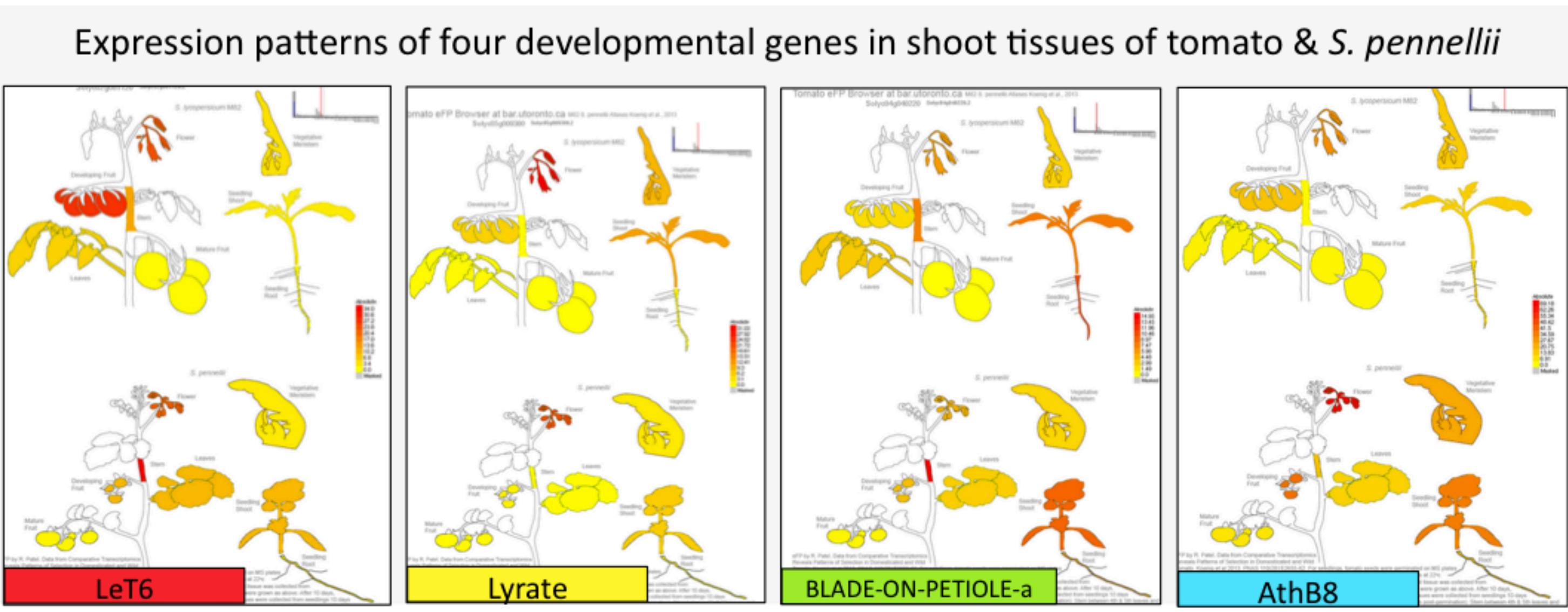
Donnelly A. West, Aashish Ranjan, Brad Townsley, Kaisa Kajala, & Neelima R. Sinha

ABSTRACT: Leaf shape is tremendously diverse in nature, yet overall leaf function in photosynthesis is conserved through development. How the leaf responds to stress, hormonal, genetic, and environmental cues is relatively unexplored. To dissect molecular leaf development, cells transitioning between meristematic and leaf tissue types must be separated from homogeneous, lineage-independent clusters. Using Isolation of Nuclei Tagged in specific Cell Types (INTACT) and Translating Ribosome Affinity Purification (TRAP), cell-type specific molecular data can be isolated with confidence. In addition to discrete cell population differences, these techniques can be used in conjunction with water stresses to investigate developmental plasticity in a domesticated crop (*Solanum lycopersicum* var *M82*; tomato) and its desert, wild relative (*Solanum pennellii*). These studies will enable the identification of molecular events that determine leaf shape across two morphologically distinct *Solanum* species and those that determine their relative responses to water stress conditions.

Leaf Developmental Genetics via Cell Specificity



Key Leaf Patterning Genes



LeT6

Arabidopsis ortholog: SHOOT MERISTEMLESS (STM), a *KNOTTED-LIKE1 HOMEBOX (KNOX)* gene

Expression Domain: central zone of the shoot apical meristem (SAM)

Possible Function: maintaining meristem identity

LYRATE (LYR)

Arabidopsis ortholog: JAGGED (JAG)

Expression Domain: in the SAM, incipient leaf primordia (P0), leaflet initiation sites

Possible Function: a *KNOX* repressor gene; allows organ initiation¹

BLADE-ON-PETIOLE-a (BOPa)

Arabidopsis ortholog: BLADE-ON-PETIOLE-1

Expression Domain: boundary zones in leaf primordia²; mature leaf tissue mesophyll

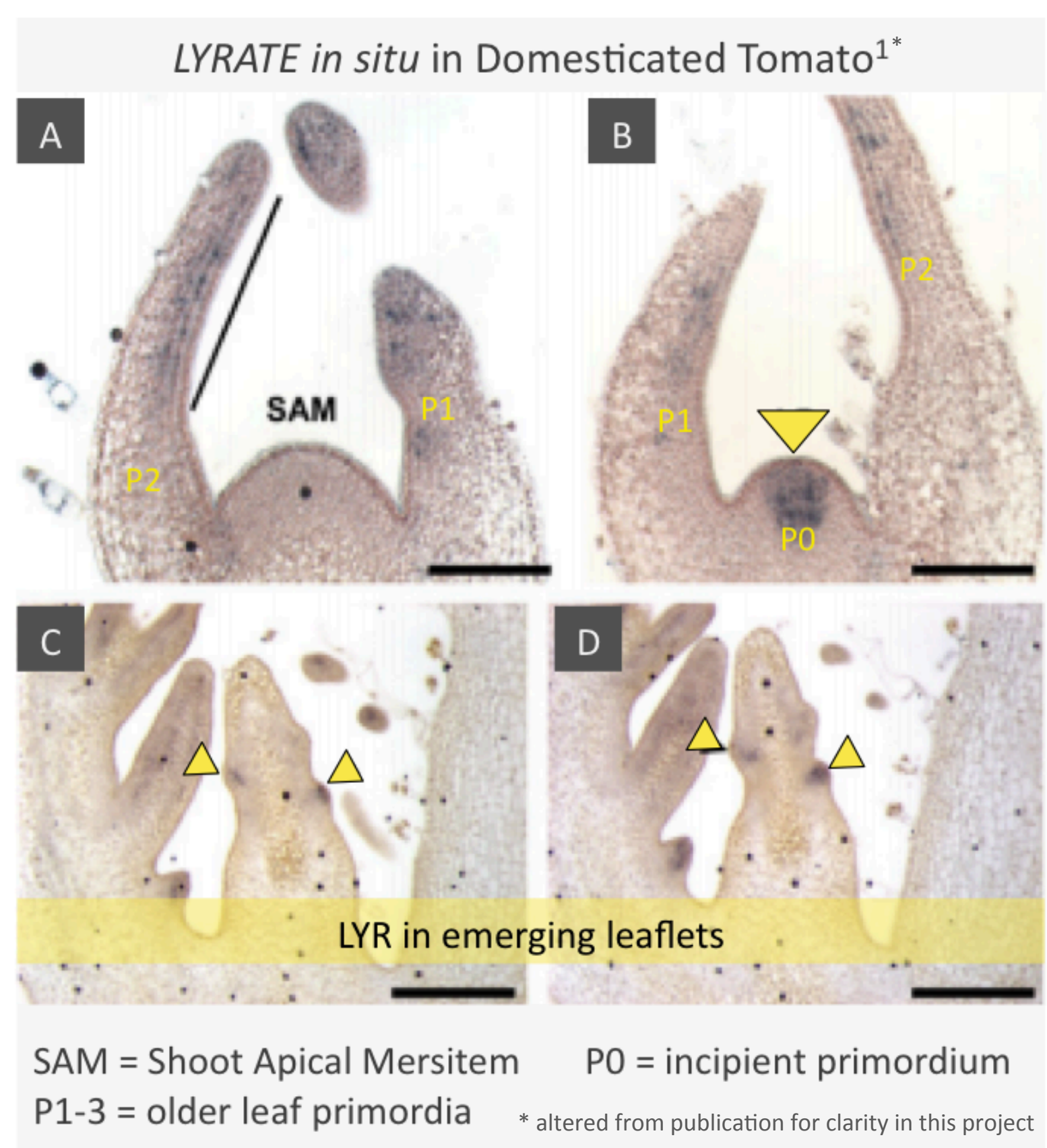
Possible Function: repression of JAG; further organization of leaf tissue cell types


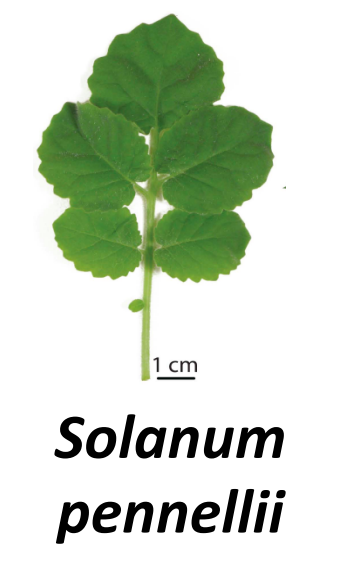
AthB8

Arabidopsis ortholog: ATHB8; class III homeodomain-leucine zipper (*HD-ZIPIII*)

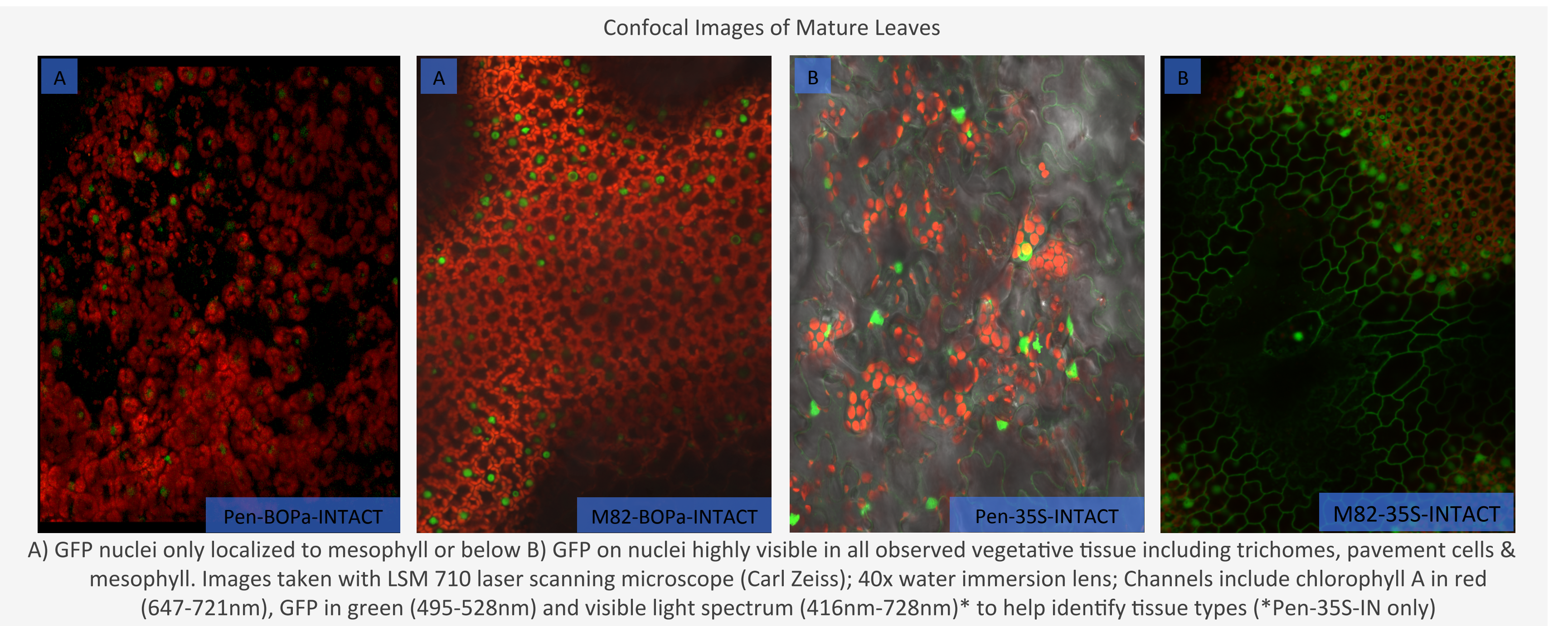
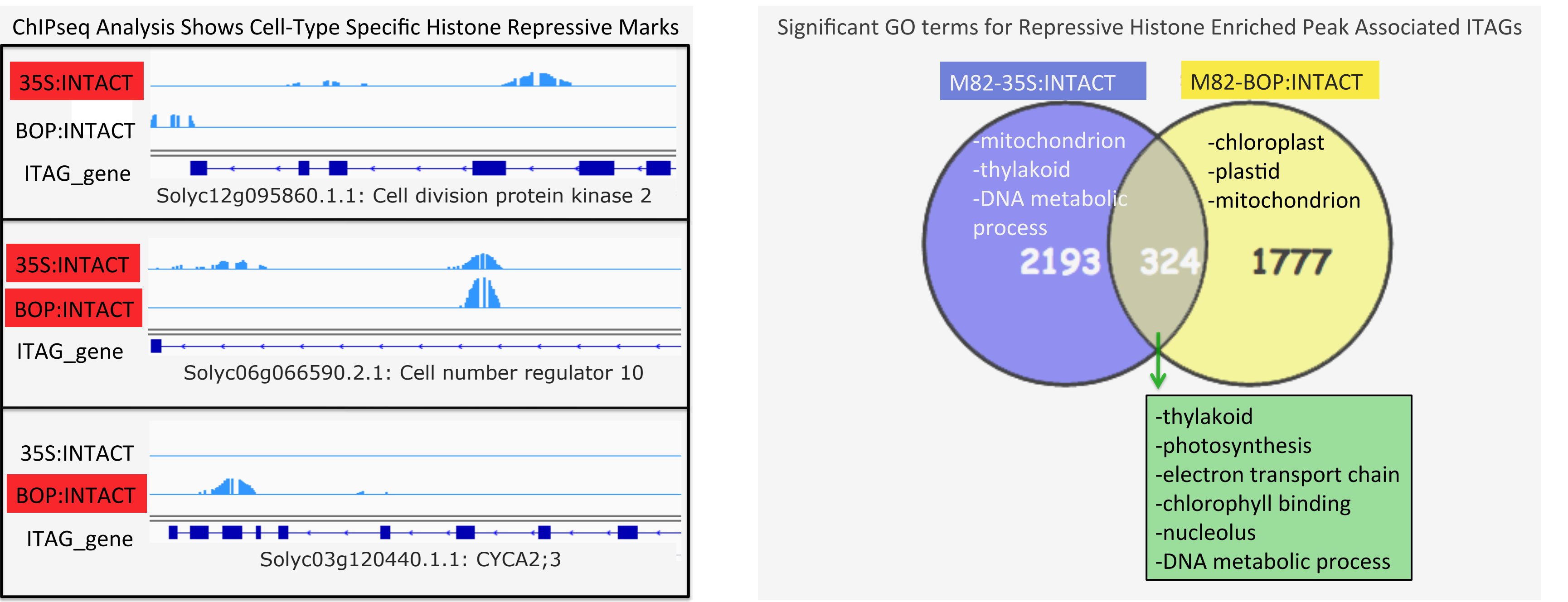
Expression Domain: in leaf primordia, along vascular track, after auxin accumulation before differentiation

Possible Function: cell fate commitment to become vasculature tissue³



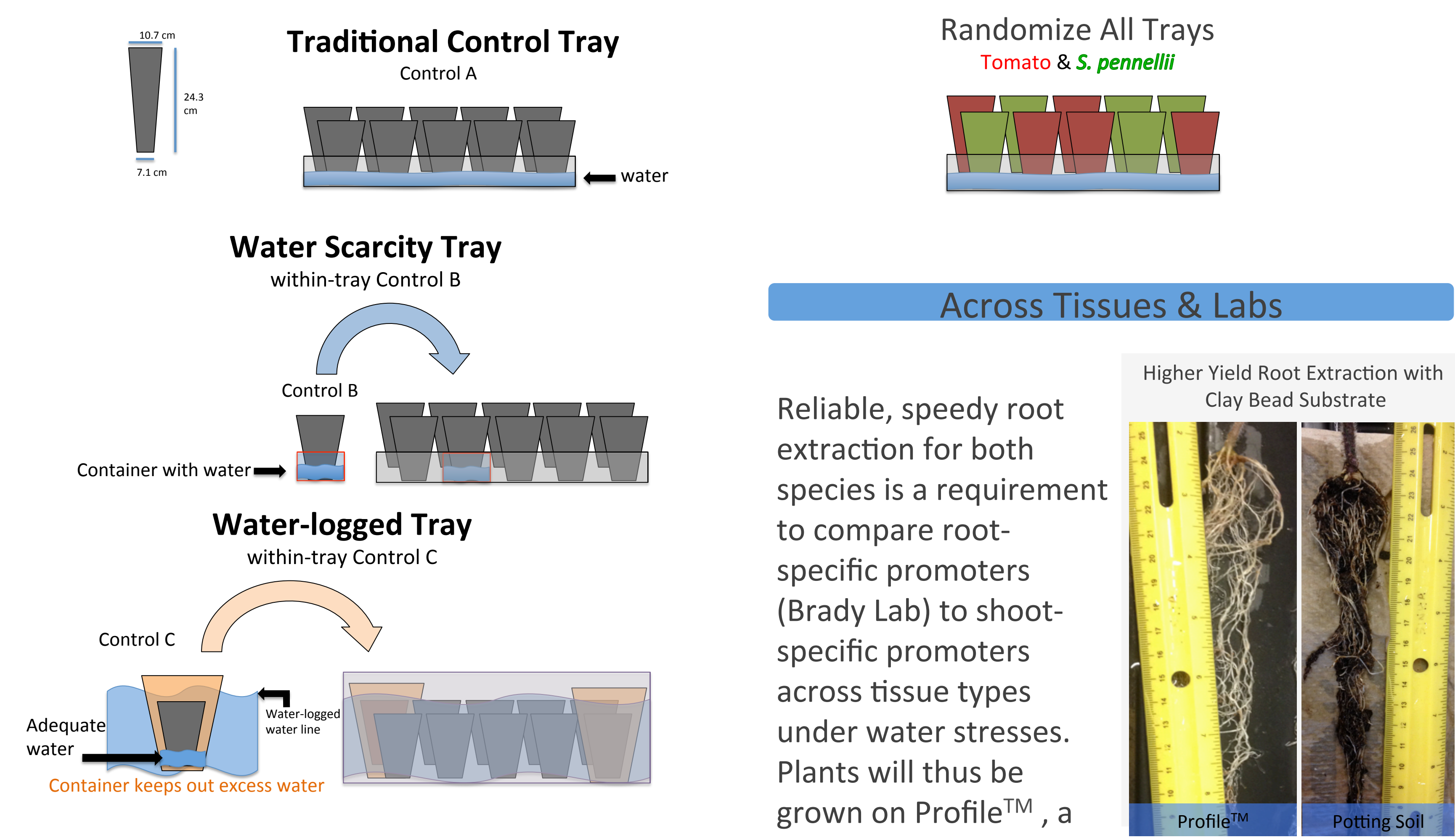
Planned Transgenic Lines for Leaf Development		
 <i>Solanum lycopersicum</i>	INTACT - ChIP & Nuclear Transcriptome	LeT6
		LYR ✓
		BOPa ✓
		AthB8 ✓
	TRAP - Translatome & Ribosomal Footprints	LeT6 LYR ✓ BOPa ✓ AthB8
 <i>Solanum pennellii</i>	INTACT - ChIP & Nuclear Transcriptome	LeT6
		LYR ✓
		BOPa ✓
		AthB8
	TRAP - Translatome & Ribosomal Footprints	LeT6 LYR BOPa AthB8

Preliminary ChIP seq Results: 35S vs BOPa in Mature Leaves



Leaf Developmental Genetics under Water Stress

Once T3 lines are established, water stress experiments will be conducted over at least 3 biological replicates with 20 plants per replicate. In order to maintain growth conditions and minimize pest influences to compare between constructs, promoters and root & shoot systems, stress experiments will be conducted in growth chambers. Because positional effects have been demonstrated⁴, within-tray controls will be utilized



Acknowledgements:
UC Davis – Sinha Lab, Brady Lab, Maloof Lab, CEF / Greenhouse Staff, & Integrative Genetics & Genomics Group. Other Collaborators - Bailey-Serres Lab, Deal Lab
Funding – National Science Foundation Plant Genome Project Cell Fate Plasticity Grant Award #1238243

References:
1. David-Schwartz R, Koenig D, Sinha NR. Plant Cell. 2009 Oct;21(10):3093-104. Epub 2009 Oct 9.PMID: 19820188
2. Ichihashi,Y., Aguilar-Martínez, J.A., Farhi,M., Chitwood, D.H., Kumar,R., Millon, L.V., Peng, J., Maloof, J.N., and N.R. Sinha. PNAS 2014 111 (25) E2616-E2621; published ahead of print June 9, 2014, doi:10.1073/pnas.1402835111
3. Baïma, S. et al. Plant Physiology 2001. 126:643–655.
4. Chitwood, DH, Kumar, R, Headland, LR, Ranjan, A, Covington, MF, Ichihashi, Y, Fulop, D, Jimenez-Gomez, JM, Peng, J, Maloof, JN, Sinha, NR (2013). Plant Cell. 25: 2465-81.

Or follow us on Twitter: Donnelly @CatsCauseTypos & Neelima @PlantDevo