

# Reproduction control tools for *Eucalyptus*

## Knockout of flowering and meiosis genes using CRISPR/Cas9 -- Effects on male-sterility and vegetative development

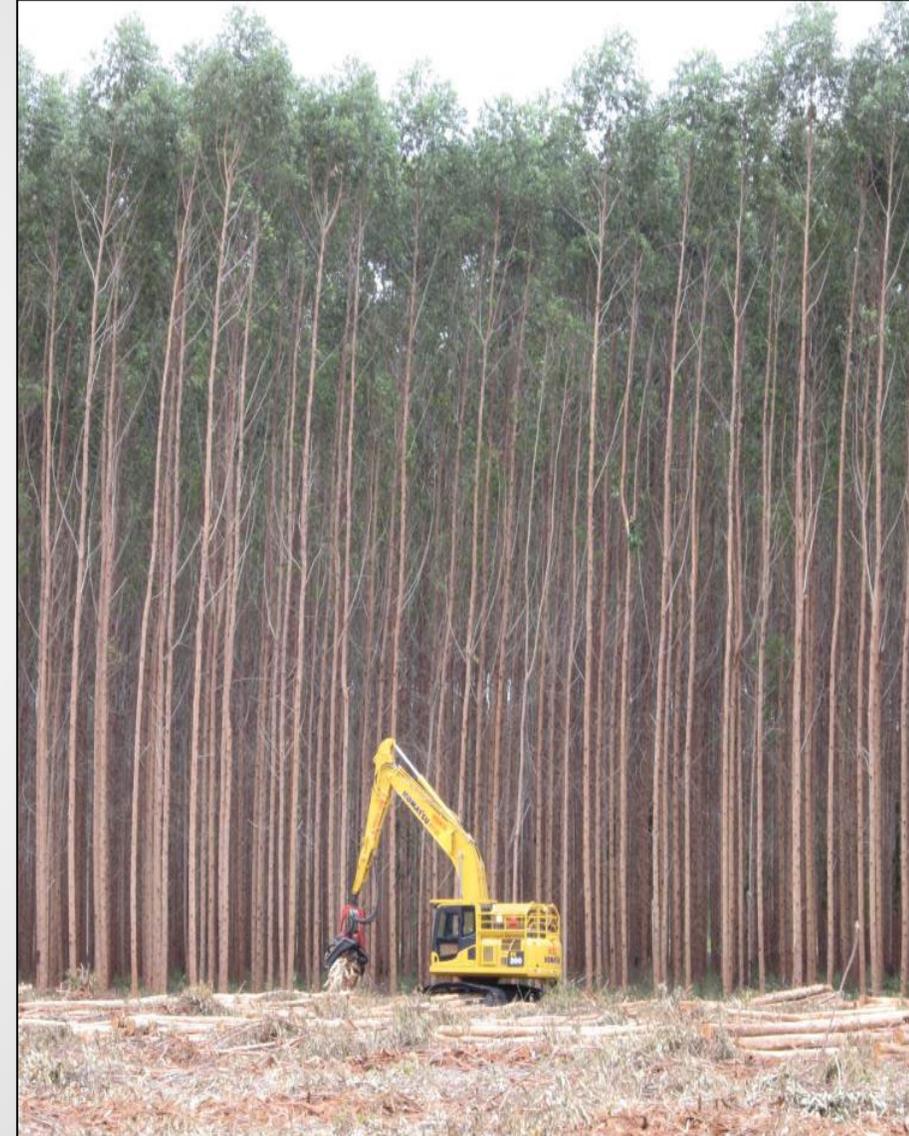
Steve Strauss, Oregon State University



Michael Nagle,  
Ph.D. Candidate

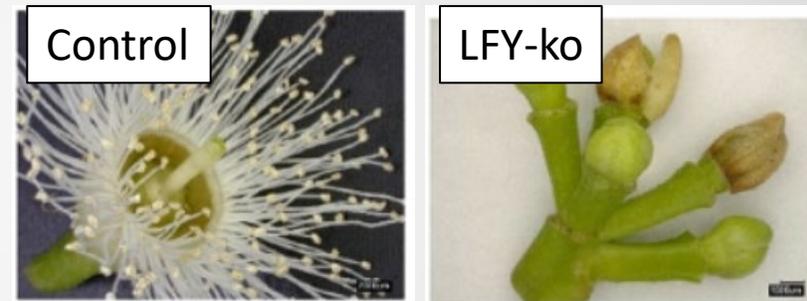
# Rationale

- *Eucalyptus* of global significance
  - Uses include fiber, energy, wood, ornamentals
- Spread of *Eucalyptus* often raises social/environmental concerns about invasiveness
- Transgenic traits tend to increase public and regulatory concerns from dispersal - within and outside native range
- Hybrids in common use: Male-sterility may facilitate seed production
- General study goal: Use CRISPR to knock-out of selected floral genes in both early-flowering and wild-type genetic backgrounds: male-reproduction and vegetative growth/form



# Floral modification a major emphasis of work in lab over many years

- Knockout of *Eucalyptus* ortholog *LEAFY* (*LFY*) using CRISPR/Cas
- Studied in rapid-flowering FT-overexpression lines
- High biallelic mutation rates, near 100% when transformation successful
- Floral disruption – nearly complete lack of flowers



Estefania Elorriaga, Ph.D.

Plant Biotechnology Journal

Plant Biotechnology Journal (2021) 19, pp. 1743–1755

doi: 10.1111/pbi.13588

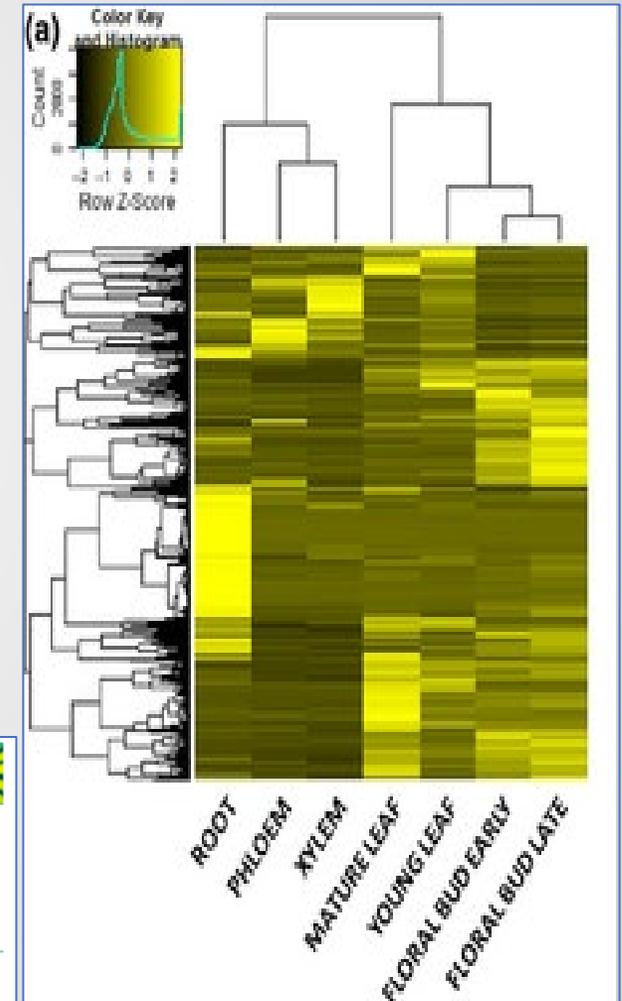
**Genetic containment in vegetatively propagated forest trees: CRISPR disruption of *LEAFY* function in *Eucalyptus* gives sterile indeterminate inflorescences and normal juvenile development**

Estefania Elorriaga<sup>1,a</sup>, Amy L. Klocko<sup>2</sup>, Cathleen Ma<sup>1</sup>, Marc du Plessis<sup>3</sup>, Xinmin An<sup>4</sup>, Alexander A. Myburg<sup>5</sup> and Steven H. Strauss<sup>1,\*</sup> 

<sup>1</sup>Department of Forest Ecosystems and Society, Oregon State University, Corvallis, OR, USA

# Strategy for selection of new target genes – male, female, and complete sterility goals

- Began with list of floral-specific genes from Vining et al. 2014 floral transcriptome
- Selected targets critical for reproduction, but not vegetative development
- Ran BLAST, Smith-Waterman alignment and examined function in Arabidopsis orthologs (literature review)
- Also examined Arabidopsis meiotic genes directly

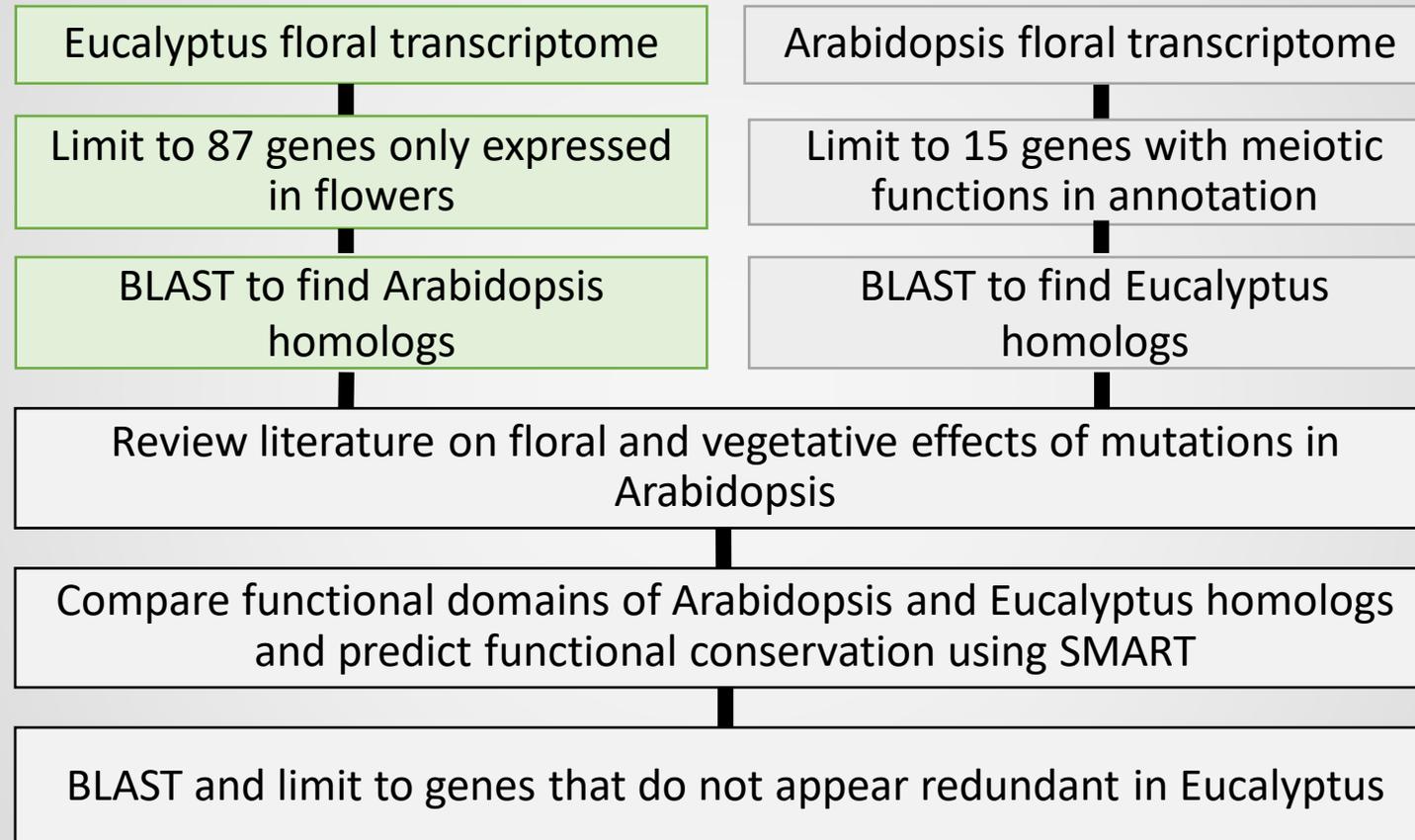


Research New Phytologist 

The floral transcriptome of *Eucalyptus grandis*

Kelly J. Vining<sup>1</sup>, Elisson Romanel<sup>2</sup>, Rebecca C. Jones<sup>3</sup>, Amy Klocko<sup>1</sup>, Marcio Alves-Ferreira<sup>4</sup>, Charles A. Hefer<sup>5</sup>, Vindhya Amarasinghe<sup>1,6</sup>, Palitha Dharmawardhana<sup>6</sup>, Sushma Naithani<sup>6</sup>, Martin Ranik<sup>7</sup>, James Wesley-Smith<sup>8</sup>, Luke Solomon<sup>9</sup>, Pankaj Jaiswal<sup>6</sup>, Alexander A. Myburg<sup>7</sup> and Steven H. Strauss<sup>10</sup>

# Overview of methods for gene target selection



***We selected three new eucalypt gene targets for transformation***

Two expected to give male sterility, and one bisexual sterility but with ~normal flowers

# Three gene targets selected for knockout

- Male-sterility

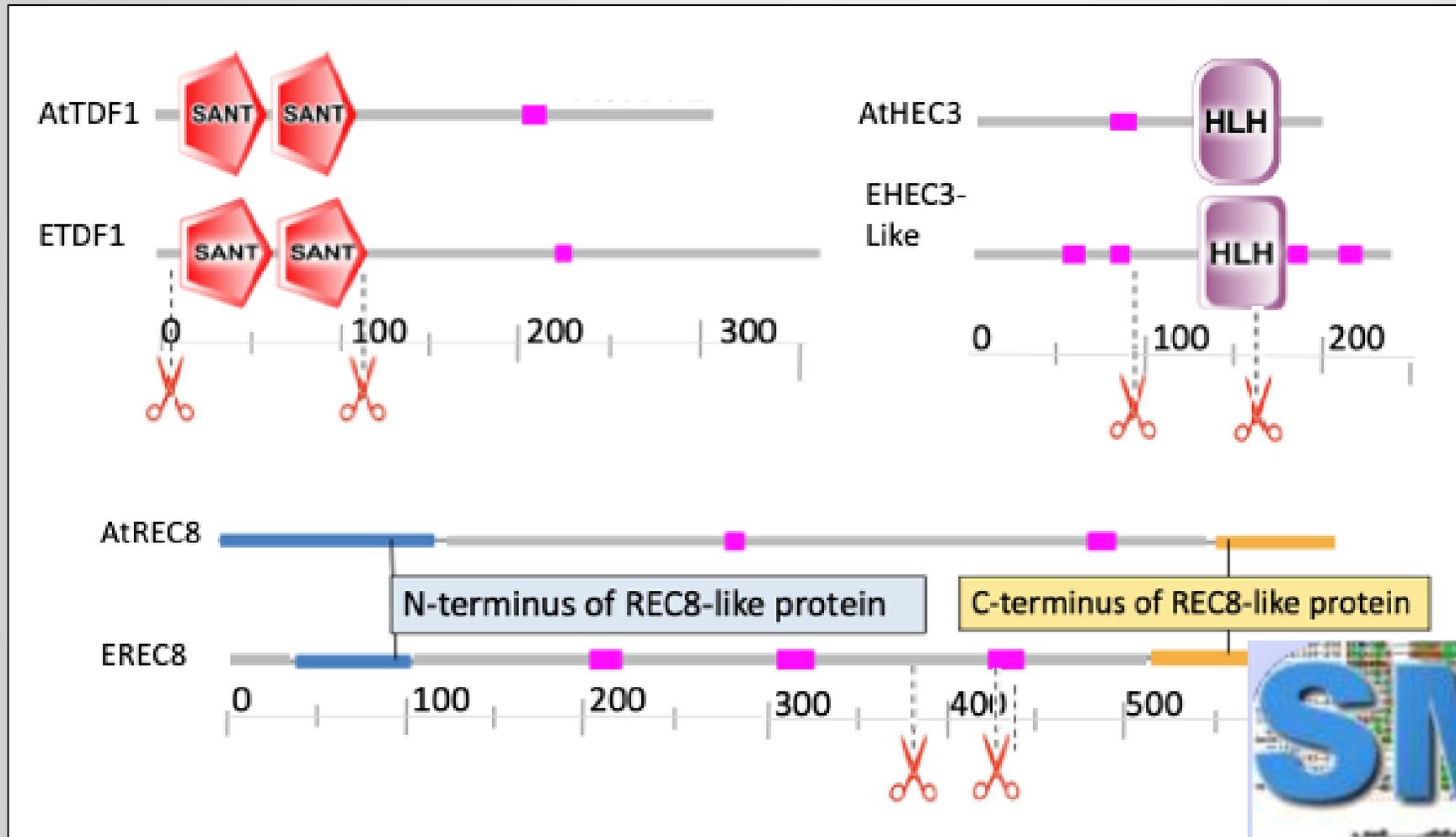
- *Eucgr.I02017*: Ortholog of male flowering transcription factor *TAPETAL DEVELOPMENT AND FUNCTION 1*
- *Eucgr.H04946*: Ortholog of flowering transcription factor *HECATE 3*

- Complete sterility

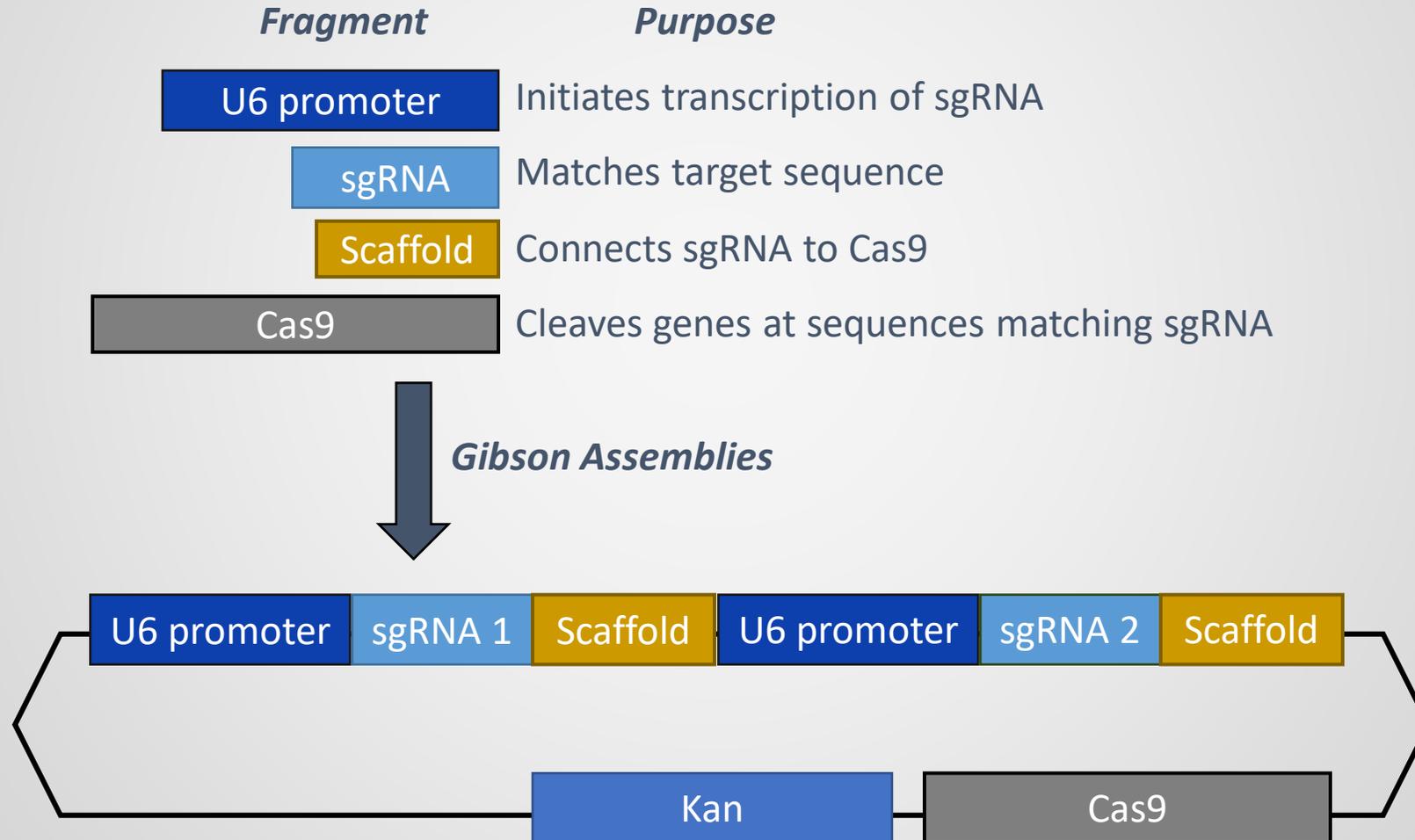
- *Eucgr.G03083*: Ortholog of meiotic cohesin *REC8 (SYNAPTIC 1)* so likely to produce ecologically functional flowers with pollen and nectar?



# Peptide domains predicted with SMART informed selection of sgRNA targets



# Fragments assembled via Gibson Assembly



# Transgenic CRISPR plants generated

- Use of *Agrobacterium* as gene transfer vector, producing stably transgenic CRISPR plants
- Early-flowering and wild-type backgrounds of model genotype-SP7 (provided by Futuragene) employed
- Allele-specific PCR and amplicon sequencing used to verify biallelic knockouts (frameshifts)



Plant Biotechnology  
Journal

aab SEB  
Association of Arabidopsis Biologists Society for Experimental Biology

Plant Biotechnology Journal (2015), pp. 1–12

doi: 10.1111/pbi.12431

## ***FT* overexpression induces precocious flowering and normal reproductive development in *Eucalyptus***

Amy L. Klocko<sup>1</sup>, Cathleen Ma<sup>1</sup>, Sarah Robertson<sup>1</sup>, Elahe Esfandiari<sup>1</sup>, Ove Nilsson<sup>2</sup> and Steven H. Strauss<sup>1,\*</sup>

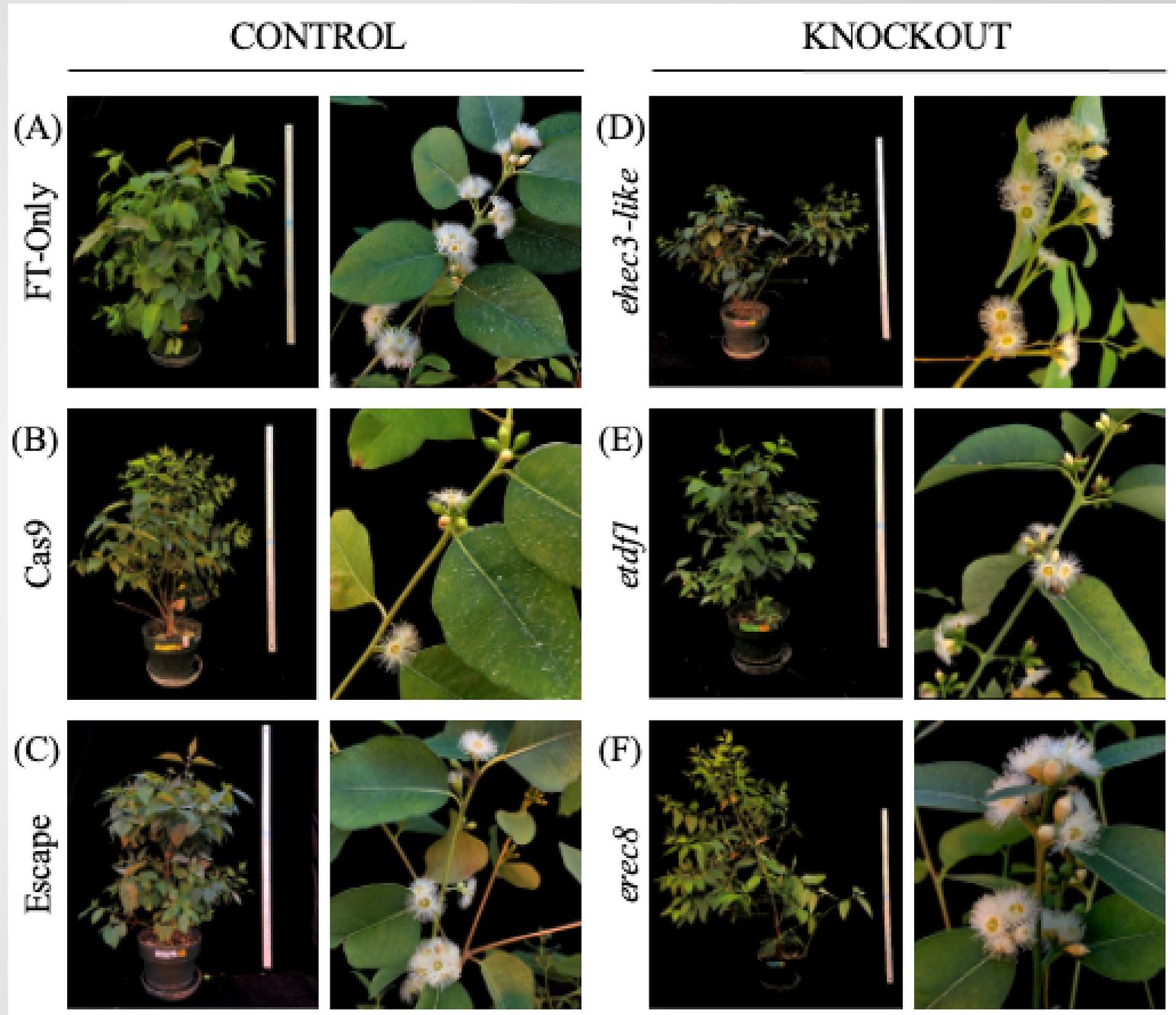
<sup>1</sup>Department Forest Ecosystems & Society, Oregon State University, Corvallis, OR, USA

<sup>2</sup>Department of Forest Genetics and Plant Physiology, Umeå Plant Science Centre, Swedish University of Agricultural Sciences, Umeå, Sweden

# High rates of mutations across gene targets

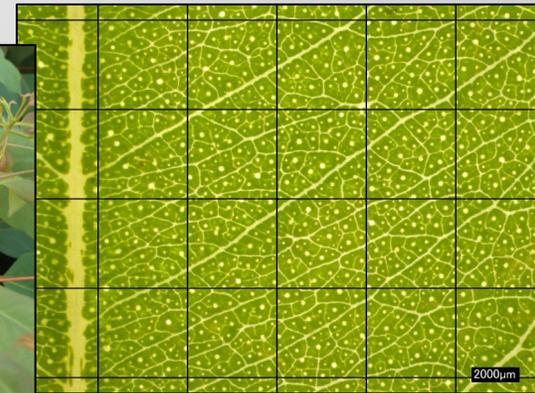
		<b><u>TDF1</u></b>	<b><u>REC8</u></b>	<b><u>HEC3</u></b>
<b>Specific mutation types</b>	Deletions	41.67%	51.14%	58.33%
	Insertions	9.72%	20.45%	13.89%
	Inversions	5.56%	0.00%	22.22%
	Compound mutations	2.78%	1.14%	1.39%
<b>Summary statistics</b>	Mutations / sgRNA sites	56.94%	72.73%	84.72%
	Frameshifts / alleles	55.56%	77.27%	88.89%

Visual inspection revealed no differences in vegetative, floral development in early flowering trees (normal bushy phenotypes)

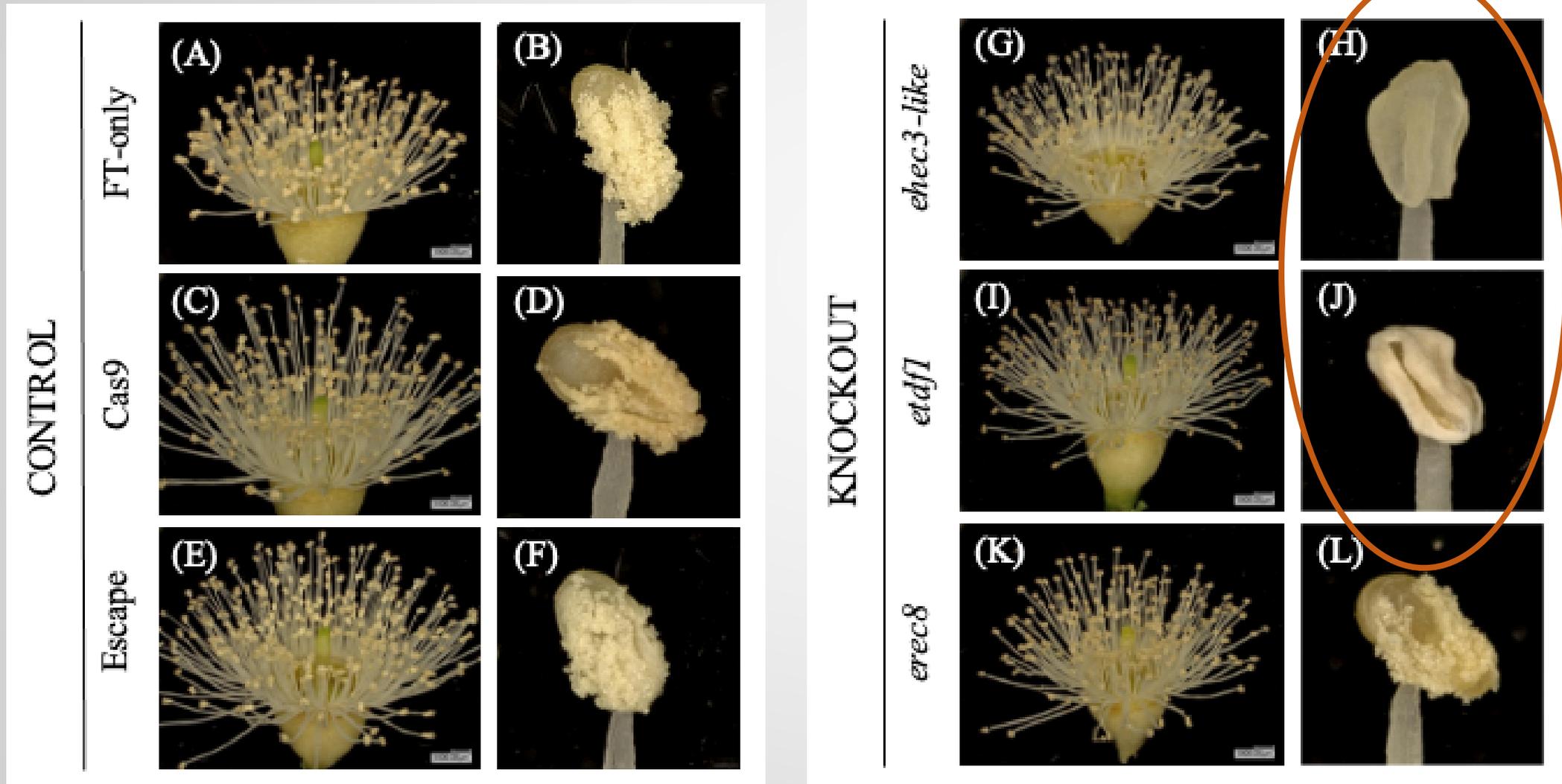


# Plants without FT overexpression were closely evaluated for effects of knockout on vegetative traits

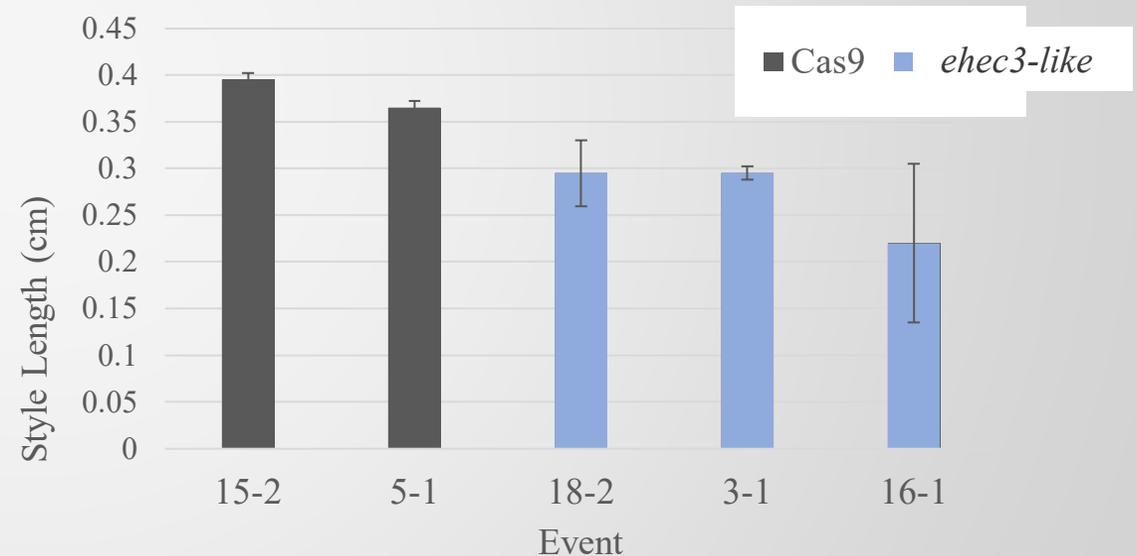
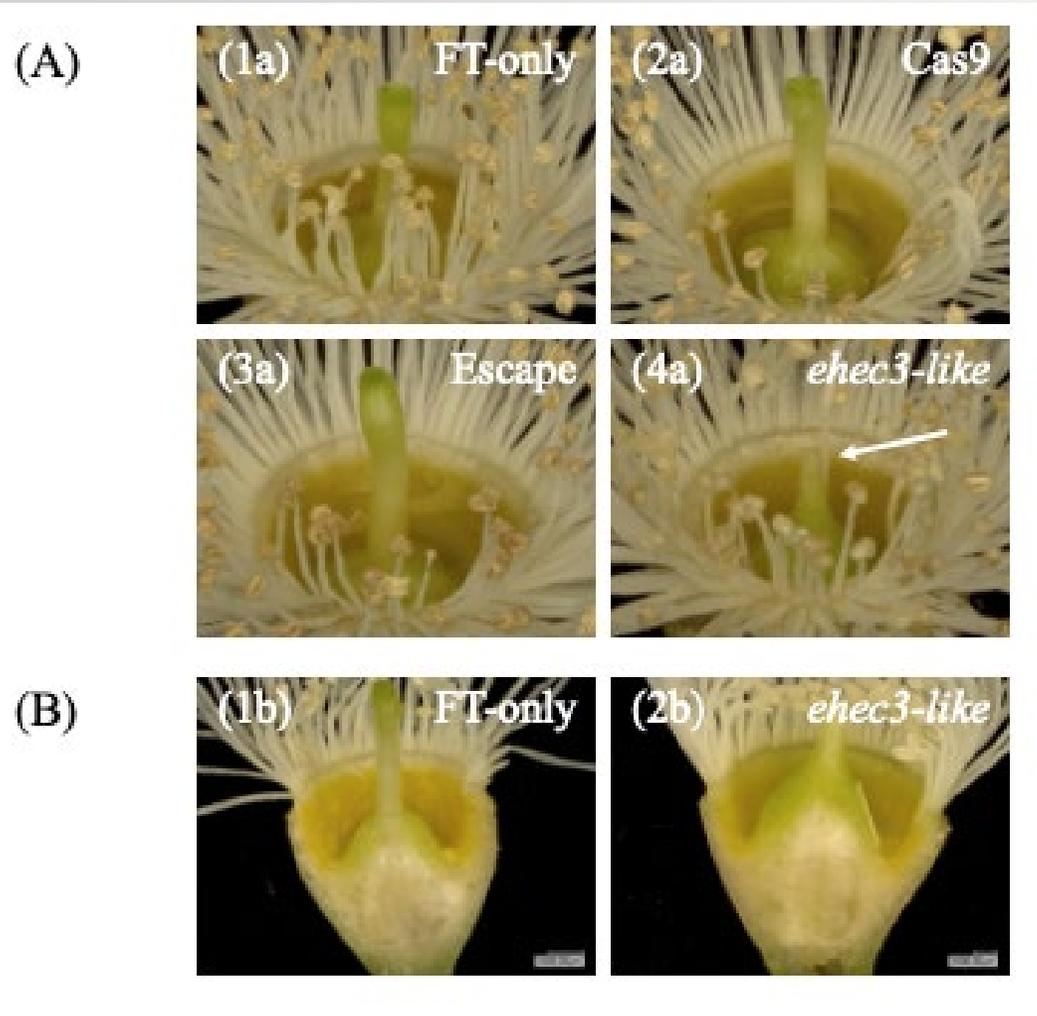
- Randomized greenhouse trials: KOs and Cas9 only and WT controls
- Measured markers of vegetative development: Leaf area, mass, SPAD (chlorophyll), oil duct count, plant diameter, tree height and stem diameter
- No statistically significant differences were found in any traits measured



*ehec3-like* and *tdf1* knockouts feature anthers lacking any pollen production

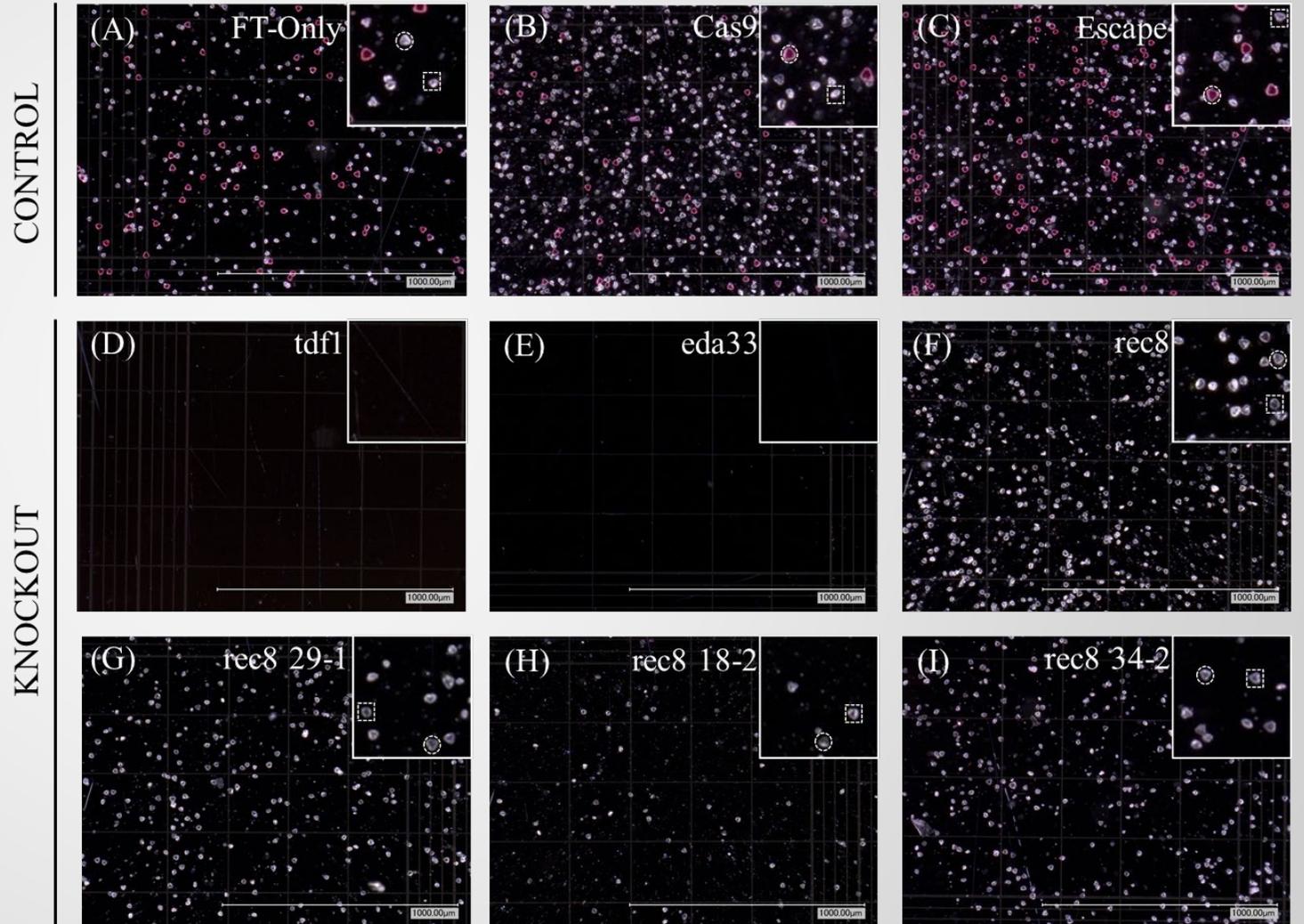


# Absence of stigma in *ehec3-like* knockout flowers, along with statistically significant reduction in style length



# Male infertility via lack of (viable) pollen for all knockouts

- Collection, staining and microscopy of pollen allows visualization of pollen presence and viability across samples
- Putatively viable pollen appears red



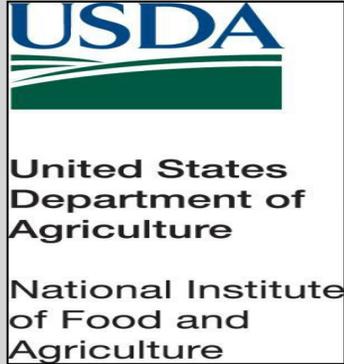
# Future work: Female fertility evaluation, ecological function in *rec8* KOs, field studies

- Controlled pollination assays to determine if female flowers are sterile
- Evaluation of nectar composition and pollen nutrition to determine whether ecological value for pollinators is affected in *rec8* knockouts
- Study of flowering in plants without FT overexpression
- Field trials to study pleiotropic effects on vegetative development

# Summary

- CRISPR/Cas9 with high KO rate
- KO phenotypes in flowers homologous to predictions from *Arabidopsis*/rice
- Male-sterility resulting from all three knockouts
- Putative functional but sterile flowers from *REC8* KOs
- Normal vegetative morphology and appearance in all KOs (no FT)
- Mutation of the Eucalyptus orthologs of *TDF1*, *REC8* and *HEC3-Like* show promise for imparting male-sterility or complete sterility without vegetative pleiotropy

# Thanks to our funders and collaborators



Biotechnology Risk Assessment

Grant # 2017-33522-27098

"CRISPR/Cas9 mutagenesis for genetic containment of forest trees"



GREAT TREES Consortium

Suzano, SAPPI, Arauco, Klabin, SweTree,

Thanks to Futuragene/Suzano for the model eucalypt genotype we transformed

# Thanks to the group



**Steve Strauss**  
Director TBGRC,  
Professor



**Cathleen Ma**  
Transformation &  
Greenhouse  
Experiments



**Amanda  
Goddard**  
Program & Field  
Manager



**Greg  
Goralogia**  
Postdoc,  
Flowering &  
Gene Editing



**Kate  
Peremyslova**  
GWAS,  
Transformation  
Experiments



**Xavier  
Tacker**  
Pollen viability



**Michael Nagle**  
PhD Candidate, GWAS,  
Transformation Genes



**Chris**  
Post-doc  
Hops  
transformation



**Nathan Ryan**  
Masters  
GRF genes



**Alexa Nino de  
Rivera**  
Tissue culture &  
greenhouse technician



**Megan  
McEldowney**  
Tissue culture &  
GWAS technician

20



**Surbhi  
Nahata**  
CRISPR  
mutations,  
greenhouse trial