



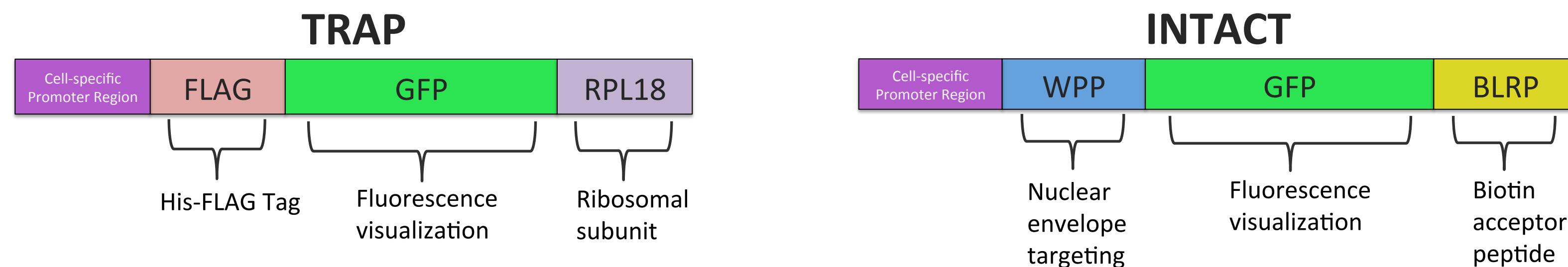
# How do cells respond to stress and developmental cues?

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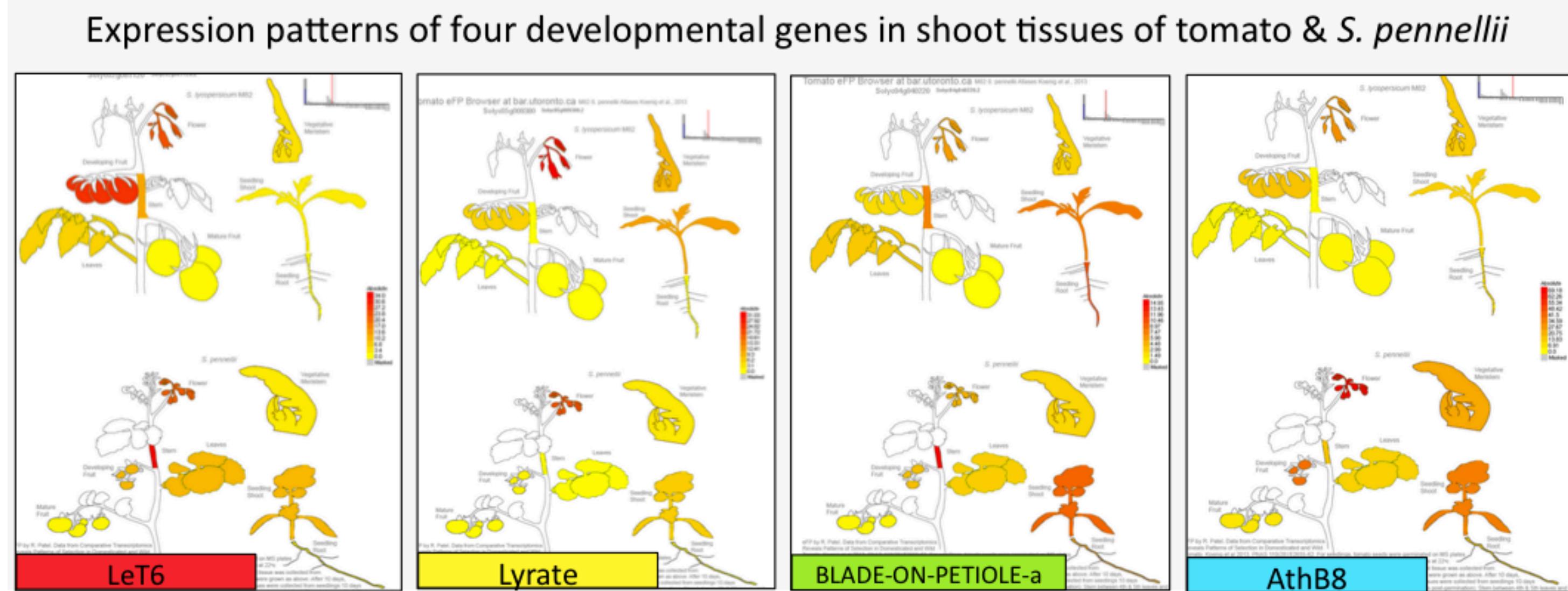


**ABSTRACT:** Leaf shape is tremendously diverse in nature, yet overall leaf function in photosynthesis is conserved through development. How the leaf responds to stress, hormonal, genetic, and environmental cues is relatively unexplored. To dissect molecular leaf development, cells transitioning between 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 HOMEOBOX (KNOX) gene  
**Expression Domain:** central zone of the shoot apical meristem (SAM)  
**Possible Function:** maintaining meristem identity

#### LYRATE (LYR)

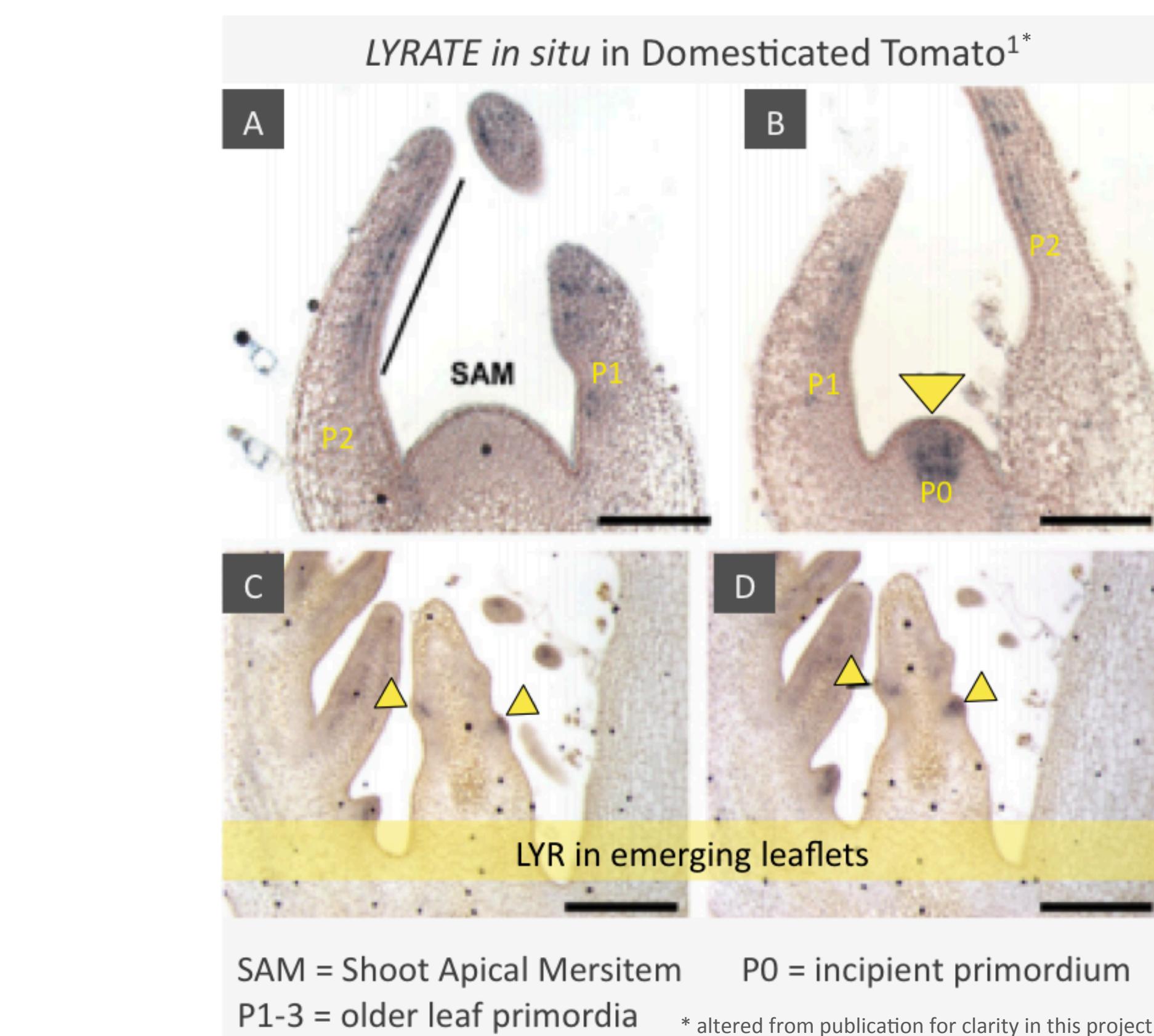
**Arabidopsis ortholog:** JAGGED (JAG)  
**Expression Domain:** in the SAM, incipient leaf primordia (PO), leaflet initiation sites  
**Possible Function:** a KNOX repressor gene; allows organ initiation<sup>1</sup>

#### BLADE-ON-PETIOLE-a (BOPa)

**Arabidopsis ortholog:** BLADE-ON-PETIOLE-1  
**Expression Domain:** boundary zones in leaf primordia<sup>2</sup>; mature leaf tissue mesophyll  
**Possible Function:** repression of JAG; further organization of leaf tissue cell types

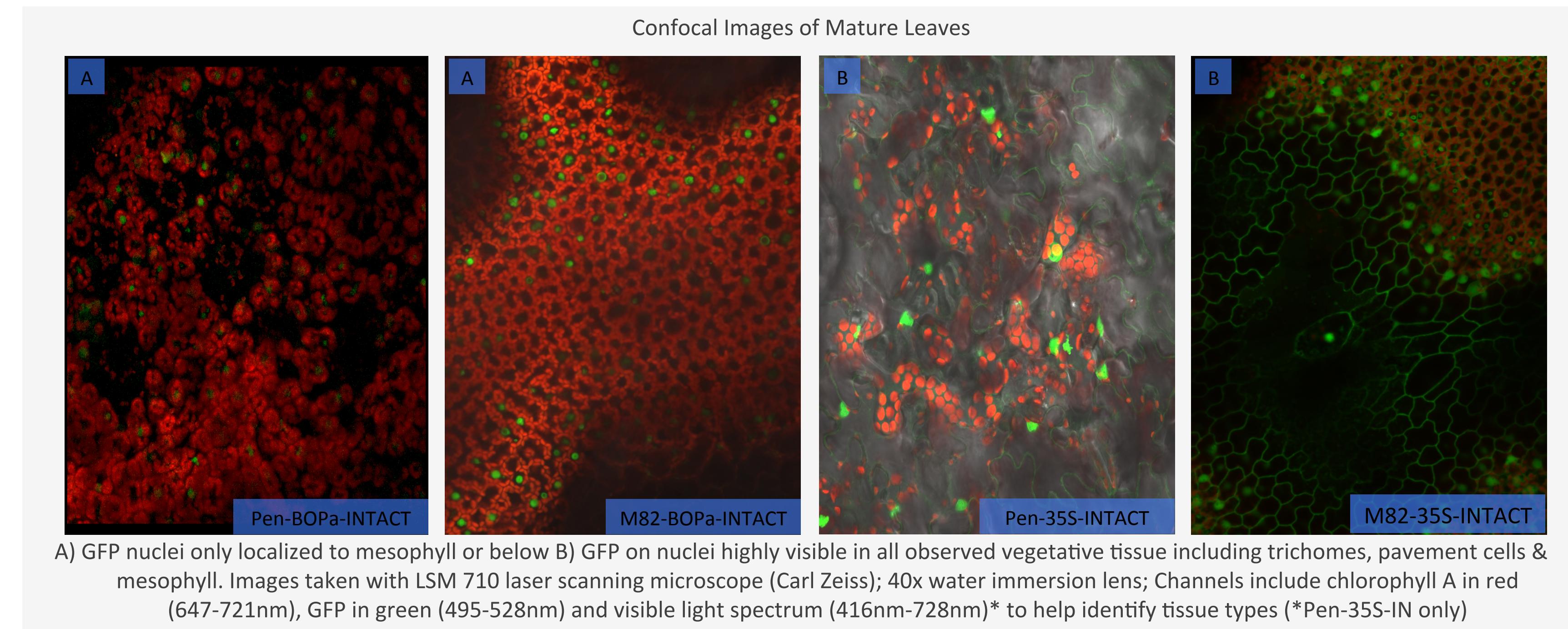
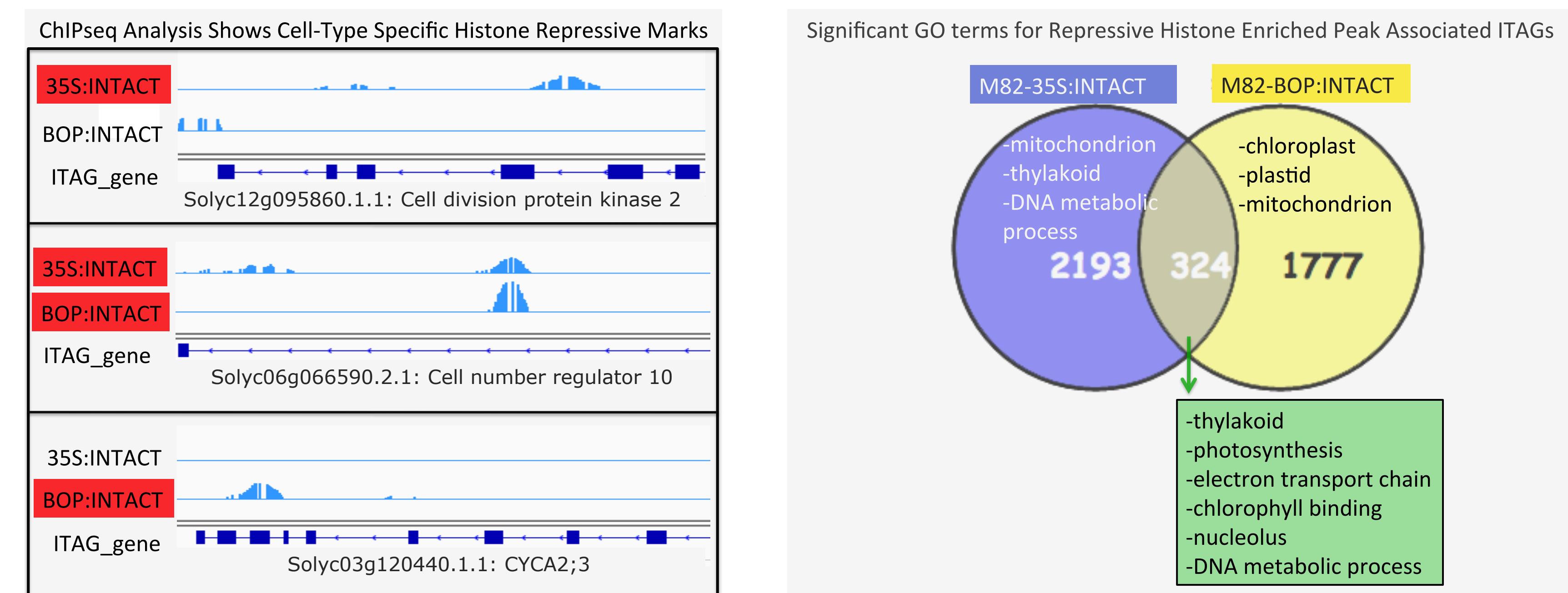
#### AthB8

**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 tissue<sup>3</sup>



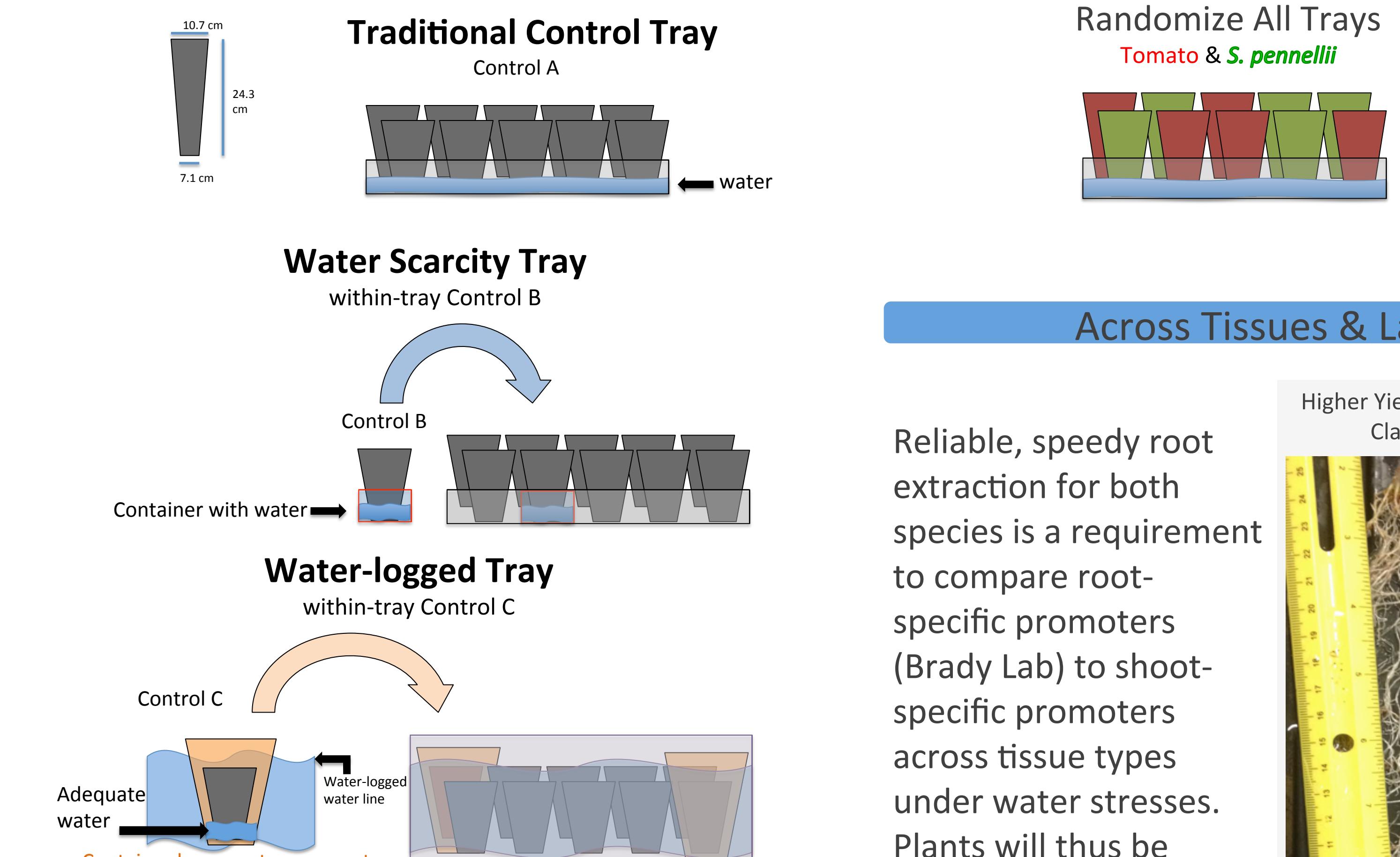
Planned Transgenic Lines for Leaf Development	
	INTACT - ChiP & Nuclear Transcriptome
	LeT6
	LYR ✓
	BOPa ✓
AthB8 ✓	
TRAP - Translatome & Ribosomal Footprints	
LeT6	
LYR ✓	
BOPa ✓	
AthB8	
	INTACT - ChiP & Nuclear Transcriptome
	LeT6
	LYR ✓
	BOPa ✓
AthB8	
TRAP - Translatome & Ribosomal Footprints	
LeT6	
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 demonstrated<sup>4</sup>, within-tray controls will be utilized



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## Across Tissues & Labs



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Plasticity	Sinha Lab	Tomato eQTL

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